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NBS Special Publication 500-117, Volume 2 Selection and Use of
General-Purpose Programming Languages - Program Examples

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Languages - Program Examples

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# Selection and Use of General-Purpose Programming Languages Volume 2 - Program Examples <br> John V. Cugini <br> Institute for Computer Sciences and Technology National Bureau of Standards 

ABSTRACT
Programming languages have been and will continue to be an important instrument for the automation of a wide variety of functions within industry and the Federal Government. Other instruments, such as program generators, application packages, query languages, and the like, are also available and their use is preferable in some circumstances.

Given that conventional programming is the appropriate technique for a particular application, the choice among the various languages becomes an important issue. There are a great number of selection criteria, not all of which depend directly on the language itself. Broadly speaking, the criteria are based on 1) the language and its implementation, 2) the application to be programmed, and 3) the user's existing facilities and software.

This study presents a survey of selection factors for the major general-purpose languages: Ada*, BASIC, C, COBOL, FORTRAN, Pascal, and PL/I. The factors covered include not only the logical operations within each language, but also the advantages and disadvantages stemming from the current computing environment, e.g., software packages, microcomputers, and standards. The criteria associated with the application and the user's facilities are explained. Finally, there is a set of program examples to illustrate the features of the various languages.

This volume includes the program examples. Volume 1 contains the discussion of language selection criteria.

Key words: Ada; alternatives to programming; BASIC; C; COBOL; FORTRAN; Pascal; PL/I; programming language features; programming languages; selection of programming language.

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```

In this volume, we shall illustrate the general style of each of the languages with a program. These programs are only examples; they do not attempt to demonstrate the full capability of each language. On the other hand, the application chosen is complex enough that the programs do make significant use of several important language features, such as reading a file, interacting with a user, recursion, data abstraction, manipulation of arrays, pointers, and character strings, and some numeric calculation. $0 f$ particular note are the language features for modularizing a program of moderate size (about 1000 lines). While no application can be completely language-neutral, this variety of requirements implies a relatively unbiased example. Finally, the application deals with a well-known realm (family relationships) in order to facilitate understanding of the programs.

All of the programs solve the same problem, i.e., they accept the same input and produce output as nearly equivalent as possible. The input is a file of people, one person per record, and a series of user queries. In the file, each person's father and mother (if known), and spouse (if any) are identified. Given this information, the user may then specify any two persons in the file, and the program computes and displays the relationship (e.g., brother-in-law, second cousin) between those two. Also, based on the number and degree of common ancestors, the expected value for the proportion of common genetic material between the two is computed and displayed.

The algorithms and data structures employed are roughly equivalent, but differ in detail owing to the language differences being illustrated. Generally, user-defined names are capitalized and language-defined keywords and identifiers are written in lower-case. In all the programs a directed graph is simulated, with the vertices representing people and the edges representing different types of direct relationships. The only direct relationships are parent, child, and spouse. Starting at one vertex, a search is conducted to find the shortest path to the other vertex. The types of edges encountered along the path, together with some additional information, determine the relationship. For instance, if the shortest path between Xl and X4 is that $X 1$ is child of $X 2$, $X 2$ is spouse of $X 3$, and $X 3$ is parent of $X 4$, this would show that $X 1$ and $X 4$ are step-siblings. It is assumed that the input file has already been validated and is correct. The user ${ }^{-}$s requests, however, are checked. The algorithm to determine the shortest path is adapted from [Baas78]. The overall algorithm is expressed by the pseudo-code below.

All of the programs, except the one in BASIC, have compiled and executed on at least one language processor which implements the corresponding standard or base document. The COBOL program, while conforming to both COBOL-74 and COBOL-8x, is essentially a COBOL-74 program, since it does not exploit any of the new COBOL-8x features.

```
            Figure 1 - Algorithm for Program Examples
for each record in input PEOPLE file do
    establish entry in PERSON array
    for all previous entries do
        compare this entry to previous, looking for
            immediate relationships: parent, child, or spouse
        if relationship found
            establish link (edge) between these two persons
        end if
    end for
end for
graph is now built
while not request to stop
    prompt and read next request
exit while-block if request to stop
    if syntax of request OK
        search for requested persons
        if exactly one of each person found
            if lst person = 2nd person
                            display "identical to self"
            else
                    find shortest path between the two persons
                    if no such path
                        display "unrelated"
                    else
                        analyze path for named relationships:
                        path initially composed of parent, child,
                        spouse edges
                                resolve child-parent and child-spouse-parent
                                    to sibling
                                    resolve child-child-... and parent-parent-...
                                    to descendant (child*) or ancestor (parent*)
                                    resolve child*-sibling-parent* to cousin,
                                    child*-sibling to nephew,
                                    sibling-parent* to uncle
                                    display consolidated relationships
                                    compute proportion of common genetic material:
                                    traverse ancestors of personl, zeroing out
                                    traverse ancestors of personl, marking and
                                    accumulating genetic contribution
                                    traverse ancestors of person2, accumulating
                                    overlap with personl
                                    display results
                        end if
            end if
        else
            display "duplicate name" or "not found"
        end if
    else
        display "invalid request"
    end if
end while
display "done"
```

Figure 2 - Input Data
This figure shows some of the input data with which the program examples were tested. The format of each record is:

| Position | Contents |
| ---: | :--- |
| -20 | Name of person |
| $21-23$ | Unique 3-digit identifier of person |
| 24 | Gender of person |
| $25-27$ | Identifier of father ( 000 if unknown) |
| $28-30$ | Identifier of mother ( 000 if unknown) |
| $31-33$ | Identifier of spouse ( 000 if none or unknown) |

Example of Input Data:

| John Smith | 001M000000002 |
| :---: | :---: |
| Mary Smith | 002F003000001 |
| Wilbur Finnegan | 010 M 000000011 |
| Mary Finnegan | 011 F 000000010 |
| James Smith | 020M001002022 |
| Wilma Smith | 022F010011020 |
| Marvin Hamlisch | 031 m 000032000 |
| Melvin Hamiisch | $033 \mathrm{M000032000}$ |
| Martha Hamlisch | 032 F 048043034 |
| Murgatroyd Whatsis | $034 \mathrm{M000000032}$ |
| Bentley Whatsis | 035 M 034036000 |
| Myrna Whozat | 036F000000000 |
| Bosworth Whatsis | 037 M 034036000 |
| K48 | 048 M 000000043 |
| K43 | 043F041042048 |
| K41 | 041 M 000000042 |
| K42 | 042F000000041 |
| K46 | 046 M 045000000 |
| K45 | 045M048043000 |
| K47 | 047 M 044000000 |
| K44 | 044M041042000 |
| Velorus Davis | 085M000000086 |
| Goldie Beacon | 083F085086082 |
| Ross Beacon | 082M000000083 |
| Velma Davis | 086F000000085 |
| Floyd Davis | 088M085084087 |
| Cindy Davis | 084F000000000 |
| David Beacon | 121 M 081120000 |
| Norma Cousins | 053 F 082083055 |
| Carmine Cousins | 051 M 000000052 |
| Maria Cousins | 052F000000051 |
| James Cousins | 054M051052000 |
| C. John Cousins | 055 M 051052053 |
| John Cousins | $073 \mathrm{M055053074}$ |
| Janet Cousins | 074 F 140141073 |
| Richard Cousins | 077 M 073074000 |
| Paul Cousins | 078 MO 073074000 |
| Marie Cousins | 079 F 073074000 |

Figure 3 - Queries and Output

This figure gives some examples of the results of running the programs.

```
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
;
    Incorrect request format: null field preceding semicolon.
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
x;x;x
    Incorrect request format: must be exactly one semicolon.
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
x; x
    First person not found.
    Second person not found.
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
        111 ; 111
    Christopher Delmonte is identical to himself.
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
G6;John Smith
    G6 is not related to John Smith
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
Carmine Cousins;111
    Duplicate names for first person - use numeric identifier.
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
163;145
    Shortest path between identified persons:
    Linda Lackluster is child of
    Millie Lackluster is child of
    Anna Pittypat is parent of
    Margaret Madison is spouse of
    Richard Madison
    Victoria Pisces
    is child of
    is parent of
    is parent of
Elzbieta Gotsocks
Condensed path:
Linda Lackluster is niece of
Richard Madison is uncle of
Elzbieta Gotsocks
Proportion of common genetic material=0.00000E+00
```

```
    Figure 3 - Queries and Output (continued)
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
094;145
    Shortest path between identified persons:
    Nancy Powers is child of
    Maxine Powers is child of
    Floyd Davis is child of
    Velorus Davis is parent of
    Goldie Beacon is parent of
    Norma Cousins is parent of
    John Cousins is spouse of
    Janet Cousins is child of
    Richard Madison is child of
    Victoria Pisces is parent of
    Maria Gotsocks is parent of
    Elzbieta Gotsocks
    Condensed path:
    Nancy Powers is 2nd half-cousin-in-law of
    Janet Cousins is cousin of
    Elzbieta Gotsocks
    Proportion of common genetic material = 0.00000E +00
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
036;033
    Shortest path between identified persons:
    Myrna Whozat is parent of
    Bentley Whatsis is child of
    Murgatroyd Whatsis is spouse of
    Martha Hamlisch is parent of
    Melvin Hamlisch
    Condensed path:
    Myrna Whozat
    is mother of
    Bentley Whatsis is step-brother of
    Melvin Hamlisch
    Proportion of common genetic material=0.00000E+00
    Enter two person-identifiers (name or number),
    separated by semicolon. Enter "stop" to stop.
031;033
    Shortest path between identified persons:
Marvin Hamlisch is child of
Martha Hamlisch is parent of
Melvin Hamlisch
Condensed path:
Marvin Hamlisch is half-brother of
Melvin Hamlisch
Proportion of common genetic material=2.50000E-01
```

```
Figure 3 - Queries and Output (continued)
```

Enter two person-identifiers (name or number), separated by semicolon. Enter "stop" to stop.
145;090
Shortest path between identified persons:
Elzbieta Gotsocks is child of
Maria Gotsocks is child of
U. Pisces
is parent of
Richard Madison is parent of
Janet Cousins is spouse of
John Cousins is child of
Norma Cousins is child of
Goldie Beacon is child of
Velorus Davis is parent of
Floyd Davis
Maxine Powers
is parent of
is spouse of
Tim Powers
Condensed path:
Elzbieta Gotsocks is cousin-in-law of
John Cousins is half-cousin-in-law once removed of
Tim Powers
Proportion of common genetic material $=0.00000 \mathrm{E}+00$
Enter two person-identifiers (name or number),
separated by semicolon. Enter "stop" to stop.
L6;R9
Shortest path between identified persons:
L6 is child of
L5 is child of
L4 is child of
L3 is child of
L2 is child of
L1 is child of
L0 is parent of
R1 is parent of
R2 is parent of
R3 is parent of
R4 is parent of
R5 is parent of
R6 is parent of
R7 is parent of
R8 is parent of
R 9
Condensed path:
L6 is $5 t h$ half-cousin 3 times removed of
R 9
Proportion of common genetic material $=3.05176 \mathrm{E}-05$

## Figure 3 - Queries and Output (continued)

Enter two person-identifiers (name or number), separated by semicolon. Enter "stop" to stop. W1; R14
Shortest path between identified persons:
Wl is spouse of
L0 is parent of
R1 is parent of
R2 is parent of
R3 is parent of
R4 is parent of
R5 is parent of
R6 is parent of
R7 is parent of
R8 is parent of
R9 is parent of
R10 is parent of
R11 is parent of
R12 is parent of
R13 is parent of
R14
Condensed path:
W 1
is great*12-grand-step-father of
R 14
Proportion of common genetic material $=0.00000 \mathrm{E}+00$
Enter two person-identifiers (name or number), separated by semicolon. Enter "stop" to stop. X8; L 6
Shortest path between identified persons:
X8 is child of

X7 is child of
X6 is child of
X5 is child of
X4 is child of
X3 is spouse of
R4 is child of
R3 is child of
R2 is child of
R1 is child of
L0 is parent of
L1 is parent of
L2 is parent of
L3 is parent of
L4 is parent of
L5 is parent of
L 6
Condensed path:
X8 is great*3-grand-step-son of
R4 is 3rd half-cousin 2 times removed of
L 6
Proportion of common genetic material $=0.00000 \mathrm{E}+00$

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Figure 3 - Queries and Output (continued)
Enter two person-identifiers (name or number), separated by semicolon. Enter "stop" to stop. G5; G6
Shortest path between identified persons:
G 5
G 6
Condensed path:
G5 is mother of
G 6
Proportion of common genetic material $=5.62500 \mathrm{E}-01$
Enter two person-identifiers (name or number), separated by semicolon. Enter "stop" to stop. stop
End of relation-finder.
2.0 ADA

```
---- first compilation-unit \#l is package of global types and objects
package RELATION_TYPES_AND_DATA is
    MAX PERSONS : constant integer := 300;
    NAME_LENGTH : constant integer := 20;
    -- every PERSON has a unique 3-digit IDENTIFIER
    IDENTIFIER LENGTH : constant integer :=3;
    BUFFER LENGTH : constant integer :=60;
    subtype NAME RANGE is integer range 1.. NAME LENGTH;
    subtype IDENTIFIER RANGE is integer range 1..IDENTIFIER LENGTH;
    subtype BUFFER_RANGE is integer range 1..BUFFER_LENGTH;
    subtype NAME TYPE is string (NAME RANGE);
    subtype BUFFER TYPE is string (BUFFER RANGE) ;
    subtype MESSAGE_TYPE is string (1..40);
    subtype INDEX_TYPE is integer range O..MAX_PERSONS;
    subtype COUNTER is integer range 0..integer'last;
    subtype DIGIT_TYPE is character range \({ }^{\prime} 0^{\prime} . .^{\prime} 9^{\prime}\);
    type REAL is digits 6;
type IDENTIFIER TYPE is array (IDENTIFIER RANGE) of DIGIT TYPE;
-- each PERSON- \(\bar{s}\) record in the file identifies at most three
-- others directly related: father, mother, and spouse
type GIVEN IDENTIFIERS is (FATHER_IDENT, MOTHER IDENT, SPOUSE IDENT);
type RELATIVE_ARRAY is array ( \(\bar{G} I V E N\) IDENTIFIERS) of IDENTIFIER_TYPE;
NULL_IDENT : constant IDENTIFIER_TYPE := "000";
REQUEST_OK \(O\) : constant MESSAGE_TYPE \(\quad:=\)
"Request OK
;
REQUEST_TO_STOP : constant BUFFER_TYPE :=
    "stop
type GENDER TYPE is (MALE, FEMALE);
type RELATIŌN TYPE is (PARENT, CHILD, SPOUSE, SIBLING, UNCLE,
    NEPHEW, COUSIN, NULL RELATION);
- directed edges in the graph are of a given subtype
subtype EDGE TYPE is RELATION_TYPE range PARENT..SPOUSE;
-- A node in the graph (= PERSON) has either already been reached,
-- is immediately adjacent to those reached, or farther away.
type REACHED TYPE is (REACHED, NEARBY, NOT_SEEN);
-- each PERSON has a linked list of adjacent nodes, called neighbors
type NEIGHBOR_RECORD;
type NEIGHBOR POINTER is access NEIGHBOR RECORD;
type NEIGHBOR_RECORD is
    record
        NEIGHBOR INDEX : INDEX TYPE;
        NEIGHBOR EDGE : EDGE TYPE;
        NEXT NEI \(\bar{G} H B O R\) : NEIGHBOR POINTER;
    end record;
```

-- All relationships are captured in the directed graph of which
-- each record is a node.
type PERSON_RECORD is
record
-- static information - filled from PEOPLE file:
NAME : NAME TYPE; IDENTIFIER : IDENTIFIER_TYPE; GENDER : GENDER TYPE; -- IDENTIFIERs of immediate relatives - father, mother, spouse RELATIVE_IDENTIFIER : RELATIVE_ARRAY; -- head of linked list of ad jacent nodes NEIGHBOR_LIST_HEADER : NEIGHBOR_POINTER;
-- data used when traversing graph to resolve user request: DISTANCE FROM SOURCE : REAL; PATH PREDECES $\bar{S} O R$ : INDEX TYPE; EDGE ${ }^{-}$TO PREDECESSOR : EDGE TYPE; REAC $\overline{H E D}$ - STATUS : REAC $\overline{H E D}$ TYPE;

- data used to compute common genetic material DESCENDANT_IDENTIFIER : IDENTIFIER_TYPE; DESCENDANT GENES : REAL;
end record;
-- the PERSON array is the central repository of information
- about inter-relationships.

PERSON : array (INDEX_TYPE) of PERSON_RECORD;

- utility to truncate or fill with spaces
procedure COERCE_STRING (SOURCE : in string; TARGET : in out string);
end RELATION_TYPES_AND_DATA;
-- - - - - END SPECIFICATION - - BEGIN BODY - - - - - - -
package body RELATION_TYPES_AND_DATA is
procedure COERCE STRING ( $\overline{\mathrm{SOUR}} \overline{\mathrm{C} E}$ : in string; TARGET : in out string) is MANY_SPACES : constant string (1..100) :=
$\because$

$$
" \&
$$

";
begin
if SOURCE ${ }^{-1}$ length < TARGET ${ }^{-}$length then
TARGET (TARGET first..TARGET ${ }^{-}$first + SOURCE $^{-}$length - 1) := SOURCE;

MANY SPACES (1..TARGET'1ength - SOURCE ${ }^{\prime}$ length);
else -- SOURCE longer than TARGET
TARGET := SOURCE (SOURCE-first..SOURCE'first + TARGET'length - 1);
end if;
end COERCE STRING;
end RELATION_TYPES_AND_DATA;

```
--- new compilation-unit #2: main line of execution RELATE
with RELATION TYPES AND DATA, text_io, sequential_io;
use RELATION_TYPES AND DATA, text_io;
procedure RELATE is
    -- this is the format of records in the file to be read in
    type FILE GENDER is ('M`, 'F`);
    type FILE PERSON RECORD is
        record
            NAME : NAME TYPE;
            IDENTIFIER : IDENTIFIER TYPE;
            -- 'M' for MALE and 'F' for FEMAL\overline{LE}
            GENDER : FILE GENDER;
            RELATIVE IDENTIFIER : RELATIVE ARRAY;
        end record;
    -- Instantiate generic package for file IO.
    package PEOPLE IO is
        new sequential_io (ELEMENT_TYPE => FILE_PERSON_RECORD);
    -- These variables are used when establishing the PERSON array
    -- from the PEOPLE file.
    PEOPLE : PEOPLE IO . FILE TYPE;
    PEOPLE RECORD : FILE PERSON RECO\overline{RD;}
    CURREN\overline{T}, NUMBER_OF PERSONS
                            : INDEX TYPE;
PREVIOUS_IDENT, CURRENT IDENT
    : TDENTIFIER TYPE;
    RELATIONSHIP : GIVEN_IDENTIFIERS;
    -- These variables are used to accept and resolve requests for
    -- RELATIONSHIP information.
    BUFFER_INDEX, SEMICOLON LOCATION
            : BU\overline{FFER RANGE;}
    REQUEST BUFFER : BUFFER TYPE;
    PERSON1 IDENT, PERSON2 IDENT
    : NA\overline{ME TYPE;}
    PERSON1 FOUND, PERSON2 FOUND
    : COUNTER;
    : MESSAGE TYPE;
    ERROR MESSAGE : MESSAGE 
    : INNDEX TYPE;
```

```
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```

```
-- declare procedures directly invoked from RELATE:
procedure LINK RELATIVES (FROM_INDEX : in INDEX_TYPE;
                                    RELATIONSHIP : in GIVEN_IDENTIFIERS;
                                    TO_INDEX : in INDEX_TYPE)
    is separate;
procedure PROMPT AND READ is separate;
procedure CHECK \overline{REQUEST (REQUEST STATUS : out MESSAGE TYPE;}
                                SEMICOLON LOCATION : out BUFFER \overline{RANGE)}
    is separate;
procedure BUFFER_TO_PERSON (PERSON ID : in out NAME_TYPE;
                        START L̄OCATION,
                        STOP LOCATION : in BUFFER_RANGE)
    is separate;
procedure SEARCH FOR REQUESTED PERSONS
            (PERSON1_IDENT, \overline{PERSON2_IDENT : in NAME_TYPE;}
            PERSON1-INDEX, PERSON2_INDEX : out INDE\overline{X TYPE;}
            PERSON1_FOUND, PERSON2_FOUND : in out COUNTER)
        is separate;
procedure FIND RELATIONSHIP (TARGET INDEX, SOURCE_INDEX : in INDEX TYPE)
    is separate;
```

-- *** execution of main sequence begins here *** --
begin
PEOPLE IO . open (PEOPLE, PEOPLE IO . IN FILE, "PEOPLE.DAT");
-- CURRENT location in array being fillē
CURRENT := 0;
-- This loop reads in the PEOPLE file and constructs the PERSON
- array from it (one PERSON = one record $=$ one array entry).
-- As records are read in, links are constructed to represent the
-- PARENT-CHILD or SPOUSE RELATIONSHIP. The array then implements
-- a directed graph which is used to satisfy subsequent user
-- requests. The file is assumed to be correct - no validation
-- is performed on it.
READ IN PEOPLE:
whīle not PEOPLE IO . end of file (PEOPLE) loop
PEOPLE IO • reād (PEOPLE, $\overline{\text { PEOPLE RECORD) ; }}$
CURRENT := CURRENT+1;
-- copy direct information from file to array
PERSON (CURRENT) . NAME $:=$ PEOPLE RECORD . NAME;
PERSON (CURRENT) • IDENTIFIER $:=$ PEOPLE RECORD • IDENTIFIER;
if PEOPLE RECORD . GENDER $={ }^{-} M^{\prime}$ then
PERSON (CURRENT) - GENDER := MALE;
else
PERSON (CURRENT) - GENDER := FEMALE;
end if;
PERSON (CURRENT) • RELATIVE IDENTIFIER :=
PEOPLE_RECORD • RELATIVE_IDENTIFIER;
-- Location of adjacent persōns as yet undetermined
PERSON (CURRENT) - NEIGHBOR LIST HEADER := null;
-- Descendants as yet undetērmined
PERSON (CURRENT) • DESCENDANT IDENTIFIER : = NULL IDENT;
CURRENT_IDENT := PERSON (CURRENT) • IDENTIFIER;

```
    -- Compare this PERSON against all previously entered PERSONs
    -- to search for RELATIONSHIPs.
COMPARE TO PREVIOUS:
    for PREVIOUS in 1..CURRENT-1 loop
        PREVIOUS IDENT := PERSON (PREVIOUS) . IDENTIFIER;
        RELATIONS
            -- Search for father, mother, or spouse relationship in
            -- either direction between this and PREVIOUS PERSON.
            -- Assume at most one RELATIONSHIP exists.
TRY ALL RELATIONSHIPS:
            100p
                if PERSON (CURRENT) . RELATIVE IDENTIFIER (RELATIONSHIP) =
                PREVIOUS IDENT
            then
                LINK RELATIVES (CURRENT, RELATIONSHIP, PREVIOUS);
                exit TRY ALL RELATIONSHIPS;
        else
            if CURRENT IDENT =
                    PERSON (PREVIOUS) . RELATIVE IDENTIFIER (RELATIONSHIP)
            then
                    LINK RELATIVES (PREVIOUS, RELATIONSHIP, CURRENT);
                    exit TRY_ALL_RELATIONSHIPS;
            end if;
        end if;
        if RELATIONSHIP < SPOUSE IDENT then
            RELATIONSHIP := GIVEN IDENTIFIERS` succ(RELATIONSHIP);
        else
            exit TRY ALL RELATIONSHIPS;
        end if;
        end loop TRY ALL RELATIONSHIPS;
    end loop COMPARE TO PREVIOUS;
    end loop READ IN PEOPELE;
    NUMBER_OF PER\overline{SON\overline{S}}:= CURRENT;
    PEOPLE_IO-. close (PEOPLE);
    -- PERSON array is now loaded and edges between immediate relatives
    -- (PARENT-CHILD or SPOUSE-SPOUSE) are established.
    -- While-1oop accepts requests and finds RELATIONSHIP (if any)
    -- between pairs of PERSONs.
```

```
READ AND PROCESS REQUEST:
    100p
        PROMPT AND READ;
    exit READ_AND PROCESS REQUEST when REQUEST BUFFER = REQUEST TO_STOP;
        CHECK RE}QUE\overline{ST (ERROR_MESSAGE, SEMICOLON_D_LOCATION);
        -- Syntax check of request completed. Now either display error
        - message or search for the two PERSONs.
        if ERROR_MESSAGE = REQUEST_OK then
            -- Request syntactically correct -
                            -- search for requested PERSONs.
            BUFFER TO PERSON (PERSON1 IDENT, 1, SEMICOLON LOCATION - 1);
            BUFFER TO PERSON (PERSON2 IDENT, SEMICOLON LOCATION + 1, BUFFER LENGTH);
            SEARCH_FOR_REQUESTED_PERSONS (PERSON1_IDENT, PERSON2 IDENT,
                        PERSON1 INDEX, PERSON2 INDEX,
                        PERSON1 FOUND, PERSON2 FOUND);
            if (PERSON1 FOUND = 1) and (PERSON2 FOUND = 1) then
                -- Exact\overline{ly one match for each PERSON - proceed to}
                - determine RELATIONSHIP, if any.
                if PERSON1 INDEX = PERSON2 INDEX then
                put ( }-=|\mathrm{ PERSON (PERSON1_INDEX) . NAME &
                " is identical to ");
                if PERSON (PERSON1 INDEX) . GENDER = MALE then
                        put_line("himse\overline{1f.");}
                else
                    put line("herself.");
                end if;
                else
                        FIND RELATIONSHIP (PERSON1 INDEX, PERSON2 INDEX);
                end if;
            else -- either not found or more than one found
                if PERSON1_FOUND = 0 then
                put line (" First person not found.");
                elsif PERSON1 FOUND > 1 then
                    put_line (" Duplicate names for first person - use" &
                        " numeric identifier.");
                end if;
                if PERSON2 FOUND = 0 then
                    put line (" Second person not found.");
                elsif PERSON2 FOUND > 1 then
                    put_line (" Duplicate names for second person - use" &
                                    " numeric identifier.");
                end if;
            end if; -- processing of syntactically legal request
        else
            put_line (" Incorrect request format: " & ERROR_MESSAGE);
        end if;
    end loop READ_AND_PROCESS_REQUEST;
    put line (" End of relation-finder.");
end RELATE;
```

```
--- new compilation-unit #3: procedures under RELATE
separate (RELATE)
procedure LINK_RELATIVES (FROM_INDEX : in INDEX TYPE;
                                    RELATIONSHIP : in GIVEN IDENTIFIERS;
                                    TO INDEX : in INDEX TYPE) is
    -- establishes cross-indexing between immediately related PERSONs.
    procedure LINK_ONE_WAY (FROM_INDEX : in INDEX TYPE;
                            THIS_EDGE : in EDGE TYPE;
                            TO INDEX : in INDE\overline{X TYPE) is}
        -- Establishes the NEIGHBOR RECORD from one PERSON to another
    NEW_NEIGHBOR : NEIGHBOR_POINTER;
    begin
        NEW NEIGHBOR := new NEIGHBOR RECORD
            `(NEIGHBOR_INDEX => T\overline{O}_INDEX,
                NEIGHBOR EDGE }=>\mathrm{ THITS EDGE,
                        NEXT NEIG\overline{GBOR => PERSO}N (FROM INDEX) . NEIGHBOR LIST_HEADER);
        PERSON (FROM_\overline{INDEX) . NEIGHBOR_LIST_HEAD\overline{DER := NEW_NEIGHBO\overline{R}};};\mathbf{}\mathrm{ _}
    end;
begin -- execution of LINK RELATIVES
    if RELATIONSHIP = SPOUSE IDENT then
            LINK ONE WAY (FROM INDEX, SPOUSE, TO INDEX);
            LINK ONE WAY (TO INDDEX, SPOUSE, FROM INDEX);
    else =- RELATIONS\overline{HIP is father or mother}
        LINK ONE WAY (FROM INDEX, PARENT, TO INDEX);
        LINK_ONE_WAY (TO_INDEX, CHILD, FROM_INDEX);
    end if;
end LINK_RELATIVES;
separate (RELATE)
procedure PROMPT_AND_READ is
    -- Issues prompt for user-request, reads in request,
    -- blank-fills buffer, and skips to next line of input.
    LAST_FILLED : natural;
begin
    put_line (" ");
    put_line (" ----------------------------------------------------");
    put line (" Enter two person-identifiers (name or number),");
    put_line (" separated by semicolon. Enter ""stop"" to stop.");
    get_line (REQUEST BUFFER, LAST FILLED);
    COERCE STRING (" "', REQUEST_BUFFER (LAST_FILLED+1..BUFFER_LENGTH));
end PROM\overline{PT_AND_READ;}
```

```
separate (RELATE)
procedure CHECK_REQUEST (REQUEST STATUS : out MESSAGE TYPE;
                                    SEMICOLON_LOCATION : out BUFFER_ \(\bar{R} A N G E)\) is
    -- Performs syntactic check on request in buffer.
    SEMICOLON COUNT : COUNTER;
    PERSON1_FIELD_EXISTS, PERSON2_FIELD_EXISTS
        : boolean;
begin
    REQUEST STATUS := REQUEST_OK;
    SEMICOLŌN LOCATION := 1;
    PERSON1_FİELD_EXISTS := false;
    PERSON2 FIELD_EXISTS := false;
    SEMICOLON COUNT : \(=0\);
    for BUFFER INDEX in BUFFER RANGE loop
        if REQUE \(\overline{\mathrm{S}} \mathrm{T}\) BUFFER (BUFFE \(\bar{R}\) INDEX) /= - then
            if REQUEST BUFFER (BUFFER INDEX) \(=\) '; ' then
                SEMICOLON LOCATION := \(\bar{B} U F F E R\) INDEX;
                SEMICOLONCOUNT := SEMICOLON COUNT +1 ;
            else -- Chēck for non-blanks befōre/after semicolon.
                if SEMICOLON COUNT < 1 then
                    PERSON1 F \(\bar{I} E L D\) EXISTS := true;
                else
                    PERSON2_FIELDEXISTS := true;
                end if;
            end if;
        end if;
    end loop;
    -- set REQUEST STATUS, based on results of scan of REQUEST BUFFER.
    if SEMICOLON COUNT /= 1 then
        REQUEST ST̄ATUS := "must be exactly one semicolon. ";
    elsif not PERSON1_FIELD_EXISTS then
        REQUEST STATUS \({ }^{-}:=\)"nūll field preceding semicolon. ";
    elsif not PERSON2 FIELD EXISTS then
        REQUEST_STATUS \({ }^{-}:=\)"nū1l field following semicolon. ";
    end if;
end CHECK_REQUEST;
separate (RELATE)
procedure BUFFER_TO_PERSON (PERSON_ID : in out NAME_TYPE;
                                    Start Location,
                                    STOP LOCATION : in BUFFER RANGE) is
    - fills in the PERSON ID from the designated portion
    -- of the REQUEST BUFFER.
    FIRST_NON_BLANK : BUFFER_RANGE;
begin
    FIRST_NON BLANK := START LOCATION;
    while \({ }^{-}\)REQ \(\bar{U} E S T\) BUFFER ( \(F I \overline{R S T}\) _NON_BLANK) \(=\) - - loop
        FIRST_NON_BLANK := FIRST_- NON_BLANK +1 ;
    end loop;
    COERCE_STRING (REQUEST BUFFER (FIRST_NON_BLANK..STOP_LOCATION),
                                    PERSON_ID);
end BUFFER_TO_PERSON;
```

```
separate (RELATE)
procedure SEARCH FOR REQUESTED PERSONS
                                    (PERSON\1_IDENT, \overline{PERSON2_IDENT : in NAME TYFE;}
                                    PERSON1-INDEX, PERSON2_INDEX : out INDE\overline{X} TYEE;
                            PERSON1 FOUND, PERSON2 FOUND : in out COUNTRI) te
    -- SEARCH FOR_REQUESTED_PERSONS scans through the PERSON arteg.
    - looking for the two requested PERSONs. Match may be by,
    -- or unique IDENTIFIER-number.
    THIS_IDENT : NAME_TYPE;
begin
    PERSON1 FOUND := 0;
    PERSON2 FOUND := 0;
    PERSON1 INDEX := 0;
    PERSON2 INDEX := 0;
SCAN ALL PERSONS:
    for
        -- THIS IDENT contains CURRENT PERSON's numeric IDENTTY 1me
        -- left-justified, padded with blanks.
        COERCE STRING (" ", THIS IDENT);
        for IDENTIFIER INDEX in IDDENTIFIER RANGE loop
            THIS IDENT (\overline{IDENTIFIER INDEX) :=}
                            PERSON (CURRENT) - IDENTIFIER (IDENTIFIER INDEX);
        end loop;
        -- allow identification by name or number.
        if (PERSON1_IDENT = THIS_IDENT) or
            (PERSON1_IDENT = PERSONN (CURRENT) . NAME)
        then
            PERSON1 FOUND := PERSON1_FOUND + 1;
            PERSON1_INDEX := CURRENT;
        end if;
        if (PERSON2_IDENT = THIS IDENT) or
            (PERSON2_IDENT = PERSŌN (CURRENT) . NAME)
        then
            PERSON2 FOUND := PERSON2 FOUND + 1;
            PERSON2_INDEX := CURRENT;
        end if;
    end loop SCAN ALL PERSONS;
end SEARCH_FOR__REQUESTED_PERSONS;
```

```
separate (RELATE)
procedure FIND RELATIONSHIP (TARGET INDEX, SOURCE INDEX : in INDEX TYPE) is
    -- Finds shōrtest path (if any) bētween two PERS̄ONs and
    -- determines their RELATIONSHIP based on immediate relations
    - traversed in path. PERSON array simulates a directed graph,
    -- and algorithm finds shortest path, based on following
    -- weights: PARENT-CHILD edge \(=1.0\)
    -- \(\quad\) SPOUSE-SPOUSE edge \(=1.8\)
    type SEARCH_TYPE is (SEARCHING, SUCCEEDED, FAILED);
    SEARCH STATUS : SEARCH TYPE;
    THIS_NODE, ADJACENT_NODE, BEST_NEARBY_INDEX, LAST_NEARBY_INDEX
    : INDEX TYPE;
    NEARBY_NODE : array (INDEX_TYPE) of INDEX_TYPE;
    THIS EDGE : EDGE TYPE;
    THIS \({ }^{-}\)NEIGHBOR : NEIG \(\overline{H B} O R\) POINTER;
    RELAT̄IONSHIP : GIVEN_IDĒNTIFIERS;
    MINIMAL_DISTANCE : REAL;
    procedure PROCESS_ADJACENT_NODE (BASE_NODE, NEXT_NODE : in INDEX TYPE;
                                    NEXT_BASE_EDGE \(\quad\) : in EDGE_TYPE)
        is separate;
    procedure RESOLVE PATH_TO ENGLISH is separate;
    procedure COMPUTE_COMMON_GENES (INDEX1, INDEX2 : in INDEX_TYPE)
        is separate;
begin -- execution of FIND_RELATIONSHIP
    -- initialize PERSON-array for processing -
    - mark all nodes as not seen
    for PERSON INDEX in 1..NUMBER OF PERSONS loop
        PERSON (PERSON_INDEX) - REACHED_STATUS := NOT_SEEN;
    end loop;
    THIS_NODE := SOURCE_INDEX;
    - mark source node as REACHED
    PERSON (THIS NODE) - REACHED STATUS := REACHED;
    PERSON (THIS_NODE) . DISTANCE_FROM_SOURCE : \(=0.0\);
    -- no NEARBY nodes exist yet
    LAST NEARBY INDEX := 0;
    if THIS NODE \(=\) TARGET INDEX then
        SEARC̄ STATUS := SŪCCEEDED;
    else
        SEARCH STATUS := SEARCHING;
    end if;
```

```
    -- Loop keeps processing closest-to-source, unREACHED node
    -- until target REACHED, or no more connected nodes.
SEARCH FOR TARGET:
    while
        - Process all nodes adjacent to THIS_NODE
        THIS_NEIGHBOR := PERSON (THIS NODE) . NEIGHBOR_LIST_HEADER;
        while THIS NEIGHBOR /= null löop
            PROCESS ADJACENT NODE (THIS NODE,
                                    THIS NEIGHBOR • NEIGHBOR INDEX,
                                    THIS NEIGHBOR . NEIGHBOR EDGE);
        THIS_NEIGHBOR := THIS_NEIGH\overline{BOR . NEXT_NEIGHBOR;}
        end 10op;
        -- All nodes adjacent to THIS NODE are set. Now search for
        - shortest-distance unREACHED (but NEARBY) node to process next.
        if LAST NEARBY INDEX = O then
            SEARC\overline{CH}STAT\overline{US := FAILED;}
        else -- determine next node to process
            MINIMAL DISTANCE := 1.0e+18;
            for PER\overline{SON INDEX in 1..LAST NEARBY INDEX loop}
                if PERSO\overline{N} (NEARBY NODE (PERRSON_INDDEX)) . DISTANCE_FROM_SOURCE
                    < MINIMAL DISTANNCE
            then
                BEST NEARBY INDEX := PERSON INDEX;
                MINIMAL DISTANCE :=
                    PERSŌN (NEARBY NODE (PERSON_INDEX)) . DISTANCE FROM_SOURCE;
                end if;
        end loop;
        -- establish new THIS NODE
        THIS NODE := NEARBY NO
        -- c\overline{7}
        PERSON (THIS NODE) . REACHED STATUS := REACHEN;
        -- remove TH\overline{IS NODE from NEA\overline{R}BY list}
        NEARBY NODE (BEST_NEARBY INDEX) := NEARBY NODE (LAST_NEARBY_INDEX);
        LAST NEARBY INDEX }\mp@subsup{\}{}{-}== LAST NEARBY INDEX - \overline{i}
        if THIS NOD\overline{E}= TARGET INDE\overline{X then}
            SEAR\overline{CH}STATUS := SUUCCEEDED;
        end if;
        end if;
    end loop SEARCH FOR TARGET;
    -- Shortest path between PERSONs now established. Next task is
    -- to translate path to English description of RELATIONSHIP.
    if SEARCH STATUS = FAILED then
        put_liñe (- - & PERSON (TARGET INDEX) . NAME & " is not related to " &
                PERSON (SOURCE_INDEX) . NAME);
    else -- success - parse path to find and display RELATIONSHIP
        RESOLVE PATH TO ENGLISH;
        COMPUTE_COMMON_\overline{GENES (SOURCE_INDEX, TARGET_INDEX);}
    end if;
end FIND_RELATIONSHIP;
```

```
    new compilation-unit #4: procedures under FIND RELATIONSHIP
separate (RELATE . FIND RELATIONSHIP)
procedure PROCESS_ADJACENT_NODE (BASE_NODE, NEXT_NODE : in INDEX TYPE;
                                    NEXT BASE EDGE : in EDGE TYPE) is
    -- NEXT_NODE is adjacent to last-REA
    -- if NEXT NODE already REACHED, do nothing.
    -- If previously seen, check whether path thru BASE_NODE is
    -- shorter than current path to NEXT_NODE, and if so re-link
    - next to base.
    -- If not previously seen, link next to base node.
    WEIGHT_THIS_EDGE, DISTANCE_THRU_BASE_NODE : REAL;
    procedure LINK_NEXT_NODE_TO_BASE_NODE is
        - link next to base by re-setting its predecessor index to
        -- point to base, note type of edge, and re-set distance
        - as it is through base node.
    begin -- execution of LINK_NEXT_NODE_TO BASE_NODE
        PERSON (NEXT NODE) . DISTANCE F\overline{ROM SOUURCE := - DISTANCE THRU BASE_NODE;}
        PERSON (NEXT-NODE) . PATH PREDECES\overline{SOR := BASE NODE ;}
        PERSON (NEXT NODE) . EDGE TO PREDECESSOR := NEXT_BASE EDGE;
    end LINK NEXT N
begin - execution of PROCESS ADJACENT NODE
    if PERSON (NEXT NODE) . REAC\overline{HED STATUS /= REACHED then}
        if NEXT BASE EDGE = SPOUSE then
        WEIG\overline{HT THIS EDGE := 1.8;}
        else
            WEIGHT_THIS_EDGE := 1.0;
        end if;
        DISTANCE THRU BASE NODE := WEIGHT THIS EDGE +
            PERSO\overline{N (BAASE NODDE) . DISTANCE FROM-SOURCE;}
        if PERSON (NEXT NODE) . REACHED STATUS = NOT SEEN then
                PERSON (NEXT NODE) . REACHED STATUS := NEARBY;
                LAST NEARBY ITNDEX := LAST NEARBY INDEX + 1;
                NEAR\overline{BY} NODE (LAST NEARBY I\
                LINK NEXT NODE TO BASE NODDE;
        else =- REACHED STATUS = NEARBY
            if DISTANCE THRU BASE NODE
                    < PERSON}\mp@subsup{}{}{-}(NEXT\_NODE ) . DISTANCE_FROM_SOURCE
                then
                    LINK_NEXT_NODE_TO_BASE_NODE;
                end if;
        end if;
    end if;
end PROCESS_ADJACENT_NODE;
```

separate (RELATE . FIND_RELATIONSHIP)
procedure RESOLVE PATH TO ENGLISH is
-- RESOLVE PATH_TO_ENGLĪSH condenses the shortest path to a
-- series of RELATIONSHIPs for which there are English

- descriptions.
-- Key persons are the ones in the RELATIONSHIP path which remain
-- after the path is condensed.

```
type SIBLING TYPE is (STEP, HALF, FULL);
type KEY PERSON_RECORD (RELATION_TO_NEXT : RELATION_TYPE := PARENT) is
    record
            PERSON INDEX : INDEX TYPE;
            GENERAT}ION_GAP : COUNTER;
            PROXIMITY - : SIBLING_TYPE;
            case RELATION TO NEXT is
                when COUSIN }\mp@subsup{}{}{-}>\mp@subsup{\mathrm{ COUSIN_RANK : COUNTER;}}{}{-
                    when others => null;
            end case;
        end record;
```

-- these variables are used to generate KEY PERSONs
GENERATION COUNT : COUNTER;
THIS COUSIN RANK : COUNTER;
THIS_PROXIMITTY : SIBLING_TYPE;
-- these variables are used to condense the path
KEY_PERSON : array (INDEX_TYPE) of KEY_PERSON_RECORD;
KEY_RELATION, LATER KEY RELATION, PRIMARY_RELATION,
$\bar{N} E X T$ PRIMARY RELATION : RELATION TYPE;
KEY_INDEX, LATER_KEY_INDEX, PRIMARY_INDEX
: INDEX_TYPE;
ANOTHER ELEMENT_POSSIBLE : boolean;
function FULL SIBLING (INDEX1, INDEX2 : in INDEX TYPE)
return boolean is
-- Determines whether two PERSONs are full siblings, i.e.,
- have the same two parents.
begin
return
PERSON (INDEX1) • RELATIVE IDENTIFIER (FATHER IDENT) /= NULL IDENT and
PERSON (INDEX1) • RELATIVE_IDENTIFIER (MOTHER_IDENT) /= NULL_IDENT and
PERSON (INDEXI) - RELATIVE IDENTIFIER (FATHER IDENT) =
PERSON (INDEX2) . RELATIVE IDENTIFIER (FATHER IDENT) and
PERSON (INDEXI) - RELATIVE IDENTTIFIER (MOTHER IDEN̄T) =
PERSON (INDEX2) • RELATIVE_IDENTIFIER (MOTHER_IDENT);
end FULL SIBLING;

```
    procedure CONDENSE KEY PERSONS (AT INDEX : in INDEX TYPE;
                        GAP_SIZE : in COUNTER) is
        -- CONDENSE KEY PERSONS condenses superfluous entries from the
        -- KEY PERSON array, starting at AT_INDEX.
    RECEIVE INDEX, SEND_INDEX : INDEX TYPE;
    begin
        RECEIVE_INDEX := AT_INDEX;
        loop
            RECEIVE INDEX := RECEIVE INDEX + 1;
            SEND INDEX := RECEIVE INDEX + GAP SIZE;
            KEY PERSON (RECEIVE INDE\overline{X}) := KEY PERSON (SEND INDEX);
        exit when KEY_PERSON (SEND_INDEX) - RELATION_TO_\overline{NEXT = NULL_RELATION;}
        end loop;
end CONDENSE KEY PERSONS;
    procedure DISPLAY RELATION (FIRST INDEX, LAST INDEX, PRIMARY_INDEX
                        : in INDEX TYPE)
            is separate;
begin -- execution of RESOLVE PATH TO ENGLISH
    put line (" Shortest path between identified persons: ");
    THIS NODE := TARGET INDEX;
    KEY INDEX := 1;
    -- Display path and initialize KEY PERSON array from path elements.
TRAVERSE SHORTEST PATH:
    whi\overline{1}e THIS NODE /= SOURCE INDEX loop
        put ( - & PERSON (THIS_NODE) . NAME & " is ");
        case PERSON (THIS NODE)-. EDGE TO_PREDECESSOR is
        when PARENT =>
            put_line ("parent of");
            KEY PERSON (KEY_INDEX) :=
                (PERSON INDE\overline{X}}\quad=>\mathrm{ THIS_NODE,
                GENERATION_GAP }\quad>1\mathrm{ ,
                PROXIMITY - }\quad>\mathrm{ FULL,
                RELATION_TO_NEXT => PARENT);
            when CHILD =>
            put_line ("child of");
            KEY PERSON (KEY INDEX) :=
                (PERSON_INDE\overline{X}}\quad=>\mathrm{ THIS_NODE,
                        GENERATION_GAP => 1,
                        PROXIMITY => FULL,
                        RELATION TO_NEXT => CHILD);
            when SPOUSE =>
            put_line ("spouse of");
            KEY_PERSON (KEY INDEX) :=
                    (PERSON_INDE\overline{X}}\quad=>\mathrm{ THIS_NODE,
                        GENERATIION_GAP }\quad>0\mathrm{ ,
                        PROXIMITY - => FULL,
                        RELATION_TO_NEXT => SPOUSE);
        end case;
        KEY INDEX := KEY INDEX + 1;
        THI\overline{S}NODE := PER\overline{SON (THIS NODE) . PATH PREDECESSOR;}
    end loop TRAVERSE_SHORTEST PATH;
```

```
    put_line(` - & PERSON (THIS_NODE) . NAME);
    KEY PERSON (KEY INDEX) :=
    (PERSON_INDE\overline{X}}\quad=>\mathrm{ THIS_NODE,
        GENERATION_GAP }=>\mathrm{ ( 0,
        PROXIMITY }\quad>\mathrm{ FULL,
        RELATION TO NEXT => NULL RELATION);
KEY PERSON (TEY INDEX + 1) :=
    (PERSON INDE\overline{X}}\quad=>0\mathrm{ ,
        GENERAT\ION_GAP => 0,
        PROXIMITY - }\quad>\mathrm{ FULL,
        RELATION TO NEXT => NULL RELATION);
    -- Resolve C\overline{HIL}\overline{D}-PARENT and CHILD-SPOUSE-PARENT relations
    - to SIBLING relations.
    KEY INDEX := 1;
FIND_SIBLINGS:
    while KEY PERSON (KEY INDEX) . RELATION_TO NEXT /= NULL_RELATION loop
        if KEY PERSON (KEY INDEX) . RELATION TO NEXT = CHILD then
            LATER_KEY_RELATİON := KEY PERSON (KEY`_INDEX + 1) . RELATION_TO_NEXT;
            if LATER_\overline{KEY_RELATION = PARRENT then}
                -- found either full or half SIBLINGs
                    if FULL SIBLING (KEY PERSON (KEY INDEX) . PERSON INDEX,
                    KEY_PERSON (KEY_INDEX + 2) . PERSON_INDEX)
                    then
                        THIS PROXIMITY := FULL;
            else
                        THIS PROXIMITY := HALF;
                    end if;
            KEY PERSON (KEY INDEX) :=
                        (PERSON INDE\overline{X}}\quad=>\mathrm{ KEY_PERSON (KEY_INDEX) . PERSON_INDEX,
                        GENERATION GAP => 0,
                        PROXIMITY - }\quad>\mathrm{ THIS_PROXIMITY,
                                RELATION TO NEXT => SIBLİNG);
                    CONDENSE_KEY_PER\
            elsif (LATER KE\overline{Y}}\mathrm{ RELATION = \
                    (KEY PERSON (KEY_INDEX + 2) . RELATION TO_NEXT = PARENT)
            then - found step-SIB\overline{L}INGs
                    KEY PERSON (KEY INDEX) :=
                    (PERSON_INDE\overline{X}}\quad=>\mathrm{ KEY_PERSON (KEY_INDEX) . PERSON_INDEX,
                                    GENERATMION_GAP }=>0\mathrm{ 0,
                                    PROXIMITY - => STEP,
                                    RELATION TO NEXT => SIBLING);
                    CONDENSE KEY PE\overline{RSONS (KEY INDEX, 2);}
            end if; -- LATER_KEY_RELATION = PARENT
        end if; -- RELATION
        KEY_INDEX := KEY_INDE\overline{X}+-
    end 1oop FIND_SIBLİNGS;
```

```
- moolve CHILD-CHILD-... and PARENT-PARENT-... relations to
# Hreet descendant or ancestor relations.
#f% InmX := 1;
@R DESCENDANTS:
    4HE ETERSON (KEY INDEX) - RELATION TO NEXT /= NULL RELATION loop
    4% (%ERSON (KEY_INDEX) . RELATION-TO_NEXT = CHILD) or
    (WY_PeRSON (KEY_INDEX) - RELATION_TO_NEXT = PARENT)
    4mma
        LATER IKEY INDEX := KEY INDEX + 1;
        *1le KEY PERSON (LATER KEY INDEX) . RELATION TO NEXT =
            KEY PERSON (KEY INDEX) . RELATION TO NEXT loop
            LATER K\overline{EY_INDEX := LATER_\overline{KEY_INDEX + 1;}}\mathbf{~}=\mp@code{I}
        men leop;
        _mmAATION COUNT := LATER KEY INDEX - KEY INDEX;
            * EmHERATION COUNT > 1 then - - compress generations
            TM PERSON (KEY INDEX) . GENERATION GAP := GENERATION COUNT;
            CX_酐NSE KEY PERSONS (KEY_INDEX, GENERATION_COUNT - 1);
                if;
            縺 -- if RELATION TO NEXT = CHILD or PARENT
            M变 := KEY INDEX + -1;
            TTHOANCESTORS_OR_DESCENDANTS;
```

```
-- Resolve CHILD-SIBLING-PARENT to COUSIN,
-- CHILD-SIBLING to NEPHEW,
-- SIBLING-PARENT to UNCLE.
```

KEY INDEX : $=1$;
FIND COUSINS NEPHEWS UNCLES:
while KEY PERSON (KEY INDEX) - RELATION TO NEXT /= NULL RELATION 1oop LATER KEY RELATION $\overline{:}=$ KEY PERSON (KEY INDEX + 1) - RELATION TO NEXT; if (KĒY PERSON (KEY INDEX) . RELATION TO NEXT = CHILD) and (LATER KEY RELATĪON = SIBLING)
then -- COUSIN or NEPHEW
if KEY PERSON (KEY INDEX + 2) . RELATION TO NEXT = PARENT then - found COUSIN
if KEY PERSON (KEY INDEX) - GENERATION GAP < KEY_PERSON (KEY_INDEX + 2) • GENERATION_GAP
then
THIS COUSIN RANK : =
$\bar{K} E Y$ PER $\bar{S} O N$ (KEY_INDEX) • GENERATION_GAP;
else
THIS COUSIN RANK :=
$\bar{K} E Y$ PERS $\bar{S} O N$ (KEY_INDEX + 2) . GENERATION_GAP;
end if;
KEY PERSON (KEY INDEX) : =
(PERSON INDE $\bar{X} \quad \Rightarrow$ KEY_PERSON (KEY INDEX) • PERSON INDEX, GENERATION GAP =>
abs (KEY PERSON (KEY INDEX) . GENERATION GAP KEY_PERSON (KEY_INDEX + 2) - GENERATION_GAP), PROXIMITY $\quad \Rightarrow$ KEY PERSON (KEY INDEX + 1) • PROXXIMITY, RELATION TO NEXT $\Rightarrow$ COUSIN, COUSIN_RANK $\quad \Rightarrow$ THIS_COUSIN_RANK); CONDENSE KEY PERSONS (KEY_INDEX, 2); else -- found $\bar{N} E P H E W$ KEY PERSON (KEY INDEX) : = (PERSON INDE $\bar{X} \quad \Rightarrow$ KEY PERSON (KEY INDEX) . PERSON INDEX, GENERATION GAP $\Rightarrow \mathrm{KEY}^{-}$PERSON (KEY ${ }^{-}$INDEX) • GENERATION GAP, PROXIMITY - $\quad \Rightarrow$ KEY_PERSON (KEY_INDEX + 1) • PROXIMITY, RELATION TO NEXT $\Rightarrow$ NEP $\overline{H E W}$ ); CONDENSE KEY_PERSONS (KEY_INDEX, 1);
end if;
elsif KEY PERSON (KEY INDEX) • RELATION_TO_NEXT = SIBLING and LATER KEY RELATION = PARENT
then -- found UNCLE
KEY PERSON (KEY_INDEX) : =
(PERSON INDE $\bar{X} \quad \Rightarrow$ KEY PERSON (KEY INDEX) • PERSON INDEX, GENERATION GAP $\Rightarrow \mathrm{KEY}^{-}$PERSON (KEY INDEX + 1) . GENEERATION GAP, PROXIMITY - $\quad \Rightarrow$ KEY PERSON (KEY INDEX) . PROXIMITY, RELATION TO NEXT $\Rightarrow$ UNC $\overline{L E}$ ); CONDENSE KEY_PERSONS (KEY INDEX, 1);
end if;
KEY INDEX := KEY INDEX + 1;
end 100p FIND_COUSINS_NEPHEWS_UNCLES;
-- Loop below will pick out valid adjacent strings of elements

- to be displayed. KEY_INDEX points to first element,
-- LATER_KEY_INDEX to las̃t element, and PRIMARY INDEX to the
- element which determines the primary English word to be used.
-- Associativity of adjacent elements in condensed table
- is based on English usage.

KEY_INDEX := 1;
put line (" Condensed path:");
CONSOLIDATE ADJACENT PERSONS:
while $\bar{K} E Y$ PERSON (KEY INDEX) - RELATION TO NEXT /= NULL RELATION $100 p$
KEY RELATION $:=\overline{\mathrm{KEY}}$ PERSON (KEY_INDEX $\overline{\text { ) }}$. RELATION_TO_NEXT;
LATER_KEY_INDEX := KEY_INDEX;
PRIMA $\bar{R} Y$ IÑDEX $:=K^{\prime} Y^{-}$INDEX;
if KEY PERSON (KEY INDEX + 1) . RELATION TO NEXT /= NULL_RELATION then

- $\overline{s e e k}$ multi-e $\overline{1} e m e n t ~ c o m b i n a t i o n ~$

ANOTHER ELEMENT POSSIBLE := true;
if KEY RELATION $=$ SPOUSE then LATER KEY INDEX := LATER_KEY_INDEX + 1; PRIMARY $\overline{\mathrm{R}}$ INDEX $:=$ LATER $^{-}$KEY ${ }^{-}$INDEX; if (KEY_PERSON (LATER_KE $\bar{Y}$ INDEX) - RELATION_TO_NEXT = SIBLING) or (KEY_PERSON (LATER_KEY_INDEX) • RELATION_TO_NEXT = COUSIN) then $=-$ Nothing can folīow SPOUSE-SIbling or ${ }^{-}$SPOUSE-COUSIN ANOTHER_ELEMENT_POSSIBLE := false;
end if;
end if;
-- PRIMARY INDEX is now correctly set. Next if-statement

- determines if a following SPOUSE relation should be
-- appended to this combination or left for the next
- combination.
if ANOTHER ELEMENT POSSIBLE and
(KEY_PERSON (PRTMMARY_INDEX + 1) • RELATION_TO_NEXT = SPOUSE) - Only a SPOUSE can follow a Primary
then
-- check primary preceding and following SPOUSE.
PRIMARY RELATION :=
KEY_PERSON (PRIMARY_INDEX) . RELATION_TO_NEXT;
NEXT Pर्RIMARY RELATION $\overline{:}=$
KĒY PERSON (PRIMARY INDEX + 2) . RELATION TO NEXT;
if (NEXT PRIMARY RELATİON = NE PHEW or
NEXT_PRTMARY_RELATION = COUSIN or
NEXT PRIMARY RELATION $=$ NULL RELATION)
or ( $\operatorname{P\overline {R}}$ IMARY RELATION $=$ NEPHEW $)$
or ( (PRIMA $\bar{R} Y$ RELATION $=$ SIBLING or
PRIMARY ${ }^{-}$RELATION $=$PARENT)
and NEXT_PRIMARY_RELATION /= UNCLE )
then -- append following SPOUSE with this combination. LATER_KEY_INDEX := LATER_KEY_INDEX + 1; end if;
end if;
end if; -- multi-element combination
DISPLAY RELATION (KEY_INDEX, LATER_KEY_INDEX, PRIMARY_INDEX);
KEY_INDEX := LATER_KEY_INDEX + 1 ;
end 10̄op CONSOLIDATE ${ }^{-}$ADJĀCENT PERSONS;
put_line ( ${ }^{-}$- \& PERSŌN (KEY PERSSON (KEY_INDEX) . PERSON_INDEX) . NAME);
end; -- RESOLVE_PATH_TO_ENGLISH

```
--- new compilation-unit #5: procedures under RESOLVE_PATH_TO_ENGLISH
separate (RELATE . FIND_RELATIONSHIP . RESOLVE PATH_TO_ENGLISH)
procedure DISPLAY_RELATION (FIRST INDEX, LAST INDEX, PRIMARY_INDEX
    : in INDEX TYPE) is
    -- DISPLAY RELATION takes 1, 2, or \overline{3}}\mathrm{ adjacent elements in the
    -- condensed table and generates the English description of
    -- the relation between the first and last + l elements.
    INLAW : boolean;
    THIS PROXIMITY : SIBLING TYPE;
    THIS GENDER : GENDER TYPE;
    FIRST RELATION, LAST RELATION, PRIMARY RELATION
                            : RELATION TYPE;
THIS_GENERATION GAP, THIS_COUTSIN_RANK
                    : COUNTER;
    -- need to instantiate package to display integer values
    package COUNTER_IO is
        new integer_io (COUNTER);
```

```
begin -- execution of DISPLAY RELATION
    FIRST RELATION := KEY PERSON (FIRST INDEX) . RELATION TO_NEXT;
    LAST_\overline{RELATION := KEY_PERSON (LAST_INDEX) . RELATIONTO_NEXT;}
    PRIMARRY_RELATION := KEY_PERSON (PRIMARY_INDEX) . RELATION_TO_NEXT;
    -- set THIS PROXIMITY
    if ((PRIMAR\overline{Y}_RELATION = PARENT) and (FIRST RELATION = SPOUSE)) or
        ((PRIMARY_RELATION = CHILD) and (LAST_\overline{RELATION = SPOUSE))}
    then
        THIS PROXIMITY := STEP;
    elsif P\overline{RIMARY RELATION = SIBLING or}
            PRIMARY RELATION = UNCLE or
            PRIMARY RELATION = NEPHEW or
            PRIMARY_RELATION = COUSIN
    then
        THIS_PROXIMITY := KEY_PERSON (PRIMARY_INDEX) . PROXIMITY;
    else
        THIS_PROXIMITY := FULL;
    end if;
    -- set THIS GENERATION GAP
    if PRIMARY \overline{RELATION = \overline{PARENT or}}\mathbf{T}\mathrm{ - }
        PRIMARY RELATION = CHILD or
        PRIMARY RELATION = UNCLE or
        PRIMARY RELATION = NEPHEW or
        PRIMARY - RELATION = COUSIN
    then
        THIS GENERATION GAP := KEY PERSON (PRIMARY_INDEX) . GENERATION_GAP;
    else
        THIS GENERATION_GAP := 0;
    end if;
    -- set INLAW
    INLAW := false;
    if (FIRST_RELATION = SPOUSE) and
        (PRIMA\overline{RY}_RELATION = SIBLING or
        PRIMARY RELATION = CHILD or
        PRIMARY RELATION = NEPHEW or
        PRIMARY_RELATION = COUSIN)
    then
        INLAW := true;
    elsif (LAST_RELATION = SPOUSE) and
            (PRIMARY RELATION = SIBLING or
            PRIMARY RELATION = PARENT or
            PRIMARY RELATION = UNCLE or
            PRIMARY_RELATION = COUSIN)
    then
        INLAW := true;
    end if;
    -- set THIS COUSIN RANK
    if PRIMARY \overline{RELATION}=\mathrm{ COUSIN then}
        THIS COUSIN RANK := KEY PERSON (PRIMARY INDEX) . COUSIN RANK;
    end if;
```

```
- parameters are set - now generate display.
put (" " & PERSON (KEY_PERSON (FIRST_INDEX) • PERSON_INDEX) • NAME &
    " is ");
if PRIMARY RELATION = PARENT or
    PRIMARY RELATION = CHILD or
    PRIMARY RELATION = UNCLE or
    PRIMARY RELATION = NEPHEW
then
    -- display generation-qualifier
    if THIS GENERATION GAP >= 3 then
        put ("great");
        if THIS GENERATION_GAP > 3 then
            put ("*");
            COUNTER_IO . put (THIS_GENERATION_GAP - 2, width => 1);
            end if;
            put ("-");
    end if;
    if THIS GENERATION GAP >= 2 then
        put ("grand-");
        end if;
elsif (PRIMARY RELATION = COUSIN) and then (THIS COUSIN RANK > 1) then
    COUNTER IO - put (THIS COUSIN RANK, width => \overline{1});
    case THIS_COUSIN RANK mod 10 is
        when 1 => put ("st ");
        when 2 }\quad>\mathrm{ put ("nd ");
        when 3 }\quad>\mathrm{ put ("rd ");
        when others => put ("th ");
    end case;
end if;
if THIS PROXIMITY = STEP then
    put ("step-");
elsif THIS PROXIMITY = HALF then
    put ("hālf-");
end if;
```

```
    THIS_GENDER := PERSON (KEY_PERSON (FIRST INDEX) . PERSON_INDEX) . GENDER;
    case PRIMARY RELATION is
        when PARENT }=>>\mathrm{ if THIS_GENDER = MALE then put ("father");
                        else put ("mother");
                            end if;
        when CHILD => if THIS GENDER = MALE then put ("son");
                        else put ("daughter");
                                end if;
        when SPOUSE => if THIS GENDER = MALE then put ("husband");
                        else put ("wife");
                            end if;
        when SIBLING => if THIS GENDER = MALE then put ("brother");
                        else put ("sister");
                            end if;
        when UNCLE => if THIS_GENDER = MALE then put ("uncle");
                        else
                            put ("aunt");
                            end if;
        when NEPHEW => if THIS GENDER = MALE then put ("nephew");
                        else put ("niece");
                        end if;
        when COUSIN => put ("cousin");
        when others => put ("null");
    end case;
    if INLAW then
    put ("-in-1aw");
    end if;
    if (PRIMARY RELATION = COUSIN) and (THIS_GENERATION_GAP > 0) then
        if THIS \overline{GENERATION GAP > 1 then}
            put (" ");
            COUNTER IO . put (THIS GENERATION GAP, width => 1);
            put (" Eimes removed");
        else
            put (" once removed");
        end if;
end if;
put_line (" of");
end DISPLAY RELATION;
```

```
--- new compilation-unit #6: procedures under FIND RELATIONSHIP
separate (RELATE . FIND RELATIONSHIP)
procedure COMPUTE COMMON GENES (INDEX1, INDEX2 : in INDEX TYPE) is
    -- COMPUTE_COMMON GENES assumes that each ancestor contributes
    - half of the genetic material to a PERSON. It finds common
    -- ancestors between two PERSONs and computes the expected
    - value of the PROPORTION of common material.
    COMMON PROPORTION : REAL;
    package REAL IO is
        new FLOAT \overline{IO (REAL);}
    procedure ZERO PROPORTION (ZERO INDEX : in INDEX TYPE) is
    -- ZERO PROPORTION recursively seeks out all ancestors and
    -- zeros them out.
    THIS_NEIGHBOR : NEIGHBOR_POINTER;
    begin
        PERSON (ZERO_INDEX) . DESCENDANT GENES := 0.0;
        THIS NEIGHBO\overline{R}}:= PERSON (ZERO INDEX) . NEIGHBOR LIST_HEADER
        while THIS NEIGHBOR /= null loop
            if THIS NEIGHBOR . NEIGHBOR EDGE = PARENT then
                ZERO PROPORTION (THIS NEİGHBGR . NEIGHBOR INDEX);
            end if;
            THIS_NEIGHBOR := THIS_NEIGHBOR . NEXT_NEIGHBOR;
        end loop;
    end ZERO PROPORTION;
    procedure MARK_PROPORTION (MARKER : in IDENTIFIER_TYPE;
                                    PROPORTION : in REAL;
                        MARKED INDEX : in INDEX TYPE) is
        -- MARK PROPORTION recursively seeks out all ancestors and
        -- marks them with the sender's PROPORTION of shared
        - genetic material. This PROPORTION is diluted by one-half
        -- for each generation.
    THIS_NEIGHBOR : NEIGHBOR_POINTER;
    begin
    PERSON (MARKED_INDEX) . DESCENDANT_IDENTIFIER := MARKER;
    PERSON (MARKED INDEX) . DESCENDANT GENES :=
            PERSON (MAR\overline{RED INDEX) . DESCENDĀNT GENES + PROPORTION;}
        THIS NEIGHBOR := \overline{PERSON (MARKED INDEX) . NEIGHBOR LIST HEADER;}
        whil\overline{e}
            if THIS NEIGHBOR . NEIGHBOR EDGE = PARENT then
                MARK PROPORTION (MARKER, PROPORTION / 2.0,
                                    THIS NEIGHBOR . NEIGHBOR INDEX);
            end if;
            THIS_NEIGHBOR := THIS_NEIGHBOR . NEXT_NEIGHBOR;
        end loop;
    end MARK_PROPORTION;
```

procedure CHECK COMMON PROPORTION
(COMMON PROPORTION : in out REAL; MATCH_IDENTIFIER : in IDENTIFIER_TYPE; PROPORTION : in REAL; ALREADY COUNTED : in REAL; CHECK IÑDEX : in INDEX TYPE) is
-- CHECK COMMON PROPORTION searches all the ancestors of
-- CHECK_INDEX to see if any have been marked, and if so -- adds the appropriate amount to COMMON PROPORTION.

THIS NEIGHBOR : NEIGHBOR_POINTER;
THIS_CONTRIBUTION : REAL;
begin
if PERSON (CHECK_INDEX) . DESCENDANT_IDENTIFIER = MATCH_IDENTIFIER then
-- Increment $\bar{C} O M M O N$ PROPORTION by the contribution of
-- this common ancestor, but discount for the contribution

- of less remote ancestors already counted.

THIS_CONTRIBUTION := PERSON (CHECK_INDEX) . DESCENDANT_GENES

* PROPORTION;

COMMON PROPORTION := COMMON PROPORTION

+ TH̄IS_CONTRIBUTION - AL $\bar{R} E A D Y$ COUNTED;
else
THIS CONTRIBUTION := 0.0;
end if;
THIS NEIGHBOR := PERSON (CHECK_INDEX) . NEIGHBOR_LIST_HEADER; while THIS NEIGHBOR /= null loop
if THIS NEIGHBOR . NEIGHBOR EDGE = PARENT then CHECK_COMMON PROPORTION (COMMON_PROPORTION, MATCH_ĪDENTIFIER, PROPORTĪON / 2.0,
THIS_C̄ONTRIBUTION / 4.0,
THIS_NEIGHBOR • NEIGHBOR_INDEX);
end if;
THIS_NEIGHBOR := THIS_NEIGHBOR . NEXT_NEIGHBOR;
end loop;
end CHECK COMMON PROPORTION;
begin -- COMPUTE_COMMON_GENES
-- First zero out all ancestors to allow adding. This is necessary
- because there might be two paths to an antestor.

ZERO PROPORTION (INDEX1);

- now mark with shared PROPORTION

MARK_PROPORTION (PERSON (INDEX1) . IDENTIFIER, 1.0, INDEX1);
COMMŌN PROPORTION := 0.0;
CHECK C̄OMMON PROPORTION (COMMON_PROPORTION,
PER̄SON (INDEX1) . IDENTIFIER, 1.0, 0.0, INDEX2);
put (" Proportion of common genetic material = ");
REAL IO . put (COMMON PROPORTION, fore $\Rightarrow 1$, aft $\Rightarrow 5$, $\exp \Rightarrow 3$ );
put line (" ");
end CŌMPUTE_COMMON_GENES;

### 3.0 BASIC

Because of the unavailability of a standard implementation, the BASIC program could not be tested directly. However, a syntactically non-standard version, which is believed to be logically equivalent, was tested.

```
10000 ! --- program-unit number 1 ----
10010 !
10020 program RELATE
10030 !
10040 ! declare subs to be used by this program-unit
10050 !
10060 declare external sub FIND RELATIONSHIP
10070 declare sub LINK RELATIVE\overline{S}, LINK ONE WAY, PROMPT AND READ
10080 declare sub CHEC\overline{K}_REQUEST, SEARC\overline{H_FOR_REQUESTED_ PERSONS}
10090 !
10100 option base 1
10110 !
10120 ! Define global objects
10130 !
10140 data 300
10150 read MAX PERSONS
10160 !
10170 data 1, 2 ! for truth values
10180 read TRUE, FALSE
10190 !
10200 ! each PERSON's record in the file identifies at most three
10210 ! others directly related: father, mother, and spouse
10220 data 1, 2, 3
10230 read FATHER_IDENT, MOTHER_IDENT, SPOUSE_IDENT
10240 !
10250 data M, F
10260 read MALE$, FEMALE $
10270 !
10280 data 000
10290 read NULL_IDENT$
10300 !
10310 data 1, 2, 3, 4, 5, 6, 7, 8
10320 read PARENT, CHILD, SPOUSE, SIBLING, UNCLE, NEPHEW
10325 read COUSIN, NULL_RELATION
10330 !
10340 ! A node in the graph (= PERSON) has either already been reached,
10350 ! is immediately adjacent to those reached, or farther away.
10360 data 1, 2, 3
10370 read REACHED, NEARBY, NOT SEEN
10380 !
```

```
10390 ! The following data arrays are the central repository of information
10400 ! .about inter-relationships. All relationships are captured in the
10410 ! directed graph of which each record is a node.
10420 !
10430 ! static information - filled from PEOPLE file:
10440 dim NAME$ (300), IDENTIFIER$ (300), GENDER$ (300)
10450 !
10460 ! IDENTIFIER$s of immediate relatives - father, mother, spouse
10470 dim RELATIVE_IDENTIFIER$ (300,3)
10480 !
10490 ! pointers to immediate neighbors in graph
10500 dim NEIGHBOR_COUNT (300)
10505 dim NEIGHBOR_INDEX (300,20), NEIGHBOR_EDGE (300, 20)
10510 !
10520 ! data used when traversing graph to resolve user request:
10530 dim DISTANCE FROM SOURCE (300), PATH PREDECESSOR (300)
10540 dim EDGE TO \overline{PREDECESSSOR (300), REACHED STATUS (300)}
10550 !
10560 ! data used to compute common genetic material
10570 dim DESCENDANT_IDENTIFIER$ (300), DESCENDANT_GENES (300)
10580 !
10590 data stop, Request OK
10600 read REQUEST_TO_STOP$, REQUEST_OK$
10610 !
10620 ! end initialization
10630 !
```

10640
10650
10660 \& 10670 !
10680 ! This loop reads in the PEOPLE file and constructs the person 10690 ! array from it (one person = one set of array entries).
10700 ! As records are read in, links are constructed to represent the
10710 ! PARENT-CHILD or SPOUSE RELATIONSHIP. The array then implements
10720 ! a directed graph which is used to satisfy subsequent user
10730 ! requests. The file is assumed to be correct - no validation
10740 ! is performed on it.
10750 !
10760 for CURRENT $=1$ to MAX PERSONS
read \#1, if missing then exit for, \& with "string*20, string*3, string*1, 3 of string*3": \&
NAME $\$(C U R R E N T), ~ I D E N T I F I E R \$ ~(C U R R E N T), ~ G E N D E R \$ ~(C U R R E N T), ~ \& ~$
RELATIVE_IDENTIFIER\$ (CURRENT, FATHER_IDENT), \&
RELATIVE - IDENTIFIER\$ (CURRENT, MOTHER -IDENT), \&
RELATIVE ${ }^{-}$IDENTIFIER\$ (CURRENT, SPOUSE ${ }^{-}$IDENT)
let NAME ( $\bar{C} U R R E N T)=$ rtrim (NAME (CUR $\bar{R} E N T)$ )
! Location of adjacent persons as yet undetermined
let NEIGHBOR_COUNT (CURRENT) $=0$
! Descendants as yet undetermined
let DESCENDANT IDENTIFIER\$ (CURRENT) = NULL IDENT $\$$
let CURRENT_IDENT\$ = IDENTIFIER\$ (CURRENT)
! Compare this PERSON against all previously entered PERSONs
! to search for RELATIONSHIPs.
for PREVIOUS = 1 to CURRENT - 1
let PREVIOUS_IDENT\$ = IDENTIFIER\$ (PREVIOUS)
! Search for father, mother, or spouse relationship in
! either direction between this and PREVIOUS person.
! Assume at most one RELATIONSHIP exists.
for RELATIONSHIP = FATHER IDENT to SPOUSE IDENT
if RELATIVE IDENTIFIER $\overline{\$}$ (CURRENT, RELAT$I O N S H I P) ~ \& ~$
$=$ PREV̄IOUS IDENT\$ then
call LINK_RELĀTIVES (CURRENT, RELATIONSHIP, PREVIOUS) exit for
elseif RELATIVE_IDENTIFIER\$ (PREVIOUS, RELATIONSHIP) \& = CURRENT IDENT\$ then call LINK_RELATIVES (PREVIOUS, RELATIONSHIP, CURRENT) exit for
end if
next RELATIONSHIP
next PREVIOUS
(品

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11080 ! Do-loop accepts requests and finds relationship (if any)
11090 ! between pairs of PERSONs.
11110 do
11120 call PROMPT AND READ
11130 if REQUEST BUFFERS = REQUEST TO STOP\$ then exit do
11140 call CHECK_REQUEST (ERROR_MESSAGE\$, PERSON1_IDENT\$, PERSON2_IDENT\$)
11150 !
11160 ! Syntax check of request completed. Now either display error
11170 ! message or search for the two PERSONs.
11180 !
11190 if ERROR MESSAGE $\$=$ REQUEST_OK\$ then
11200 ! request syntactically correct
11210 call SEARCH FOR REQUESTED PERSONS (PERSON1 IDENT\$, PERSON2 IDENT\$, \&
\&
\&
11220
11230
11240
11250
11260
11270
11280
11290
11300
11310
11320
11330
\&
\&
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\&
\&
11340
11350
11360
11370
11380
11390
11400
11410
11420
11430
11440
11450
11460
11470
11480
11490
11500
11510
11520 1oop
11530 print " End of relation-finder."
11540 stop
11550 !
11560 ! end of main line of execution; internal subs follow

```
11570 !
11580 sub LINK RELATIVES (FROM INDEX, RELATIONSHIP, TO INDEX)
11590 ! esta\overline{b}lishes cross-in\overline{d}exing between immediate\overline{ly related PERSONs.}
11600 !
11610 if RELATIONSHIP = SPOUSE IDENT then
11620 call LINK ONE WAY (FROMM INDEX, SPOUSE, TO INDEX)
11630 cal1 LINK_ONE WAY (TO_INDEX, SPOUSE, FROM_INDEX)
11640 else ! RELA
11650 call LINK ONE WAY (FROM INDEX, PARENT, TO INDEX)
11660 call LINK_ONE WAY (TO_INDEX, CHILD, FROM INDEX)
11670 end if
11680 end sub
11690 !
11700 sub LINK ONE WAY (FROM INDEX, THIS EDGE, TO INDEX)
11710 ! Esta\overline{b}lis\overline{hes the ne\overline{ighbor entries from one person to another}}\mathbf{|}=\overline{M}
11720 !
11730 let NEXT NEIGHBOR = NEIGHBOR COUNT (FROM INDEX) + 1
11740 let NEIG\overline{HBOR COUNT (FROM INDEXX) = NEXT NEIGHBOR}
11750 let NEIGHBOR INDEX (FROM INDEX, NEXT NEIGHBOR) = TO INDEX
11760 let NEIGHBOR_EDGE (FROM_INDEX, NEXT_NEIGHBOR) = THIS_EDGE
11770 end sub
11780 !
11790 sub PROMPT AND READ
11800 ! Issues prompt for user-request, reads in request,
11810 ! blank-fills buffer, and skips to next line of input.
11820 !
11830 print
11840 print
11850 print " Enter two person-identifiers (name or number),"
11860 print " separated by semicolon. Enter ""stop"" to stop."
11870 1ine input REQUEST BUFFER$
11880 end sub
11890 !
11900 sub CHECK REQUEST (REQUEST STATUS$, PERSON1 IDENT$, PERSON2 IDENT$)
11910 ! Performs syntactic check on request in buffer
11920 ! and fills in identifiers of the two requested persons.
11930 !
11940 let SEMICOLON LOCATION = pos (REQUEST_BUFFER$, ";")
11950 let PERSON1_IDENTS = 1trim$ (rtrim$ -&
& (REQUEST
11960 let PERSON2 IDENTS = 1trim$ (rtrim$ &
& (REQUEST BUFFER$ (SEMICOLON LOCATION + 1 : len (REQUEST BUFFERS))))
11970 if SEMICOLON LOCATION = 0 or pOs (PERSON2 IDENT$, ";") <> \overline{0}}\mathrm{ then
11980 let REQUEST STATUS$ = "must be exactly one semicolon."
11990 elseif PERSON1 IDENT$ = "" then
12000 let REQUEST STATUS$ = "null field preceding semicolon."
12010 e1seif PERSON2 IDENT$ = "" then
12020 let REQUEST_STATUS$ = "null field following semicolon."
12030 else
12040 let REQUEST_STATUS$ = REQUEST_OK$
12050 end if
12060 end sub
12070 !
```

```
12080 sub SEARCH_FOR_REQUESTED_PERSONS (PERSON1_IDENT$, PERSON2_IDENT$, &
& - PERSON1-INDEX, PERSON2 INDEX, &
& PERSON1-FOUND, PERSON2 FOUND)
12090 ! SEARCH FOR REQUESTED PERSONS scans through the PERSON array,
12100 ! looking for the two requested PERSONs. Match may be by NAME
12110 ! or unique IDENTIFIER-number
12120 !
12130 let PERSON1 FOUND = 0
12140 let PERSON2 FOUND = 0
12150 let PERSON1 INDEX = 0
12160 let PERSON2_INDEX = 0
12170 for CURRENT = 1 to NUMBER_OF_PERSONS
12180 ! allow identification
12190 if IDENTIFIER$ (CURRENT) = PERSON1 IDENT$ &
&
12200
12210 let PERSON1_FOUND = PERSON1 FOUND + 1
12220 end if
12230 if IDENTIFIER$ (CURRENT) = PERSON2 IDENT$ &
& or NAME$ (CURRENT) = PERSON2_IDENT$ then
12240 let PERSON2 INDEX = CURRENT
12250 let PERSON2_FOUND = PERSON2_FOUND + 1
12260 end if
12270 next CURRENT
12280 end sub
12290 end ! of main program unit - external procedures follow
12300 !
```

```
12310 ! --- program-unit number 2 ----
12320 !
12330 external sub FIND RELATIONSHIP &
& (TARGET INDEX, SOURCE INDEX, NUMBER OF PERSONS, &
& NAME$ (), IDENTIFIER\ (), GENDER$ (), RELATIVE IDENTIFIER$ (,), &
& NEIGHBOR COUNT (), NEIGHBOR INDEX (,), NEIGHBOR EDGE (,), &
& DISTANCE FROM SOURCE (), PATH PREDECESSOR (), - &
& EDGE TO PREDECESSOR (), REACHED STATUS (),
& DESCENDANT IDENTIFIER$ (), DESCENDANT GENES ())
12340 !
12350 !
12360 ! determines their RELATIONSHIP based on immediate relations
12370 ! traversed in path. PERSON array simulates a directed graph,
12380 ! and algorithm finds shortest path, based on following
12390 ! weights: PARENT-CHILD edge = 1.0
12400 ! SPOUSE-SPOUSE edge = 1.8
12410
12420 ! declare subs and functions to be used by this program-unit
12430 !
12440 declare external sub COMPUTE COMMON GENES
12450 declare sub PROCESS_ADJACENT NODE, LINK NEXT NODE_TO BASE_NODE
12460 declare sub RESOLVE PATH TO ENGLISH, CONDENSE KEY PE F
12465 declare sub DISPLAY RELATION
12470 declare function SI\overline{BLING PROXIMITY}
12480 !
12483 option base l
12487 !
12490 ! Define global objects
12500 !
12510 data 300
12520 read MAX PERSONS
12530 !
12540 data 1, 2 ! for truth values
12550 read TRUE, FALSE
12560
12570 ! each PERSON`s record in the file identifies at most three
12580 ! others directly related: father, mother, and spouse
12590 data 1, 2, 3
12600 read FATHER IDENT, MOTHER IDENT, SPOUSE IDENT
12610 !
12620 data M, F
12630 read MALE$, FEMALE $
12640 !
1 2 6 5 0 \text { data 000}
12660 read NULL_IDENT$
12670 !
12680 data 1, 2, 3, 4, 5, 6, 7, 8
1 2 6 9 0 \text { read PARENT, CHILD, SPOUSE, SIBLING, UNCLE, NE PHEW}
12695 read COUSIN, NULL RELATION
12700 !
12710 ! A node in the graph (= PERSON) has either already been reached,
12720 ! is immediately adjacent to those reached, or farther away.
12730 data 1, 2, 3
12740 read REACHED, NEARBY, NOT SEEN
12750 !
```

```
12760 data 1, 2, 3 ! values for search status
12770 read SEARCHING, SUCCEEDED, FAILED
12780 !
12790 data 1, 2, 3 ! values for sibling proximity
12800 read STEP, HALF, FULL
12810 !
12820 ! The following arrays contain information on key persons.
12830 ! Key persons are the ones in the RELATIONSHIP path which remain
12840 ! after the path is condensed.
12850 !
12860 dim RELATION TO NEXT (300), PERSON INDEX (300), GENERATION GAP (300)
12870 dim PROXIMIT\overline{Y}}(\overline{3}00), COUSIN RANK (\overline{300}
12880 !
12890 ! keeps track of current NEARBY nodes in graph search
12900 dim NEARBY_NODE (300)
12910 !
12920 ! begin main line of execution of FIND RELATIONSHIP
12930 !
12940 ! initialize PERSON-array for processing -
12950 ! mark all nodes as not seen
12960 for THIS NODE = 1 to NUMBER OF PERSONS
12970 let REACHED_STATUS (THIS_NODE) = NOT_SEEN
12980 next THIS NODE
12990 !
13000 let THIS NODE = SOURCE INDEX
13010 ! mar\overline{k}}\mathrm{ source node as REACHED
13020 let REACHED_STATUS (THIS_NODE) = REACHED
13030 let DISTANCE_FROM_SOURCE (THIS_NODE) = 0
13040 ! no near\overline{b}y no\overline{des exist yet}\mp@subsup{}{}{-}
13050 let LAST NEARBY INDEX = 0
13060 if THIS N
13070 let SEARCH STATUS = SUCCEEDED
13080 else
13090 let SEARCH STATUS = SEARCHING
13100 end if
13110 !
```



```
13570 sub PROCESS ADJACENT NODE (BASE NODE, NEXT NODE, NEXT BASE EDGE)
13580 ! NEXT NODE is ad \(\bar{j}\) acent to lāst-REACHED node (= BASE NODE).
13590 ! if NEXT NODE already REACHED, do nothing.
13600 ! If previously seen, check whether path thru BASE NODE is
13610 ! shorter than current path to NEXT NODE, and if so re-link
13620 ! next to base.
13630 ! If not previously seen, link next to base node.
13640 !
13650 if NEXT BASE EDGE \(=\) SPOUSE then
13660 let \(\overline{\mathrm{W} E I G H T}\) _THIS_EDGE \(=1.8\)
13670 else
13680 let WEIGHT_THIS_EDGE \(=1.0\)
13690 end if
13700 !
13710 if REACHED STATUS (NEXT NODE) 〈〉 REACHED then
13720 let DISTANCE THRU BASE NODE
\& \(\quad=\) WEIGHT \(^{-}\)THIS_EDGE \({ }^{-}+\)DISTANCE FROM SOURCE (BASE NODE)
13760 let LAST NEARBY INDEX = LAST NEARBY INDEX + 1
13770 let NEAR \(\bar{B} Y\) NODE (LAST NEARBY_INDEX) \(=\) NEXT NODE
13780 ! link next to base by re-setting its predecessor index to
13790 ! point to base, note type of edge, and re-set distance
13800 ! as it is through base node.
13810 let DISTANCE FROM SOURCE (NEXT NODE) = DISTANCE THRU BASE NODE
13820 let PATH PREDECESSOR (NEXT NODE) = BASE NODE
13830 let EDGE \({ }^{-}\)TO PREDECESSOR (NEXT \({ }^{-}\)NODE) \(=\)NEXT \({ }^{-}\)BASE EDGE
13840 else ! \(\overline{R E A C} H E D ~ S T A T U S ~=~ N E A R B \bar{Y}\)
13850 if DISTANCE THRU BASE NODE < DISTANCE FROM SOURCE (NEXT NODE) then
13860 ! link nex \(\bar{t}\) to \(\bar{b}\) ase by re-setting it \(\bar{s}\) predecessor index to
13870 ! point to base, note type of edge, and re-set distance
13880 ! as it is through base node.
13890 let DISTANCE FROM SOURCE (NEXT_NODE) = DISTANCE THRU_BASE_NODE
13900 let PATH PREDECESSOR (NEXT NODE) = BASE NODE
13910 let EDGE TO PREDECESSOR (NEXT NODE) = NEXT BASE EDGE
13920 end if
13930 end if
13940 end if
13950 end sub
13960 !
```

```
13970 sub RESOLVE_PATH_TO_ENGLISH
13980 ! RESOLVE PAT\overline{H}T\overline{O}}\mathrm{ ENGLISH condenses the shortest path to a
13990 ! series of RELAT\overline{I}ONSHIPs for which there are English
14000 ! descriptions.
14010 !
14020 ! Key persons are the ones in the RELATIONSHIP path which remain
14030 ! after the path is condensed.
14040 !
14050 print " Shortest path between identified persons: "
14060 let THIS NODE = TARGET INDEX
14070 ! priñt path and in\overline{i}tialize KEY_PERSON array from path elements,
14080 ! as shortest path is traversed.
14090 let KEY INDEX = 1
14100 do unti\overline{1}}\mathrm{ THIS NODE = SOURCE INDEX
14110 let PERSON_INDEX (KE\overline{Y}_INDEX) = THIS_NODE
14120 let PROXIMITTY (KEY_INDEX) = FULL
14130 let RELATION_TO_NEXT (KEY_INDEX) = EDGE_TO_PREDECESSOR (THIS_NODE)
14140 print " "; NAMME\ (THIS NODE); tab(23); "is '";
14150 if EDGE TO PREDECESSOR (THIS NODE) = SPOUSE then
14160 let GENERATION_GAP (KEY_INDEX) = 0
14170 print "spouse of"
14180 else
14190 let GENERATION GAP (KEY INDEX) = 1
14200 if EDGE TO PREDECCESSOR (THIS NODE) = PARENT then
14210 print "parent of"
14220 else ! edge is child-type
14230 print "child of"
14240 end if
14250 end if
14260 let KEY INDEX = KEY INDEX + 1
14270 let THI\overline{S}}\mathrm{ NODE = PATH}\mathrm{ PREDECESSOR (THIS NODE)
14280 loop
14290 print " "; NAME$ (THIS NODE)
14300 let PERSON INDEX (\overline{KEY INDEX) = THIS NODE}
14310 let RELATIO\overline{N TO NEXT (KEY INDEX) = NULL RELATION}
14320 let RELATION_TO-NEXT (KEY_INDEX + 1) = NULL_RELATION
14330 !
```

```
14340 ! Resolve CHILD-PARENT and CHILD-SPOUSE-PARENT relations
14350 ! to SIBLING relations.
14360 let KEY INDEX = 1
14370 do unti\overline{l}}\mathrm{ RELATION TO NEXT (KEY INDEX) = NULL RELATION
14380 if RELATION TO- NE\overline{XT (KEY INDEX) = CHILD then}
14390 let LATE\overline{R}K\overline{EY RELATION = RELATION_TO_NEXT (KEY_INDEX + 1)}
14400 if LATER KEY \overline{RELATION = PARENT then}
14410 ! found either full or half SIBLINGs
14420 let GENERATION_GAP (KEY INDEX) = 0
14430 let. RELATION_TO_NEXT (KEY_INDEX) = SIBLING
14440 let PROXIMITY - (KEY INDEX) = &
&
&
14450
14460 else
14470 if LATER KEY RELATION = SPOUSE and
&
&
14480
14490
14500 let RELATION T\overline{O_NEXT (KEY_INDEX) = SIBLING}
14510 let PROXIMIT\overline{Y - (KEY_INDEX) = STEP}
14520 call CONDENSE_KEY_PERSONS (KEY_INDEX, 2)
14530 end if
14540 end if
14550 end if
14560 let KEY_INDEX = KEY_INDEX + 1
14570 1oop
14580 !
14590 ! Resolve CHILD-CHILD-... and PARENT-PARENT-... relations to
14600 ! direct descendant or ancestor relations.
14610 let KEY INDEX = 1
14620 do unti\overline{1} RELATION TO NEXT (KEY INDEX) = NULL_RELATION
14630 if RELATION_TO_NEXT (KEY_INDEX) = CHILD or &
& RELATION_TO NEXT (KEY_INDEX) = PARENT then
14640 let LATER KEY INDEX = KEY INDEX + 1
14650 do while \overline{RELATION_TO_NEXT (LATER_KEY_INDEX) &}
&
14660
14670 100p
14680 let GENERATION COUNT = LATER KEY INDEX - KEY INDEX
14690 if GENERATION \overline{COUNT > 1 then -! compress generations}
14700 let GENERATTION GAP (KEY INDEX) = GENERATION COUNT
14710 call CONDENSE_KEY_PERSONS (KEY_INDEX, GENERATION_COUNT - 1)
14720 end if
14730 end if
14740 let KEY_INDEX = KEY_INDEX + 1
14750 loop
14760 !
```

```
14770 ! Resolve CHILD-SIBLING-PARENT to COUSIN,
```

14780 !
14790 !

```
\[
\begin{array}{cl}
\text { Resolve } & \text { CHILD-SIBLING-PARENT } \\
\text { CHILD-SIBLING } & \text { to NEPHEW, } \\
\text { SIBLING-PARENT } & \text { to UNCLE. }
\end{array}
\]
```

```
14800 let KEY INDEX = 1
```

14800 let KEY INDEX = 1
14810 do unti\overline{1}}\mathrm{ RELATION TO NEXT (KEY INDEX) = NULL RELATION
14820 let LATER KEY \overline{RELATTION = RELATION_TO_NEXT``(KEY_INDEX + 1)}
14830 if RELATION T\overline{O} NEXT (KEY INDEX) = CHĪLD - \&
\&
14840
14850
14860
14870
14880
14890
14900
14910
14920
14930
14940
14950
14960
14970
14980
14990
15000
\&
15010
15020
\&
15030
15040
15050
15060
15070 let KEY INDEX = KEY INDEX + 1
15080 10op
15090 !

```
```

15100 ! Loop below will pick out valid ad jacent strings of elements
15110 ! to be printed. KEY_INDEX points to first element,
15120 ! LATER_KEY_INDEX to last element, and PRIMARY_INDEX to the
15130 ! element which determines the primary English word to be used.
15140 ! Associativity of adjacent elements in condensed table
15150 ! is based on English usage.
15160 print " Condensed path:"
15170 let KEY INDEX = 1
15180 do unti\ RELATION TO NEXT (KEY INDEX) = NULL RELATION
15190 let KEY RELATION = RELATION TO NEXT (KEY INDEX)
15200 let LATER KEY INDEX, PRIMAR\overline{Y}}\mathrm{ INDEX = KEY INDEX
15210 if RELATIO\overline{N}_T\overline{O}}\mathrm{ NEXT (KEY_INDEX + 1) <> NUULL_RELATION then
15220 ! seek multi-element combination
15230 let ANOTHER ELEMENT POSSIBLE = TRUE
15240 if KEY RELATION = SPOUSE then
let LATER KEY INDEX = LATER KEY INDEX + 1
let PRIMA\overline{R}Y I\overline{NDEX = LATER KEY INDEX}
if RELATION TO NEXT (LATER \overline{KEY INDEX) = SIBLING or \&}
RELATION-TO-NEXT (LATER KEY-INDEX) = COUSIN then
! nothīng can follow \overline{spouse-sibling or spouse-cousin}
let ANOTHER ELEMENT POSSIBLE = FALSE
end if
end if
! PRIMARY INDEX is now correctly set. Next if-statement
determines if a following SPOUSE relation should be
appended to this combination or left for the next
combination.
if RELATION TO NEXT (PRIMARY INDEX + 1) = SPOUSE and \&
ANOTHER ELEMENT POSSIBLE \ TRUE then
! Onl\overline{y a SPOUSE can follow a Primary}
! check primary preceding and following SPOUSE.
let PRIMARY RELATION = RELATION TO NEXT (PRIMARY INDEX)
let NEXT_PRTMARY_RELATION = RELATION_TO-NEXT (PRIMARY - INDEX + 2)
if (NEXT_PRIMARY_RELATION = NEPHEW -or- \&
NEXT_PRIMARY_RELATION = COUSIN or \& \&
NEXT PRIMARY RELATION = NULL RELATION) \&
or (P\overline{RIMARY RELATION = NEPHEW\overline{)}}\mathrm{ (')}\&\&
or ( (PRIMARY RELATION = SIBLING or \&
PRIMARY RELATION = PARENT) \&
and NEXT PRIMARY_RELATION > UNCLE ) then
! append following SPOUSE with this combination
let LATER_KEY_INDEX = LATER_KEY_INDEX + I
end if
end if
end if ! multi-element combination
call DISPLAY RELATION (KEY_INDEX, LATER_KEY INDEX, PRIMARY_INDEX)
let KEY_INDE }\overline{\textrm{X}}=\mathrm{ LATER_KEY_\
1oop
!
15510 print " "; NAME\$ (PERSON_INDEX (KEY_INDEX))
15520 end sub
15530 ! end of RESOLVE_PATH_TO_ENGLISH
15540

```
```

15550 function SIBLING PROXIMITY (INDEX1, INDEX2)
15560 ! Determines whether two PERSONs are full siblings, i.e.,
15570 ! have the same two parents.
15580 if RELATIVE IDENTIFIER\$ (INDEX1, FATHER IDENT) <> NULL IDENT\$ and \&
\& RELATIVE_IDENTIFIER\$ (INDEX1, MOTHER_IDENT) <> NULL_IDENT\$ and \&
\& RELATIVE_IDENTIFIER\$ (INDEXI, FATHER_IDENT) = - \&
\& RELATIVE_IDENTIFIER\$ (INDEX2, FATHERIIDENT) and \&
\& RELATIVE_IDENTIFIER\$ (INDEX1, MOTHER_IDENT) = \&
\& RELATIVE_IDENTIFIER\$ (INDEX2, MOTHER_IDENT) then
15590 let SIBLİNG_PROXIMITY = FULL
15600 else
15610 let SIBLING PROXIMITY = HALF
15620 end if
15630 end function ! SIBLING PROXIMITY
15640 !
15650 sub CONDENSE KEY PERSONS (AT INDEX, GAP SIZE)
15660 ! CONDENSE_KEY_PERSONS condenses superffluous entries from the
15670 ! key persōn array entries, starting at AT_INDEX
15680 let RECEIVE_INDEX = AT_INDEX
15690 do
15700 let RECEIVE_INDEX = RECEIVE_INDEX + 1
15710 let SEND_INDEX = RECEIVE_INDEX + GAP_SIZE
15720 let RELAT

```

```

15740 let GENERA\overline{TION_GAP (RECEIVE_INDEX) = GENERATION_GAP (SEND_INDEX)}
15750 let PROXIMITY - (RECEIVE - INDEX) = PROXIMITY - (SEND INDEX)
15760 let COUSIN RANK (RECEIVE_INDEX) = COUSIN RANK (SEND_INDEX)
15770 1oop until RELATION_TO_NEXT (SEND_INDEX) = NULL_\overline{RELATION}
15780 end sub
15790 !
15800 sub DISPLAY RELATION (FIRST INDEX, LAST INDEX, PRIMARY INDEX)

```

```

15820 ! condensed table and generates the English description of
15830 ! the relation between the first and last + 1 elements.
15840 !
15850 let FIRST_RELATION = RELATION_TO NEXT (FIRST INDEX)
15860 let LAST_\overline{RELATION = RELATION_TO_NEXT (LAST_INDEX)}
15870 let PRIMA\overline{RY_RELATION = RELATION_TO_NEXT (PRIMARMYINDEX)}
15880 !
15890 ! set THIS PROXIMITY
15900 if (PRIMARY RELATION = PARENT and FIRST RELATION = SPOUSE) or \&
\& (PRIMARY_RELATION = CHILD and LAST_\overline{RELATION = SPOUSE) then}
15910 let THIS_PROXIMITY = STEP
15920 else
15930 if PRIMARY_RELATION = SIBLING or \&
\& PRIMARY_RELATION = UNCLE or \&
\& PRIMARY_RELATION = NEPHEW or \&
\& PRIMARY RELATION = COUSIN then
15940 let THIS_PROXIMITY = PROXIMITY (PRIMARY_INDEX)
15950 else
15960 let THIS_PROXIMITY = FULL
15970 end if
15980 end if
15990 !

```

```

16380 elseif PRIMARY RELATION = COUSIN and THIS_COUSIN RANK > 1 then

```

16390
16400
16410
16420
16430
16440
16450
16460
16470
16480
16490
16500 end
16510 !
16520 if THIS PROXIMITY \(=\) STEP then
16530 print "step-";
16540 elseif THIS PROXIMITY = HALF then
16550 print "half-";
16560 end if
16570 !
16580 let THIS_GENDER\$ = GENDER\$ (PERSON_INDEX (FIRST_INDEX))
16590 select case PRIMARY RELATION
16600 case 1 ! PARENT
16610 if THIS GENDER\$ = MALE\$ then print "father"; else print "mother";
16620 case 2 ! CHILD
16630 if THIS GENDERS = MALE\$ then print "son"; else print "daughter";
16640 case 3 T SPOUSE
16650 if THIS GENDER \(=\) MALE \(\$\) then print "husband"; else print "wife";
16660 case 4 ! SIBLING
16670 if THIS GENDER\$ = MALE\$ then print "brother"; else print "sister";
16680 case 5 ! UNCLE
16690 if THIS GENDER\$ = MALE\$ then print "uncle"; else print "aunt";
16700 case 6 ! NEPHEW
16710 if THIS GENDER\$ = MALES then print "nephew"; else print "niece";
16720 case 7 ! COUSIN
16730 print "cousin";
16740 case else
16750 print "null";
16760 end select
16770 !
16780 if INLAW = TRUE then print "-in-law";
16790 !
16800 if PRIMARY RELATION \(=\) COUSIN and THIS GENERATION GAP \(>0\) then
16810 if THIS GENERATION GAP \(>1\) then
16820 print THIS_GENERATION_GAP; "times removed";
16830 else
16840 print " once removed";
16850 end if
16860 end if
16870 !
16880 print " of"
16890 !
16900 end sub ! end of internal sub DISPLAY RELATION
16910 end sub ! end of external sub FIND_RELATIONSHIP
16920 !

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```

16930 ! ---- program-unit number 3 ----
16940 !
16950 external sub COMPUTE COMMON GENES (INDEX1, INDEX2, IDENTIFIER\$ ()
\& NEIGHBOR_COUNT (), NEIG\overline{GBOR INDEX (,), NEIGHBOR_EDGE (,),}
\& DESCENDANT IDENTIFIER\$ (), DESCENDANT GENES ())
16960 !
16970 ! COMPUTE COMMON GENES assumes that each ancestor contributes
16980 ! half of the geñetic material to a person. It finds common
16990 ! ancestors between two persons and computes the expected
17000 ! value of the PROPORTION of common material.
17010 !
17020 declare sub ZERO_PROPORTION, MARK PROPORTION, CHECK_COMMON PROPORTION
17030 !
17035 option base 1
17040 !
17045 data 1, 2, 3, 4, 5, 6, 7, 8
17050 read PARENT, CHILD, SPOUSE, SIBLING, UNCLE, NEPHEW
17055 read COUSIN, NULL_RELATION
17057 !
17060 ! Begin main line of execution of COMPUTE COMMON GENES
17065!
17070 ! First zero out all ancestors to allow adding. This is necessary
17075 ! because there might be two paths to an ancestor.
17080 call ZERO_PROPORTION (INDEX1, 0)
17090 ! now mark with shared PROPORTION
17100 cal1 MARK PROPORTION (IDENTIFIER\$ (INDEX1), 1.0, INDEX1, 0)
17110 let COMMO\overline{N PROPORTION = 0.0}
17120 ca11 CHECK COMMON_PROPORTION (COMMON PROPORTION, \&
\& ID IDENTIFIER\$ (INDEX1), 1.0, 0.0, INDEX2, 0)
17130 print using " Proportion of common genetic material = 非非非非^^^^"": \&
\& COMMON PROPORTION
17140 !
17150 ! End main line of execution of COMPUTE_COMMON_GENES
17160 !
17170 sub ZERO PROPORTION (ZERO INDEX, THIS NEIGHBOR)
17180 ! Z\overline{ERO_PROPORTION recursively seeks out all ancestors and}
17190 ! zeros them out
17200 let DESCENDANT GENES (ZERO_INDEX) = 0.0
17210 for THIS_NEIGHB
17220 if NEIGHBOR EDGE (ZERO_INDEX, THIS_NEIGHBOR})=\mathrm{ PARENT then
17230
\&
17240 end if
17250 next THIS NEIGHBOR
17260 end sub -! ZERO PROPORTION
17270 !

```


\subsection*{4.0 C}

The identifiers NULL and FILE are capitalized, even though they are supplied by the standard run-time library, because identifiers in C are case-sensitive, e.g., "null" is not equivalent to "NULL".
```

/* Bring in standard routines for run-time support */
\#include <stdio.h>
/* Global types and objects */
typedef short int BOOLEAN;
\#define TRUE 1
\#define FALSE 0
\#define EQUALS 0
\#define NULL ID "000"
\#define NULL_CHR - \0`
\#define MAX PERS 300
\#define NAME LEN 20
/* every PERSON has a unique 3-digit IDENT */
\#define ID LEN 3
\#define BUF}\mathrm{ LEN 60
/* Use "+ l" when treating type as variable-length - extra character
used to hold NULL_CHR termination character. */
typedef char NAME TYP [NAME LEN + 1];
typedef char BUF_TYPE [BUF_LEN + 1];
typedef char MSG-TYPE [40 \mp 1];
typedef char ID_TYPE [ID_LEN + 1];
typedef int INDX_TYP, COUNTER;
/* each PERSON's record in the file identifies at most thre,
others directly related: father, mother, and spouse */
typedef short int GIVEN_ID;
\#define FATHR ID 0
\#define MOTHR ID 1
\#define SPOUS ID 2
\#define MAX_G\overline{VEN 3}
typedef ID_TYPE REL_ARRY [MAX_GVEN];
\#define REQ_OK "Request OK"
\#define REQ_STOP "stop"
typedef char GNDR_TYP;
\#define MALE -M'
\#define FEMALE 'F'

```
```

    typedef unsigned int REL_TYPE;
    /* Values defined as octal powers of two to facilitate comparisons
    of one relation with several possibilities. */
    \#define PARENT 0001
\#define CHILD 0002
\#define SPOUSE 0004
\#define SIBLING 0010
\#\#define UNCLE 0020
\#define NEPHEW 0040
\#define COUSIN 0100
\#define NULL REL 0200
/* directed edges in the graph are of a given type */
typedef REL_TYPE EDG_TYPE;
/* A node in the graph (= PERSON) has either already been reached,
is immediately adjacent to those reached, or farther away. */
typedef short int REACH TY;
\#define REACHED 1
\#define NEARBY 2
\#define NOT SEEN 3
/* each PERSON has a linked list of adjacent nodes, called neighbors */
typedef struct NBR_NODE
{ INDX TYP NBR DEX;
EDG TYPE NBR EDGE;
struct NBR_NODE *NEX\overline{T}}\mathrm{ NBR;
}
NBR_REC, *NBR_PTR;
/* All relationships are captured in the directed graph of which
each record is a node. */
typedef struct
{
/* static information - filled from PEOPLE file: */
NAME TYP NAME;
ID TYPE IDENT;
GNDR TYP GENDER;
/* IDENTs of immediate relatives - father, mother, spouse */
REL ARRY REL ID;
/* head of linked list of adjacent nodes */
NBR PTR NBR HDR;
/* da\overline{ta used when traversing graph to resolve user request: */}
float DIST SRC;
INDX TYP PATH\overline{PRED;}
EDG TYPE EDG PRED;
REAC
/* data-used to compute common genetic material */
ID_TYPE DSC_ID;
f1ōat DSC-GENE;
}
PERS_REC;

```
/* the PERSON array is the central repository of information about inter-relationships. */
PERS REC PERSON [MAX PERS];
INDX_TYP NUM_PERS;
/* Key persons are the ones in the REL_SHIP path which remain after the path is condensed. */
typedef short int SIB_TYPE;
\#define STEP 1
\#define HALF 2
\#define FULL 3
typedef struct
\{ REL TYPE REL NEXT; IND \(\bar{X}\) TYP PERS DEX; COUNTER GEN GAP; SIB TYPE PROXIMTY; COUNTTER CUZ RANK;
\}
KEY_REC;
/********** Main line of execution RELATE **********/
main ()
\{ /* These variables are used when establishing the PERSON array from the PEOPLE file. */
FILE \(\quad\) fopen(), *PEOPLE;
register INDX TYP CURRENT, PREVIOUS;
ID TYPE - PREV_ID, CUR_ID;
GIVEN_ID
REL SHIP;
char INP_BUF [100];
/* These variables are used to accept and resolve requests for REL SHIP information. */
COUNTER SEMI LOC;
BUF TYPE
REQ \(\bar{B} U F ;\)
BUF TYPE
P1 IDENT, P2 IDENT;
COUNTER
P1-FOUND, P2 FOUND;
MSG TYPE
ERR MSG;
INDX_TYP
P1_INDEX, P2_INDEX;
/* *** execution of main sequence begins here *** */
```

    PEOPLE = fopen("PEOPLE.DAT", "r");
    /* This loop reads in the PEOPLE file and constructs the PERSON
        array from it (one PERSON == one record == one array entry).
        As records are read in, links are constructed to represent the
        PARENT-CHILD or SPOUSE REL_SHIP. The array then implements
        a directed graph which is used to satisfy subsequent user
        requests. The file is assumed to be correct - no validation
        is performed on it. */
    READ PEO:
for
{
/* copy direct information from file to array */
if (FXD_GETC (PERSON [CURRENT] - NAME, PEOPLE, NAME LEN)
break;
FXD GETC (PERSON [CURRENT] . IDENT, PEOPLE, ID LEN);
FXD_GETC (\&(PERSON [CURRENT] . GENDER), PEOPLE, 1);
for (REL SHIP = FATHR ID; REL SHIP < MAX GVEN; REL_SHIP++)
FXD_\overline{GETC (PERSON [CURRENT] . REL_ID [REL_SHIP], PEOPLE, ID_LEN);}
/* flush}\mathrm{ remainder of record */
fgets (INP BUF, 100, PEOPLE);
/* Location of adjacent persons as yet undetermined */
PERSON [CURRENT] . NBR HDR = NULL;
/* Descendants as yet undetermined */
strcpy (PERSON [CURRENT] . DSC ID, NULL ID);
/* Compare this PERSON against all previously entered PERSONs
to search for REL SHIPs. */
strcpy (CUR_ID, PERSON [CURRENT] . IDENT);
CMP PREV:
for (PREVIOUS = 0; PREVIOUS < CURRENT; PREVIOUS++)
{
strcpy (PREV_ID, PERSON [PREVIOUS] . IDENT);
/* Search for
either direction between this and PREVIOUS PERSON.
Assume at most one REL_SHIP exists. */
TRY RELS:
for (REL_SHIP = FATHR ID; REL_SHIP < MAX_GVEN; REL_SHIP++)
{
if (STREQ (PREV ID, PERSON [CURRENT] . REL_ID [REL_SHIP]))
LINK REL (CURRENT, REL_SHIP, PREVIOUS);
break;
}
if (STREQ (CUR_ID, PERSON [PREVIOUS] . REL_ID [REL_SHIP]))
LINK_REL (PREVIOUS, REL_SHIP, CURRENT);
break;
}
} /* end TRY RELS */
} /* end CMP P\overline{REV */}
} /* end READ PEO */
NUM PERS = CURRENT;
fclose (PEOPLE);

```
```

    /* PERSON array is now loaded and edges between immediate relatives
        (PARENT-CHILD or SPOUSE-SPOUSE) are established.
    While-1oop accepts requests and finds REL_SHIP (if any)
    between pairs of PERSONs. */
    PROC REQ:
whī1e (TRUE)
{
PROMPT (REQ BUF);
if (STREQ (\overline{REQ_BUF, REQ_STOP))}
break;
SEMI_LOC = CHK_RQST (REQ_BUF, ERR_MSG);
/* Syntax check of request completed. Now either display error
message or search for the two PERSONs. */
if (STREQ (ERR MSG, REQ OK))
{ /* Reques\overline{t}}\mathrm{ syntactically correct - search for requested PERSONs. */
REQ BUF [SEMI LOC] = NULL CHR;
BUF_PERS (REQ_BUF, 0, P1 IDENT);
BUF PERS (REQ buF, SEMI LOC + 1, P2 IDENT);
SEEK_PER (P1_IDENT, P2_IDENT, \& P1_INDEX, \& P2_INDEX,
\& P1 FOUND, \& P2 FOUND);
if (P1 FOUND == 1 \&\& P2 FOUND == 1)
/* Exactly one match for each PERSON - proceed to
determine REL_SHIP, if any. */
if (P1_INDEX == \overline{P}2_INDEX)
printf (" %1s is identical to %8s \n",
PERSON [P1 INDEX] . NAME,
(PERSON [P\overline{1}INDEX] . GENDER == MALE) ?
"himself." : "herself.");
else
FIND REL (P1 INDEX, P2 INDEX);
else /* \overline{either not found or}\mathrm{ more than one found */}
if (P1_FOUND == 0)
printf (" First person not found.\n");
else if (P1_FOUND > 1)
{
printf (" Duplicate names for first person -");
printf (" use numeric identifier.\n");
}
if (P2 FOUND == 0)
priñtf (" Second person not found.\n");
else if (P2_FOUND > 1)
{
printf (" Duplicate names for second person -");
printf (" use numeric identifier.\n");
}
} /* end processing of syntactically legal request */
else
printf (" Incorrect request format: %1s \n", ERR_MSG);
} /* end PROC_REQ loop */
printf (" End of relation-finder. \n");
}
/* End of main line of RELATE */

```
```

/* procedures under RELATE */
FXD_GETC (RECEIVER, SENDING, GET LEN)
char *RECEIVER;
FILE *SENDING;
int
GET LEN;
{ register int CHAR CNT;
for (CHAR CNT = 0;
CHAR CNT++ < GET LEN \&\& (*RECEIVER++ = getc (SENDING)) != EOF ; ) ;
if (CHAR_\overline{CNT > = GET_LEN)}
{
*RECEIVER = NULL_CHR;
return !EOF;
}
else
return EOF;
}
STREQ (STRING1, STRING2)
/* compare for equality, ignore trailing spaces */
register char *STRING1, *STRING2;
{ register char *LONGER;
for (; *STRING1 == *STRING2; STRING1++,STRING2++)
if (*STRING1 == NULL_CHR)
return TRUE;
if (*STRING1 == NULL CHR)
LONGER = STRING2;
else
if (*STRING2 == NULL CHR)
LONGER = STRING1;
else
return FALSE;
for ( ; *LONGER++ == ' '; ) ;
return (*--LONGER == NULL_CHR);
}

```
```

LINK REL (FROM DEX, REL SHIP, TO INDEX)
/* establishē cross-indexing \overline{between immediately related PERSONs. */}
register INDX TYP FROM_DEX, TO INDEX;
register GIVEN_ID REL_SHIP;
{ /* execution of LINK REL */
if (REL_SHIP == SPOUS_ID)
{
LINK_ONE (FROM_DEX, SPOUSE, TO_INDEX);
LINK_ONE (TO_INNDEX, SPOUSE, FROM_DEX);
}
else /* REL SHIP is father or mother */
{
LINK ONE (FROM DEX, PARENT, TO INDEX);
LINK_ONE (TO_INDEX, CHILD, FROM_DEX);
}
}
LINK_ONE (FROM_DEX, THIS_EDG, TO_INDEX)
/* Establishes the NBR_REC from one PERSON to another */
INDX TYP FROM DEX, TO_INDEX;
EDG TYPE THIS_EDG;
{ register NBR_PTR NEW_NBR;
NEW NBR = (NBR REC * ) calloc(1, sizeof(NBR REC));
NEW NBR -> NBR DEX = TO INDEX;
NEW_NBR -> NBR EDGE = THIS EDG;
NEW NBR -> NEXT NBR = PERSON [FROM DEX] . NBR HDR;
PERSON [FROM_DEX] . NBR_HDR = NEW NBR;
}
PROMPT (REQ BUF)
/* Issues prompt for user-request, reads in request,
blank-fills buffer, and skips to next line of input. */
BUF_TYPE REQ BUF;
{
printf (" \n");
printf (" ----------------------------------------------------------
printf (" Enter two person-identifiers (name or number),\n");
printf (" separated by semicolon. Enter \"stop\" to stop.\n");
fgets (REQ BUF, BUF_LEN, stdin);
for ( ; *REQ BUF++ != -\n' ; ) ;
*--REQ BUF =-'\0';
}

```
```

CHK RQST (REQ_BUF, REQ STAT)
/\overline{*}}\mathrm{ Performs syntactic}\mathrm{ check on request in buffer. */
BUF_TYPE REQ_BUF;
MSG_TYPE REQ_STAT;
{ COUNTER SEMI_LOC = 1,
register COUNTER BUF_\overline{DEX;}
BOOLEAN P1_EXIST = FALSE,
P2_EXIST = FALSE;
strcpy (REQ STAT, REQ_OK);
for (BUF_DEX = O; BUF_DEX < BUF_LEN \&\& REQ BUF [BUF_DEX]; BUF_DEX++)
{
if (REQ BUF [BUF DEX] != - -)
if (\overline{REQ_BUF [\overline{BUF_DEX] == ';'`)}}\mathbf{{}=\mp@subsup{}{}{\prime}
SEMI LOC = BUF DEX;
SEMI_CNT = \EMM_CNT + 1;
}
else /* Check for non-blanks before/after semicolon. */
if (SEMI_CNT < 1)
P1_EXIST = TRUE;
else
P2 EXIST = TRUE;
}
/* set REQ STAT, based on results of scan of REQ_BUF. */
if (SEMI_CNT != 1)
strcpy (REQ STAT, "must be exactly one semicolon.");
else if ( ! P1 EXIST)
strcpy (REQ STAT, "null field preceding semicolon.");
else if ( ! P2_EXIST)
strcpy (REQ_STAT, "null field following semicolon.");
return SEMI_LOC;
}
BUF PERS (REQ BUF, BUF DEX, PERS ID)
/\overline{*}}\mathrm{ fills in the PERS_ID from the designated portion
of the REQ_BUF, d\overline{leting leading blanks. */}
BUF TYPE REQ_BUF;
register COUNTER BUF_DEX;
NAME_TYP PERS_ID;
{
for ( ; REQ_BUF [BUF DEX++] == - '; ) ;
strcpy (PER\overline{S}_ID, \&REQ\_BUF [--BUF_DEX]);
}

```
```

SEEK_PER (P1_IDENT, P2_IDENT, P1_INDEX, P2_INDEX,
P1 FOUND, P2 FOUND)
/* SEEK PER scans through the PERSON array,
looking for the two requested PERSONs. Match may be by NAME
or unique IDENT-number. */
BUF TYPE IND $\bar{X}$ TYP COUNTER
\{ register INDX_TYP
*P1 INDEX $=0 ;$
*P2 ${ }^{\text {INDEX }}=0 ;$
*P1 FOUND $=0$;
*P2 FOUND $=0$;
SCAN PER:
for (CURRENT $=0$; CURRENT < NUM_PERS; CURRENT++)
\{
/* allow identification by name or number. */
if (STREQ (P1_IDENT, PERSON [CURRENT] . IDENT) II STREQ (P1-IDENT, PERSON [CURRENT] . NAME))
\{ (*P1 FOUND) ++;
*P1_INDEX = CURRENT;
\}
if (STREQ (P2 IDENT, PERSON [CURRENT] . IDENT) || STREQ (P2_IDENT, PERSON [CURRENT] . NAME))
\{
(*P2 FOUND) ++;
*P2 INDEX = CURRENT;
\}
\} /* end SCAN PER loop */
\} /* end of SEEK_PER */

```
```

FIND REL (TARG DEX, SRCE DEX)
/* Finds shortest path (if any) between two PERSONs and
determines their REL_SHIP based on immediate relations
traversed in path. PERSON array simulates a directed graph,
and algorithm finds shortest path, based on following
weights: PARENT-CHILD edge = 1.0
SPOUSE-SPOUSE edge = 1.8 */
INDX_TYP
\{ register INDX_TYP
INDX_TYP
register NBR_PTR
float
typedef short int
\#define SEARCHNG
\#define SUCCESS
\#define FAILED
SRCH_TYP SRCH_ST;
/* begin execution of FIND_REL */
/* initialize PERSON-array for processing -
mark all nodes as not seen */
for (PERS DEX = 0; PERS DEX < NUM PERS; PERS DEX++)
PERSO\overline{N [PERS DEX] - REACH ST = NOT SEEN;}
THIS_NOD = SRCE_DEX;
/* mark source node as REACHED */
PERSON [THIS NOD] . REACH ST = REACHED;
PERSON [THIS_NOD] . DIST_SRC = 0.0;
/* no NEARBY nodes exist yet */
LST_NRBY = -l;
SRC\overline{M}}\mathbf{ST}=\mathrm{ (THIS_NOD == TARG_DEX) ? SUCCESS : SEARCHNG;

```
```

    /* Loop keeps processing closest-to-source, unREACHED node
        until target REACHED, or no more connected nodes. */
    SEEKTARG:
while (SRCH ST == SEARCHNG)
{ /* Process all nodes adjacent to THIS NOD */
for (THIS NBR = PERSON [THIS NOD] . NBR \overline{HDR;}
THIS NBR != NULL;
THIS NBR = THIS NBR }->\mathrm{ NEXT NBR)
PROC ADJ (THIS NOD, THIS_NBR -> NBR_DEX, THIS_NBR -> NBR EDGE,
NRBY ND, \&LST NRBY);
/* All nodes adjacent to THIS NOD are set. Now search for
shortest-distance unREACHED (but NEARBY) node to process next. */
if (LST NRBY == -1)
SRCH ST = FAILED;
else /** determine next node to process */
{
MIN DIST = 1.0E+18;
for (PERS DEX = 0; PERS DEX <= LST NRBY; PERS DEX++)
if (PERSON [NRBY_ND [\overline{PERS_DEX]] - DIST_SRC < MIN_DIST)}
{
BEST DEX = PERS DEX;
MIN DIST = PERSON [NRBY ND [PERS DEX]] . DIST SRC;
}
/* establish new THIS NOD */
THIS NOD = NRBY ND [BEST DEX];
/* change THIS N
PERSON [THIS NOD] - REACH ST = REACHED;
/* remove THIS NOD from NEARBY list */
NRBY ND [BEST \overline{DEX] = NRBY ND [LST NRBY--];}
if (THIS NOD \equiv= TARG DEX)
SRCH_\overline{ST}= SUCCESS;
}
} /* end SEEKTARG loop */
/* Shortest path between PERSONs now established. Next task is
to translate path to English description of REL_SHIP. */
if (SRCH ST == FAILED)
print\overline{f}}\mathrm{ (" %ls is not related to %ls\n",
PERSON [TARG DEX] - NAME, PERSON [SRCE DEX] . NAME);
else /* success - parse path to find and display REL SHIP */
{
RESOLVE (SRCE DEX, TARG_DEX);
CMPT_GNS (SRCE_DEX, TARG_DEX);
}
} /* end FIND_REL */

```
```

/* procedures under FIND REL */
PROC ADJ (BASENODE, NXT NODE, N B EDGE, NRBY ND, LST NRBY)

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        If NXT NODE already REACHED, do nothing.
        If previously seen, check whether path thru BASENODE is
        shorter than current path to NXT_NODE, and if so re-link
        next to base.
        If not previously seen, link next to base node. */
    register INDX_TYP NXT NODE;
    INDX TYP - BASENODE, NRBY ND[], *LST NRBY;
    EDG_TYPE N_B_EDGE;
    { float WGHT EDG, DIST_BAS;
/* begin execution of PROC ADJ */
if (PERSON [NXT NODE] - REACH ST != REACHED)
{
WGHT EDG = (N_B_EDGE == SPOUSE) ? 1.8 : 1.0;
DIST_BAS = WGHT_EDG + PERSON [BASENODE] . DIST_SRC;
if (\overline{PERSON [NXT_NODE] - REACH_ST == NOT_SEEN)}
{
PERSON [NXT NODE] - REACH ST = NEARBY;
NRBY ND [++-}*LSST NRBY] = NXT NODE;
/* l\overline{ink next to \overline{base by re-setting its predecessor index to}}0\mathrm{ (oun}
point to base, note type of edge, and re-set distance
as it is through base node. */
PERSON [NXT NODE] . DIST SRC = DIST BAS;
PERSON [NXT NODE] . PATHPRED = BASENODE;
PERSON [NXT_NODE] . EDG PRED = N B EDGE;
}
else /* REACH ST = NEARBY */
if (DIST BAS < PERSON [NXT NODE] . DIST_SRC)

```

```

                    point to base, note type of edge, and re-set distance
                    as it is through base node. */
                        PERSON [NXT NODE] . DIST SRC = DIST BAS;
                        PERSON [NXT NODE] . PATH\overline{PRED = BASENNODE;}
                        PERSON [NXT_NODE] . EDG_PRED = N_B_EDGE;
            }
        }
    } /* end PROC ADJ */

```
```

RESOLVE (SRCE DEX, TARG_DEX)
/* RESOLVE condenses the shortest path to a
series of REL_SHIPs for which there are English
descriptions.**/
INDX TYP SRCE DEX, TARG_DEX;
{ /* these variables are used to generate KEY_PERSs */
COUNTER GENCNT;
/* these variables are used to condense the path */
KEY_REC KEY_PERS [MAX PERS];
REL_TYPE KEY_REL, LKEY_REL, PRIM_REL, NXT_PRIM;
register INDX_TYP KEYDEX;
INDX TYP - LKEY_DEX, PRIM_DEX, THIS_NOD;
BOOLEAN SEEKMORE;
/* begin execution of RESOLVE */
printf (" Shortest path between identified persons: \n");
/* Display path and initialize KEY PERS array from path elements. */
TRAVERSE:
for (THIS NOD = TARG DEX, KEY DEX = 0; THIS NOD != SRCE DEX;
THIS_NOD = PERSONN [THIS_N̄OD] . PATHPRED\overline{D}, KEY_DEX++)
{
printf (" %1s is ", PERSON [THIS NOD] . NAME);
KEY_PERS [KEY_DEX] . PERS_DEX = THTIS_NOD;
KEY_PERS [KEY-DEX] . PROXIMTY = FULL;
KEY_PERS [KEY_DEX] . REL NEXT = PERSON [THIS_NOD] . EDG_PRED;
swi\overline{tch (PERSON}[THIS_NOD]}\mathrm{ . EDG_PRED)
{
case PARENT: printf ("parent of\n");
KEY_PERS [KEY DEX] . GEN_GAP = 1;
break;
case CHILD : printf ("child of\n");
KEY_PERS [KEY DEX] . GEN_GAP = 1;
break;
case SPOUSE: printf ("spouse of\n");
KEY_PERS [KEY_DEX] . GEN_GAP = 0;
break;
} /* end switch */
} /* end TRAVERSE loop */
printf (" %1s\n", PERSON [THIS NOD] . NAME);
KEY PERS [KEY DEX] . PERS DEX = THIS NOD;
KEY_PERS [KEY DEX] . REL \overline{NEXT = NULL_REL;}
KEY_PERS [KEY_DEX + 1] . REL_NEXT = NULL_REL;

```
```

    /* Resolve CHILD-PARENT and CHILD-SPOUSE-PARENT relations
        to SIBLING relations. */
    FIND SIB:
for
{
if (KEY_PERS [KEY DEX] . REL_NEXT == CHILD)
LKEY_REL = KEY_PERS [KEY_DEX + 1] . REL_NEXT;
if (\overline{LKEY_REL == PARENT)}
{ /* found either full or half SIBLINGs */
BOOLEAN FULL_SIB();
KEY_PERS [KEY DEX] . PROXIMTY =
FULL_SIB (\overline{KEY_PERS [KEY_DEX] . PERS_DEX,}
KEY_PERS [KEY-DEX + 2] . PERS_DEX)
? FULL : HALF;
KEY PERS [KEY DEX] . GEN GAP = 0;
KEY_PERS [KEY_DEX] . REL_NEXT = SIBLING;
CONDENSE (KEY_DEX, 1, KEY_PERS);
}
else
if (LKEY REL == SPOUSE
\&\& KEYYPERS [KEY DEX + 2] . REL_NEXT == PARENT)
{ /* found step-S\overline{IbLINGs */}
KEY PERS [KEY_DEX] . GEN_GAP = 0;
KEY_PERS [KEY_DEX] . PRO\overline{XIMTY = STEP;}
KEY PERS [KEYDEX] . REL NEXT = SIBLING;
CONDENSE (KEY_DEX, 2, KEY_ PERS);
}
} /* end if REL_NEXT == CHILD */
} /* end FIND_SIB loop */
/* Resolve CHILD-CHILD-... and PARENT-PARENT-... relations to
direct descendant or ancestor relations. */
FIND ANC:
for
{
if (KEY_PERS [KEY_DEX] . REL_NEXT == CHILD ||
KEY_PERS [KEY_DEX] . REL_NEXT == PARENT)
{
for (LKEY DEX = KEY DEX + 1;
KEY_PERS [LKEY_DEX] . REL_NEXT == KEY_PERS [KEY_DEX] . REL_NEXT;
LKEY DEX++) ;
GEN CNT = LKEY DEX - KEY DEX;
if (GEN_CNT > \overline{)} /* compress generations */
{
KEY PERS [KEY DEX] . GEN GAP = GEN CNT;
CONDENSE (KEY_DEX, GEN_C\overline{NT - 1, KEY`_PERS);}
}
} /* end if REL NEXT == CHILD or PARENT */
} /* end FIND_ANC Ioop */

```

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```

. /* Resolve CHILD-SIBLING-PARENT to COUSIN,
CHILD-SIBLING to NEPHEW,
SIBLING-PARENT to UNCLE. */
FIND CUZ:
fo\overline{r}}\mathrm{ (KEY_DEX = 0; KEY_PERS [KEY_DEX] . REL_NEXT != NULL_REL; KEY_DEX++)
{
LKEY REL = KEY PERS [KEY DEX + 1] . REL_NEXT;
if (\overline{KEY_PERS [KEY DEX] . - REL_NEXT == CHĨLD \&\& LKEY_REL == SIBLING)}
if (KEY PERS [KEY DEX + 2] . REL NEXT == PARENT)
{ /* found COUSIIN */
COUNTER GAP1, GAP2;
GAP1 = KEY PERS [KEY_DEX] . GEN_GAP;
GAP2 = KEY_PERS [KEY_DEX + 2] . GEN_GAP;
KEY_PERS [KEY DEX] - PROXIMTY = KEY_PERS [KEY_DEX + 1] . PROXIMTY;
KEY_PERS [KEY_DEX] . GEN_GAP
= (GAP1 < GAP2) ? (GAP2 - GAP1) : (GAP1 - GAP2);
KEY_PERS [KEY DEX] . CUZ_RANK = (GAP1 < GAP2) ? GAP1 : GAP2;
KEY PERS [KEY DEX] . REL NEXT = COUSIN;
CONDENSE (KEY_DEX, 2, KE\overline{Y}}\mathrm{ PERS);
}
else /* found NEPHEW */
{
KEY PERS [KEY DEX] . PROXIMTY = KEY PERS [KEY DEX + 1] . PROXIMTY;
KEY_PERS [KEY_DEX] . REL NEXT = NEPMEW;
CON\overline{DENSE (KEY_DEX, 1, KEY_ PERS);}
}
} /* end COUSIN or NEPHEW */
else
if (KEY PERS [KEY DEX] . REL NEXT == SIBLING \&\& LKEY REL == PARENT)
{ /\ found UN\overline{CLE */}
KEY PERS [KEY DEX] . GEN GAP = KEY PERS [KEY DEX + 1] . GEN GAP;
KEY PERS [KEY DEX] . REL NEXT = UNCLE;
CONDENSE (KEY`DEX, 1, KE\overline{Y}}\mathrm{ PERS);
}
} /* end FIND_CUZ loop */

```
```

    /* Loop below will pick out valid adjacent strings of elements
    to be displayed. KEY DEX points to first element,
    LKEY DEX to last element, and PRIM_DEX to the
    element which determines the primary English word to be used.
    Associativity of adjacent elements in condensed table
    is based on English usage. */
    printf (" Condensed path:\n");
    CONSLIDT:
for (KEY_DEX = 0; KEY PERS [KEY DEX] . REL_NEXT != NULL_REL;
KEY_DEX = LKEY_DEX + 1)
{
KEY REL = KEY PERS [KEY DEX] . REL NEXT;
LKEY DEX = KEY DEX;
PRIM DEX = KEY DEX;
if (KEY PERS [KEY DEX + 1] . REL NEXT != NULL REL)
{ /* seek multi
SEEKMORE = TRUE;
if (KEY REL == SPOUSE)
{
PRIM DEX = ++LKEY DEX;
/* Nōthing can fo\1low SPOUSE-SIBLING or SPOUSE-COUSIN */
SEEKMORE = ! (KEY PERS [LKEY DEX] . REL_NEXT \& (SIBLING | COUSIN));
}
/* PRIM DEX is now correctly set. Next if-statement
determines if a following SPOUSE relation should be
appended to this combination or left for the next
combination. */
if (SEEKMORE \&\& KEY PERS [PRIM DEX + 1] . REL NEXT == SPOUSE)
{ /* Only a SPOUSE can fol1ow a Primary;
check primary preceding and following SPOUSE. */
PRIM REL = KEY_PERS [PRIM DEX] . REL_NEXT;
NXT PRIM = KEY_PERS [PRIM_DEX + 2] . REL_NEXT;
if ((NXT PRIM \overline{\& (NEPHEW | COUSIN | NULL REL))}
(PRTM REL == NEPHEW)
|| ((PRI\overline{M}REL \& (SIBLING | PARENT)) \&\& NXT PRIM != UNCLE ))
/* append following SPOUSE with this combination. */
LKEY DEX++;
}
} /* end multi-element combination */
SHOW REL (KEY DEX, LKEY DEX, PRIM DEX, KEY_PERS);
} /*- end CONSLIDT loop **/
printf (" %1s\n", PERSON [KEY_PERS [KEY_DEX] . PERS_DEX] . NAME);
} /* end of RESOLVE */

```
```

BOOLEAN FULL_SIB (INDEX1, INDEX2)
/* Determines whether two PERSONs are full siblings, i.e.,
have the same two parents. */
register INDX_TYP INDEX1, INDEX2;
{
return
! STREQ (PERSON [INDEXI] . REL ID [FATHR ID], NULL ID) \&\&
! STREQ (PERSON [INDEXl] . RELID [MOTHR ID], NULLID) \&\&
STREQ (PERSON [INDEXI] . REL ID [FATHR ID}
PERSON [INDEX2] . REL ID [FATHR ID]) \&\&
STREQ (PERSON [INDEX1] . REL_ID [MOTHR_ID],
PERSON [INDEX2] . REL_ID [MOTHR_ID]);
}
CONDENSE (AT_INDEX, GAP SIZE, KEY_PERS)
/* CONDENSE condenses superfluous entries from the
KEY_PERS array, starting at AT_INDEX. */
register INDX TYP AT INDEX;
COUNTER - GA\overline{P}\mathrm{ SIZE;}
KEY REC KEY PERS [];
register INDX_TYP SEND_DEX;
do
{
AT INDEX++;
SEND DEX = AT INDEX + GAP SIZE;
KEY_\overline{PERS [AT_INDEX] = KEY_PERS [SEND_DEX];}
}
while (KEY_PERS [SEND_DEX] . REL_NEXT != NULL_REL);
}

```
```

/* procedures under RESOLVE */
SHOW_REL (FRST_DEX, LAST_DEX, PRIM_DEX, KEY_PERS)
/* SHOW_REL Takes 1, 2, or 3 adjacent elements in the
condensed table and generates the English description of
the relation between the first and last + 1 elements. */
INDX TYP FRST DEX, LAST DEX, PRIM DEX;
KEY \overline{REC KEY PERS [];}
BOOLEAN INLAW;
SIB TYPE THIS PRX;
GND\overline{R}}\mathrm{ TYP THIS GND;
short int SUFFIX;
register REL_TYPE FRST_REL, LAST_REL, PRIM_REL;
COUNTER
FRST_REL = KEY PERS [FRST_DEX] . REL_NEXT;
LAST REL = KEY PERS [LAST DEX] . REL NEXT;
PRIM_REL = KEY_PERS [PRIM_DEX] . REL_NEXT;
/* set THIS PRX */
if ((PRIM REL == PARENT \&\& FRST REL = SPOUSE) ||
(PRIMREL == CHILD \&\& LAST_REL == SPOUSE))
THIS_P\overline{RX = STEP;}
else
if (PRIM REL \& (SIBLING | UNCLE | NEPHEW | COUSIN))
THIS_PRX = KEY_PERS [PRIM_DEX] . PROXIMTY;
else
THIS_PRX = FULL;
/* set THIS GAP */
if (PRIM REL \& (PARENT | CHILD | UNCLE | NEPHEW | COUSIN))
THIS_\overline{GAP = KEY_PERS [PRIM DEX] . GEN_GAP;}
else
THIS_GAP = 0;
/* set INLAW */
INLAW = FALSE;
if (FRST REL == SPOUSE \&\& (PRIM REL \& (SIBLING | CHILD | NEPHEW | COUSIN)))
INLAW = TRUE;
else
if (LAST REL == SPOUSE \&\&
(PRIMMREL \& (SIBLING | PARENT | UNCLE | COUSIN)))
INLAW = TRUE;
/* set THIS CUZ */
if (PRIM REL == COUSIN)
THIS_\overline{CUZ = KEY_PERS [PRIM_DEX] . CUZ_RANK;}
else
THIS_CUZ = 0;

```
```

/* parameters are set - now generate display. */
printf (" %ls is ", PERSON [KEY PERS [FRST DEX] . PERS DEX] . NAME);
if (PRIM REL \& (PARENT | CHILD T UNCLE | NEPHEW))
{ /* \overline{display generation-qualifier */}
if (THIS_GAP >= 3)
{
printf ("great");
if (THIS_GAP > 3)
print\overline{f}("*%1d", THIS_GAP - 2);
printf ("-");
}
if (THIS GAP >= 2)
print\overline{f ("grand-");}
}
else
if (PRIM_REL == COUSIN \&\& THIS_CUZ > 1)
{
printf ("%1d", THIS CUZ);
SUFFIX = THIS CUZ % 10;
switch (SUFFI\overline{X})
{
case 1: printf ("st "); break;
case 2: printf ("nd "); break;
case 3: printf ("rd "); break;
default: printf ("th "); break;
}
}
if (THIS PRX == STEP)
print\overline{f ("step-");}
else
if (THIS PRX == HALF)
print\overline{f ("half-");}

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```

    THIS_GND = PERSON [KEY PERS [FRST DEX] . PERS_DEX] . GENDER;
    switch (PRIM REL)
    {
        case PARENT : if (THIS_GND = MALE) printf ("father");
                        else printf ("mother");
                    break;
    case CHILD : if (THIS_GND == MALE) printf ("son");
                    else printf ("daughter");
                    break;
    case SPOUSE : if (THIS GND == MALE) printf ("husband");
                    else printf ("wife");
                        break;
    case SIBLING: if (THIS_GND == MALE) printf ("brother");
                        else printf ("sister");
                break;
    case UNCLE : if (THIS_GND == MALE) printf ("uncle");
                                else printf ("aunt");
                                break;
    case NEPHEW : if (THIS_GND == MALE) printf ("nephew");
                    else printf ("niece");
                    break;
    case COUSIN : printf ("cousin");
                break;
    default : printf ("null");
                        break;
    }
    if (INLAW)
    printf ("-in-law");
    if (PRIM REL == COUSIN && THIS_GAP > 0)
    if (TH゙IS GAP > 1)
        print\overline{f}(" %1d times removed", THIS_GAP);
    else
        printf (" once removed");
    printf (" of\n");
} /* end of SHOW_REL */

```
```

/* procedures under FIND REL */
CMPT GNS (INDEX1, INDEX2)
/* CMPT GNS assumes that each ancestor contributes
half of the genetic material to a PERSON. It finds common
ancestors between two PERSONs and computes the expected
value of the PROPORTN of common material. */
register INDX_TYP INDEX1, INDEX2;
{ float COM_PROP;
/* First zero out all ancestors to allow adding. This is necessary
because there might be two paths to an ancestor. */
ZERO_PRO (INDEX1);
/* now mark with shared PROPORTN */
MARK PRO (PERSON [INDEX1] . IDENT, 1.0, INDEX1);
COM P
CHK_COM ( \& COM_PROP, PERSON [INDEX1] . IDENT, 1.0, 0.0, INDEX2);
printf (" Proportion of common genetic material = %1.5e \n",
COM PROP);
} /* end of CM\overline{PT}GNS */
ZERO PRO (ZERO DEX)
/* ZERO PRO recursively seeks out all ancestors and
zeros them out. */
register INDX TYP ZERO_DEX;
{ register NBR_PTR THIS_NBR;
PERSON [ZERO_DEX] . DSC_GENE = 0.0;
for (THIS NB\overline{R}= PERSON [ZERO_DEX] - NBR_HDR;
THIS NBR != NULL;
THIS_NBR = THIS_NBR }->\mathrm{ NEXT_NBR)
{
if (THIS NBR -> NBR EDGE == PARENT)
ZERO_\overline{PRO (THIS_NBR M NBR_DEX);}
}
} /* end of ZERO PRO */

```
```

MARK PRO (MARKER, PROPORTN, MARK DEX)
/* MARK PRO recursively seeks out all ancestors and
marks them with the sender's PROPORTN of shared
genetic material. This PROPORTN is diluted by one-half
for each generation. */

```

ID TYPE
f1ōat INDX TYP
\{ register NBR_PTR THIS_NBR;
strcpy (PERSON [MARK DEX] . DSC ID, MARKER);
PERSON [MARK DEX] - DSC GENE += PROPORTN;
for (THIS_NB \(\bar{R}=\) PERSON [MARK_DEX] - NBR_HDR ;
THIS_NBR ! = NULL;
THIS_NBR \(=\) THIS_NBR \(\rightarrow\) NEXT_NBR)
\{
if (THIS NBR \(\rightarrow\) NBR EDGE \(==\) PARENT) MARK \(\overline{\text { PRO }}\) (MARKER, PROPORTN / 2.0, THIS_NBR \(\rightarrow\) NBR_DEX);
\}
\} /* end of MARK PRO */
CHK COM (COM PTR, MATCH ID, PROPORTN, COUNTED, CHK DEX)
\(/^{*}\) CHK COM searches \(a \overline{11}\) the ancestors of CHK DEX to see if any have been marked, and if so add \(\bar{s}\) the appropriate amount to *COM_PTR. */
float *COM PTR, PROPORTN, COUNTED;
ID TYPE
INDX TYP
\{ register NBR PTR THIS NBR; register flōat CONTR̄IB;
if (STREQ (PERSON [CHK DEX] . DSC ID, MATCH ID))
\{ /* Increment *COM PTR by the contribution of
this common ancestor, but discount for the contribution of less remote ancestors already counted. */ CONTRIB = PERSON [CHK DEX] . DSC GENE * PROPORTN; *COM PTR += CONTRIB - COUNTED;
\}
else
CONTRIB \(=0.0\);
for (THIS NBR \(=\) PERSON [CHK DEX] - NBR_HDR;
THIS \({ }^{-}\)NBR \(!=\)NULL;
THIS NBR \(=\) THIS NBR \(\rightarrow\) NEXT NBR)
\{
if (THIS NBR \(\rightarrow\) NBR EDGE \(==\) PARENT)
CHK COM (COM PTR, MATCH ID, PROPORTN / 2.0, CONTRIB / 4.0, THIS NBR \(\rightarrow\) NBR DEX);
\}
\} \(/ *\) end of CHK_COM */
```

5.0 COBOL
In keeping with the general convention of the examples, language-supplied
keywords and identifiers are written in lower case in the program. To conform
strictly to the COBOL-74 standard, however, programs must use only upper-case
letters.
*
Compilation unit number 1 ----
identification division.
program-id. RELATE.
environment division.
configuration section.
source-computer. VAX-11.
object-computer. VAX-11.
input-output section.
file-control.
select PEOPLE assign to "PEOPLE.DAT",
file status is PEOPLE-STATUS.
data division.
file section.
fd PEOPLE
label records are standard.
01 PEOPLE-RECORD.
05 NAME pic X(20).
05 IDENTIFIER pic 999.
*** "M" for MALE and "F" for FEMALE
0 5 ~ G E N D E R ~ p i c ~ X .
05 IMMEDIATE-RELATIONS.
10 RELATIVE-IDENTIFIER occurs 3 times pic 999.
working-storage section.
77 ARG-PERSON1-INDEX pic 999.
77 ARG-PERSON2-INDEX pic 999.
01 PEOPLE-STATUS.
05 STATUS-1 pic X.
88 END-OF-PEOPLE-FILE value "1".
05 STATUS-2 pic X.

* Define global objects
01 TRUTH-VALUES.
05 IS-TRUE pic X value "T".
05 IS-FALSE pic X value "F".
01 SPECIAL-IDENT-VALUE.
05 NULL-IDENT pic 999 value 000.

```
```

* each PERSON's record in the file identifies at most three
* others directly related: father, mother, and spouse
01 GIVEN-IDENTIFIERS.
0 5 ~ F A T H E R - I D E N T ~ p i c ~ 9 ~ v a l u e ~ 1 . ~
0 5 ~ M O T H E R - I D E N T ~ p i c ~ 9 ~ v a l u e ~ 2 . ~
05 SPOUSE-IDENT pic 9 value 3.
01 GENDER-TYPE.
05 MALE pic X value "M".
05 FEMALE pic X value "F".
01 RELATION-TYPE.
05 PARENT pic 9 value 1.
05 CHILD
0 5 ~ S P O U S E ~
05 SIBLING
05 UNCLE
0 5 ~ N E ~ P H E W ~
05 COUSIN
05 NULL-RELATION
pic 9 value 2.
pic 9 value 3.
pic 9 value 4.
pic 9 value 5.
pic }9\mathrm{ value 6.
pic 9 value 7.
pic 9 value 8.
    * A node in the graph (= PERSON) has either already been reached,
* is immediately adjacent to those reached, or farther away.
O1 REACHED-TYPE.
05 REACHED
05 NEARBY
0 5 ~ N O T - S E E N
pic }9\mathrm{ value 1.
pic 9 value 2.
pic 9 value 3.

```
```

* the PERSON array is the central repository of information
* about inter-relationships.
* All relationships are captured in the directed graph of which
* each record is a node.
01 PERSON-TABLE.
05 NUMBER-OF-PERSONS usage index.
05 PERSON occurs 300 times
indexed by CURRENT, PREVIOUS,
FROM-INDEX, TO-INDEX,
PERSON1-INDEX, PERSON2-INDEX.
*** static information - filled from PEOPLE file:
10 NAME pic X(20).
10 IDENTIFIER pic 999.
10 GENDER pic X.
*** IDENTIFIERs of immediate relatives - father, mother, spouse
10 IMMEDIATE-RELATIONS.
15 RELATIVE-IDENTIFIER occurs 3 times indexed by RELATIONSHIP
pic 999.
*** pointers to immediate neighbors in graph
10 NEIGHBOR-COUNT pic 99.
10 NEIGHBOR-RECORD occurs 20 times indexed by NEXT-NEIGHBOR.
15 NEIGHBOR-INDEX usage index.
15 NEIGHBOR-EDGE pic 9.
*** data used when traversing graph to resolve user request:
10 DISTANCE-FROM-SOURCE pic 99999V9.
10 PATH-PREDECESSOR usage index.
10 EDGE-TO-PREDECESSOR pic 9.
10 REACHED-STATUS : pic 9.
*** data used to compute common genetic material
10 DESCENDANT-IDENTIFIER pic 999.
10 DESCENDANT-GENES pic 9V99999999.
* These variables are used to accept and resolve requests for
* RELATIONSHIP information.
01 RELATIONSHIP-WORK-ITEMS.
05 REQUEST-BUFFER pic X(60).
88 REQUEST-TO-STOP value "stop".
0 5 ~ P E R S O N 1 - I D E N T ~ p i c ~ X ( 2 0 ) . ~
05 PERSON2-IDENT pic X(20).
05 PERSON1-FOUND pic 999.
05 PERSON2-FOUND pic 999.
0 5 ~ E R R O R - M E S S A G E ~ p i c ~ X ( 4 0 ) . ~
05 REQUEST-OK
pic X(40) value "Request OK".
01 AUXILIARY-VARIABLES.

| 05 | RELATION-LOOP-DONE | pic X. |
| :--- | :--- | :--- |
| 08 RELATION-LOOP-IS-DONE | value "T". |  |
| 05 | TEMP-INDEX | usage index. |
| 05 | THIS-EDGE | pic 9. |
| 05 | LEADING-SPACES | pic 99. |
| 05 | SEMICOLON-COUNT | pic 99. |
| 05 | CURRENT-IDENT | pic 999. |
| 05 | PREVIOUS-IDENT | pic 999. |
| 05 | TEMP-IDENT | pic X(20). |

```
```

procedure division.
MAIN-LINE.
open input PEOPLE.
read PEOPLE at end perform NULL.

* This loop reads in the PEOPLE file and constructs the PERSON
* array from it (one PERSON = one record = one array entry).
* As records are read in, links are constructed to represent the
* PARENT-CHILD or SPOUSE RELATIONSHIP. The array then implements
* a directed graph which is used to satisfy subsequent user
* requests. The file is assumed to be correct - no validation
* is performed on it.
perform READ-IN-PEOPLE thru READ-IN-PEOPLE-EXIT
varying CURRENT from 1 by 1 until END-OF-PEOPLE-FILE.
set CURRENT down by 1.
set NUMBER-OF-PERSONS to CURRENT.
close PEOPLE.
* PERSON array is now loaded and edges between immediate relatives
* (PARENT-CHILD or SPOUSE-SPOUSE) are established.
perform PROMPT-AND-READ.
* While-loop accepts requests and finds RELATIONSHIP (if any)
* between pairs of PERSONs.
perform READ-AND-PROCESS-REQUEST thru READ-AND-PROCESS-REQUEST-EXIT
until REQUEST-TO-STOP.
display " End of relation-finder.".
stop run.
READ-IN-PEOPLE.
*** copy direct information from file to array
move corresponding PEOPLE-RECORD to PERSON (CURRENT).
move IMMEDIATE-RELATIONS of PEOPLE-RECORD
to IMMEDIATE-RELATIONS of PERSON (CURRENT).
*** Location of adjacent persons as yet undetermined
move zero to NEIGHBOR-COUNT of PERSON (CURRENT).
*** Descendants as yet undetermined
move NULL-IDENT to DESCENDANT-IDENTIFIER of PERSON (CURRENT).
move IDENTIFIER of PERSON (CURRENT) to CURRENT-IDENT.
*** Compare this PERSON against all previously entered PERSONs
*** to search for RELATIONSHIPs.
perform COMPARE-TO-PREVIOUS varying PREVIOUS from 1 by 1
until PREVIOUS not < CURRENT.
read PEOPLE at end perform NULL.
READ-IN-PEOPLE-EXIT.
exit.
NULL.
exit.

```
```

    COMPARE-TO-PREVIOUS.
    move IDENTIFIER of PERSON (PREVIOUS) to PREVIOUS-IDENT.
    *** Search for father, mother, or spouse relationship in
*** either direction between this and PREVIOUS PERSON.
*** Assume at most one RELATIONSHIP exists.
move IS-FALSE to RELATION-LOOP-DONE.
perform TRY-ALL-RELATIONSHIPS
varying RELATIONSHIP from FATHER-IDENT by 1
until RELATIONSHIP > SPOUSE-IDENT or RELATION-LOOP-IS-DONE.
TRY-ALL-RELATIONSHIPS.
if RELATIVE-IDENTIFIER of PERSON (CURRENT, RELATIONSHIP) =
PREVIOUS-IDENT
set FROM-INDEX to CURRENT
set TO-INDEX to PREVIOUS
perform LINK-RELATIVES
move IS-TRUE to RELATION-LOOP-DONE
else
if CURRENT-IDENT =
RELATIVE-IDENTIFIER of PERSON (PREVIOUS, RELATIONSHIP)
set FROM-INDEX to PREVIOUS
set TO-INDEX to CURRENT
perform LINK-RELATIVES
move IS-TRUE to RELATION-LOOP-DONE.
LINK-RELATIVES.

* establishes cross-indexing between immediately related PERSONs.
if RELATIONSHIP = SPOUSE-IDENT
move SPOUSE to THIS-EDGE
perform LINK-ONE-WAY
set TEMP-INDEX to FROM-INDEX
set FROM-INDEX to TO-INDEX
set TO-INDEX to TEMP-INDEX
perform LINK-ONE-WAY
else
RELATIONSHIP is father or mother
move PARENT to THIS-EDGE
perform LINK-ONE-WAY
move CHILD to THIS-EDGE
set TEMP-INDEX to FROM-INDEX
set FROM-INDEX to TO-INDEX
set TO-INDEX to TEMP-INDEX
perform LINK-ONE-WAY.
LINK-ONE-WAY.
*** Establishes the NEIGHBOR-RECORD from one PERSON to another
add 1 to NEIGHBOR-COUNT of PERSON (FROM-INDEX).
set NEXT-NEIGHBOR to NEIGHBOR-COUNT of PERSON (FROM-INDEX).
set NEIGHBOR-INDEX of PERSON (FROM-INDEX, NEXT-NEIGHBOR)
to TO-INDEX.
move THIS-EDGE
to NEIGHBOR-EDGE of PERSON (FROM-INDEX, NEXT-NEIGHBOR).

```
```

    PROMPT-AND-READ.
    * Issues prompt for user-request, reads in request,
* blank-fills buffer, and skips to next line of input.

```
```

    display " ".
    ```
    display " ".
    display " -----------------------------------------------------
    display " -----------------------------------------------------
    display " Enter two person-identifiers (name or number),".
    display " Enter two person-identifiers (name or number),".
    display " separated by semicolon. Enter ""stop"" to stop.".
    display " separated by semicolon. Enter ""stop"" to stop.".
    move spaces to REQUEST-BUFFER.
    move spaces to REQUEST-BUFFER.
    accept REQUEST-BUFFER.
    accept REQUEST-BUFFER.
    READ-AND-PROCESS-REQUEST.
    perform CHECK-REQUEST.
*** Syntax check of request completed. Now either display error
*** message or search for the two PERSONs.
    if ERROR-MESSAGE = REQUEST-OK
            perform PROCESS-LEGAL-REQUEST
        else
            display " Incorrect request format: " , ERROR-MESSAGE.
        perform PROMPT-AND-READ.
READ-AND-PROCESS-REQUEST-EXIT.
    exit.
    CHECK-REQUEST.
* Performs syntactic check on request in buffer
* and fills in identifiers of the two requested persons.
    move zero to SEMICOLON-COUNT.
    inspect REQUEST-BUFFER tallying SEMICOLON-COUNT
    for all ";".
    if SEMICOLON-COUNT not = 1
    move "must be exactly one semicolon." to ERROR-MESSAGE
    else
        move zero to LEADING-SPACES
        inspect REQUEST-BUFFER tallying LEADING-SPACES
        for leading spaces
    add 1 to LEADING-SPACES
    unstring REQUEST-BUFFER delimited by ";"
        into PERSON1-IDENT, TEMP-IDENT
        with pointer LEADING-SPACES
    if PERSON1-IDENT = spaces
                move "null field preceding semicolon." to ERROR-MESSAGE
            else
                if TEMP-IDENT = spaces
                move "null field following semicolon." to ERROR-MESSAGE
                else
                    move zero to LEADING-SPACES
                    inspect TEMP-IDENT tallying LEADING-SPACES
                    for leading spaces
            add 1 to LEADING-SPACES
            unstring TEMP-IDENT into PERSON2-IDENT
                    with pointer LEADING-SPACES
            move REQUEST-OK to ERROR-MESSAGE.
```

PROCESS-LEGAL-REQUEST.
*** search for requested PERSONs. move zero to PERSON1-FOUND, PERSON2-FOUND.
perform SCAN-ALL-PERSONS varying CURRENT from 1 by 1
until CURRENT > NUMBER-OF-PERSONS.
if PERSON1-FOUND $=1$ and PERSON2-FOUND $=1$
Exactly one match for each PERSON - proceed to
determine RELATIONSHIP, if any.
if PERSON1-INDEX = PERSON2-INDEX if GENDER of PERSON (PERSON1-INDEX) = MALE
display " ", NAME of PERSON (PERSON1-INDEX), " is identical to himself."
else
display " ", NAME of PERSON (PERSON1-INDEX), " is identical to herself."
else
set ARG-PERSON1-INDEX to PERSON1-INDEX set ARG-PERSON2-INDEX to PERSON2-INDEX call "FINDREL" using

ARG-PERSON1-INDEX, ARG-PERSON2-INDEX, PERSON-TABLE
else
*** either not found or more than one found perform MISSING-OR-DUPLICATE-PERSONS.

SCAN-ALL-PERSONS.
if PERSON1-IDENT = NAME of PERSON (CURRENT) or IDENTIFIER of PERSON (CURRENT)
set PERSON1-INDEX to CURRENT
add 1 to PERSON1-FOUND.
if PERSON2-IDENT = NAME of PERSON (CURRENT) or IDENTIFIER of PERSON (CURRENT)
set PERSON2-INDEX to CURRENT
add 1 to PERSON2-FOUND.

MISSING-OR-DUPLICATE-PERSONS.
if PERSON1-FOUND = zero
display " First person not found."
else
if PERSON1-FOUND > 1
display " Duplicate names for first person - use",
" numeric identifier.".
if PERSON2-FOUND = zero
display " Second person not found."
else
if PERSON2-FOUND > 1
display " Duplicate names for second person - use",
" numeric identifier.".

```
*
    ---- Compilation unit number 2 ----
    identification division.
    program-id. FINDREL.
    Finds shortest path (if any) between two PERSONs and
    determines their RELATIONSHIP based on immediate relations
    traversed in path. PERSON array simulates a directed graph,
    and algorithm finds shortest path, based on following
    weights: PARENT-CHILD edge = 1.0
    SPOUSE-SPOUSE edge = 1.8
    environment division.
    configuration section.
    source-computer. VAX-11.
    object-computer. VAX-11.
    data division.
    working-storage section.
* Define global objects
    01 TRUTH-VALUES.
        05 IS-TRUE pic X value "T".
        05 IS-FALSE pic X value "F".
    * each PERSON`s record in the file identifies at most three
* others directly related: father, mother, and spouse
    01 GIVEN-IDENTIFIERS.
        0 5 ~ F A T H E R - I D E N T ~ p i c ~ 9 ~ v a l u e ~ 1 . ~
        0 5 ~ M O T H E R - I D E N T ~ p i c ~ 9 ~ v a l u e ~ 2 . ~
        05 SPOUSE-IDENT pic 9 value 3.
    01 GENDER-TYPE.
        05 MALE pic X value "M".
        05 FEMALE pic X value "F".
    01 RELATION-TYPE.
        05 PARENT pic 9 value 1.
        0 5 ~ C H I L D ~
        05 SPOUSE
        05 SIBLING
        0 5 ~ U N C L E ~
        0 5 ~ N E ~ P H E W ~
        0 5 ~ C O U S I N
        05 NULL-RELATION
```

```
pic 9 value 2.
```

pic 9 value 2.
pic 9 value 3.
pic 9 value 3.
pic 9 value 4.
pic 9 value 4.
pic 9 value 5.
pic 9 value 5.
pic 9 value 6.
pic 9 value 6.
pic 9 value 7.
pic 9 value 7.
pic 9 value 8.

```
pic 9 value 8.
```

* A node in the graph (= PERSON) has either already been reached, * is immediately adjacent to those reached, or farther away. 01 REACHED-TYPE.

| 05 | REACHED | pic 9 | value 1. |
| :--- | :--- | :--- | :--- | :--- |
| 05 | NEARBY | pic 9 | value 2. |
| 05 | NOT-SEEN | pic 9 | value 3. |

01 SEARCH-TYPE.
05 SEARCHING pic 9 value 1.
05 SUCCEEDED pic 9 value 2.
05 FAILED pic 9 value 3.
01 SIBLING-TYPE.

| 05 | STEP | pic 9 | value 1. |
| :--- | :--- | :--- | :--- |
| 05 | HALF | pic 9 | value 2. |
| 05 | FULL | pic 9 | value 3. |

01 KEY-PERSON-TABLE.
05 KEY-PERSON occurs 300 times
indexed by KEY-INDEX, LATER-KEY-INDEX, PRIMARY-INDEX, FIRST-INDEX, LAST-INDEX, RECEIVE-INDEX, SEND-INDEX.
10 RELATION-TO-NEXT pic 9.
10 PERSON-INDEX usage index.
10 GENERATION-GAP pic 999.
10 PROXIMITY pic 9. 10 COUSIN-RANK pic 999.

01 AUXILIARY-VARIABLES.
*** these variables are used to find the shortest path
05 WEIGHT-THIS-EDGE pic 99V9.
05 DISTANCE-THRU-BASE-NODE pic 99999V9.
05 SEARCH-STATUS pic 9.
05 NEARBY-NODE usage index, occurs 300 times, indexed by THIS-NEARBY-INDEX, BEST-NEARBY-INDEX, LAST-NEARBY-INDEX.
05 THIS-EDGE pic 9.
05 NEXT-BASE-EDGE pic 9.
05 MINIMAL-DISTANCE pic 9999999V9.
05 DISPLAY-BUFFER pic X(70).
05 DISPLAY-POINTER pic 99.
05 NULL-IDENT pic 999 value 000.
*** these variables are used to condense the path
05 KEY-RELATION
pic 9.
05 LATER-KEY-RELATION
pic 9.
05 PRIMARY-RELATION
pic 9.
05 FIRST-RELATION
pic 9.
05 LAST-RELATION
05 NEXT-PR IMARY-RELATION
05 GAP-SIZE
pic 9.
pic 9.
05 ANOTHER PIC 999.
05 ANOTHER-ELEMENT-POSSIBLE pic X. 88 ANOTHER-ELEMENT-IS-POSSIBLE value "T".

```
*** these variables are used to generate KEY-PERSONs and for DISPLAY
    05 GENERATION-COUNT pic 999.
    0 5 ~ T E M P - N U M B E R ~ p i c ~ 9 9 9 . ~
    0 5 ~ T H I S - C O U S I N - R A N K ~ p i c ~ 9 9 9 . ~
    0 5 ~ T H I S - P R O X I M I T Y ~ p i c ~ 9 . ~
    05 THIS-GENDER
    05 THIS-GENERATION-GAP
    05 SUFFIX-INDICATOR
    05 TWO-DIGIT-F IELD
    05 INLAW
    88 RELATION-IS-INLAW
    05 MALE-NAME-VALUES.
        10 filler pic X(8) value "father ".
        10 filler pic X(8) value "son ".
        10 filler pic X(8) value "husband ".
        10 filler pic X(8) value "brother ".
        10 filler pic X(8) value "uncle ".
        10 filler pic X(8) value "nephew ".
        10 filler pic X(8) value "cousin ".
        10 filler pic X(8) value "null ".
    05 MALE-NAME-TABLE redefines MALE-NAME-VALUES.
        10 PRIMARY-MALE-NAME pic X(8) occurs 8 times
        indexed by MALE-INDEX.
    05 FEMALE-NAME-VALUES.
        10 filler pic X(8) value "mother ".
        10 filler pic X(8) value "daughter".
        10 filler pic X(8) value "wife ".
        10 filler pic X(8) value "sister ".
        10 filler pic X(8) value "aunt ".
        10 filler pic X(8) value "niece ".
        10 filler pic X(8) value "cousin ".
        10 filler pic X(8) value "null ".
    05 FEMALE-NAME-TABLE redefines FEMALE-NAME-VALUES.
        10 PRIMARY-FEMALE-NAME pic X(8) occurs 8 times
        indexed by FEMALE-INDEX.
```

1inkage section.

```
77 PARM-TARGET-INDEX pic 999.
77 PARM-SOURCE-INDEX pic 999.
01 PERSON-TABLE.
    05 NUMBER-OF-PERSONS usage index.
    05 PERSON occurs 300 times
            indexed by INDEX1, INDEX2, TARGET-INDEX, SOURCE-INDEX,
                BASE-NODE, THIS-NODE, NEXT-NODE.
*** static information - filled from PEOPLE file:
    10 NAME pic X(20).
    10 IDENTIFIER pic 999.
    10 GENDER pic X.
*** IDENTIFIERs of immediate relatives - father, mother, spouse
    10 IMMEDIATE-RELATIONS.
            15 RELATIVE-IDENTIFIER occurs 3 times indexed by RELATIONSHIP
                                    pic 999.
*** pointers to immediate neighbors in graph
    10 NEIGHBOR-COUNT pic 99.
    10 NEIGHBOR-RECORD occurs 20 times indexed by THIS-NEIGHBOR.
            15 NEIGHBOR-INDEX usage index.
            15 NEIGHBOR-EDGE pic 9.
*** data used when traversing graph to resolve user request:
    10 DISTANCE-FROM-SOURCE pic 99999V9.
    10 PATH-PREDECESSOR usage index.
    10 EDGE-T0-PREDECESSOR pic 9.
    10 REACHED-STATUS pic 9.
*** data used to compute common genetic material
    10 DESCENDANT-IDENTIFIER pic 999.
    10 DESCENDANT-GENES pic 9V99999999.
    procedure division using
                PARM-TARGET-INDEX, PARM-SOURCE-INDEX, PERSON-TABLE.
    MAIN-LINE.
    set TARGET-INDEX to PARM-TARGET-INDEX.
    set SOURCE-INDEX to PARM-SOURCE-INDEX.
*** initialize PERSON-array for processing -
*** mark all nodes as not seen
    perform MARK-AS-NOT-SEEN varying THIS-NODE from l by l
        until THIS-NODE > NUMBER-OF-PERSONS.
    set THIS-NODE to SOURCE-INDEX.
*** mark source node as REACHED
    move REACHED to REACHED-STATUS of PERSON (THIS-NODE).
    move zero to DISTANCE-FROM-SOURCE of PERSON (THIS-NODE).
*** no nearby nodes exist yet
    set LAST-NEARBY-INDEX to 1.
    set LAST-NEARBY-INDEX down by 1.
    if THIS-NODE = TARGET-INDEX
        move SUCCEEDED to SEARCH-STATUS
    else
        move SEARCHING to SEARCH-STATUS.
```

```
*** Loop keeps processing closest-to-source, unREACHED node
***
    until target REACHED, or no more connected nodes.
    perform SEARCH-FOR-TARGET until SEARCH-STATUS not = SEARCHING.
*** Shortest path between PERSONs now established. Next task is
*** to translate path to English description of RELATIONSHIP.
    if SEARCH-STATUS = FAILED
        display " ", NAME of PERSON (TARGET-INDEX), " is not related to ",
                        NAME of PERSON (SOURCE-INDEX)
    else
        success - parse path to find and display RELATIONSHIP
        perform RESOLVE-PATH-TO-ENGLISH
        call "COMGENES" using
            PARM-SOURCE-INDEX, PARM-TARGET-INDEX, PERSON-TABLE.
    END-OF-F INDREL.
        exit program.
    MARK-AS-NOT-SEEN.
    move NOT-SEEN to REACHED-STATUS of PERSON (THIS-NODE).
    SEARCH-FOR-TARGET.
*** Process all nodes adjacent to THIS-NODE
    perform PROCESS-ADJACENT-NODE varying THIS-NEIGHBOR from 1 by 1
        until THIS-NEIGHBOR > NEIGHBOR-COUNT of PERSON (THIS-NODE).
*** All nodes adjacent to THIS-NODE are set. Now search for
*** shortest-distance unREACHED (but NEARBY) node to process next.
    if LAST-NEARBY-INDEX = zero
        move FAILED to SEARCH-STATUS
        else
*** determine next node to process
        move 9999999 to MINIMAL-DISTANCE
        perform FIND-CLOSEST-UNREACHED-NODE varying THIS-NEARBY-INDEX
            from 1 by 1 until THIS-NEARBY-INDEX > LAST-NEARBY-INDEX
*** establish new THIS-NODE
        set THIS-NODE to NEARBY-NODE (BEST-NEARBY-INDEX)
*** change THIS-NODE from being NEARBY to REACHED
        move REACHED to REACHED-STATUS of PERSON (THIS-NODE)
*** remove THIS-NODE from NEARBY list
    set NEARBY-NODE (BEST-NEARBY-INDEX) to NEARBY-NODE (LAST-NEARBY-INDEX)
    set LAST-NEARBY-INDEX down by 1
    if THIS-NODE = TARGET-INDEX
                move SUCCEEDED to SEARCH-STATUS.
```

```
PROCESS-ADJACENT-NODE.
    set BASE-NODE to THIS-NODE.
    set NEXT-NODE to NEIGHBOR-INDEX of PERSON (BASE-NODE, THIS-NEIGHBOR).
    move NEIGHBOR-EDGE of PERSON (BASE-NODE, THIS-NEIGHBOR)
    to NEXT-BASE-EDGE.
    NEXT-NODE is adjacent to last-REACHED node (= BASE-NODE).
    if NEXT-NODE already REACHED, do nothing.
    If previously seen, check whether path thru BASE-NODE is
    shorter than current path to NEXT-NODE, and if so re-link
    next to base.
    If not previously seen, link next to base node.
    if NEXT-BASE-EDGE = SPOUSE
    move 1.8 to WEIGHT-THIS-EDGE
    else
        move 1.0 to WEIGHT-THIS-EDGE.
    if REACHED-STATUS of PERSON (NEXT-NODE) not = REACHED
    add WEIGHT-THIS-EDGE, DISTANCE-FROM-SOURCE of PERSON (BASE-NODE)
        giving DISTANCE-THRU-BASE-NODE
    if REACHED-STATUS of PERSON (NEXT-NODE) = NOT-SEEN
        move NEARBY to REACHED-STATUS of PERSON (NEXT-NODE)
        set LAST-NEARBY-INDEX up by 1
        set NEARBY-NODE (LAST-NEARBY-INDEX) to NEXT-NODE
        perform LINK-NEXT-NODE-TO-BASE-NODE
    else
        REACHED-STATUS = NEARBY
        if DISTANCE-THRU-BASE-NODE
                            < DISTANCE-FROM-SOURCE of PERSON (NEXT-NODE)
                perform LINK-NEXT-NODE-TO-BASE-NODE.
    LINK-NEXT-NODE-TO-BASE-NODE.
*** link next to base by re-setting its predecessor index to
*** point to base, note type of edge, and re-set distance
*** as it is through base node.
    move DISTANCE-THRU-BASE-NODE
        to DISTANCE-FROM-SOURCE of PERSON (NEXT-NODE).
    set PATH-PREDECESSOR of PERSON (NEXT-NODE) to BASE-NODE.
    move NEXT-BASE-EDGE to EDGE-TO-PREDECESSOR of PERSON (NEXT-NODE).
FIND-CLOSEST-UNREACHED-NODE.
    set NEXT-NODE to NEARBY-NODE (THIS-NEARBY-INDEX).
    if DISTANCE-FROM-SOURCE of PERSON (NEXT-NODE) < MINIMAL-DISTANCE
        set BEST-NEARBY-INDEX to THIS-NEARBY-INDEX
        move DISTANCE-FROM-SOURCE of PERSON (NEXT-NODE) to MINIMAL-DISTANCE.
```

```
    RESOLVE-PATH-TO-ENGLISH.
*** RESOLVE-PATH-TO-ENGLISH condenses the shortest path to a
*** series of RELATIONSHIPs for which there are English
*** descriptions.
*** Key persons are the ones in the RELATIONSHIP path which remain
*** after the path is condensed.
    display ".Shortest path between identified persons: ".
    set THIS-NODE to TARGET-INDEX.
*** Display path and initialize KEY-PERSON array from path elements.
    perform TRAVERSE-SHORTEST-PATH varying KEY-INDEX from 1 by 1
    until THIS-NODE = SOURCE-INDEX.
    display " ", NAME of PERSON (THIS-NODE).
    set PERSON-INDEX of KEY-PERSON (KEY-INDEX) to THIS-NODE.
    move NULL-RELATION to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX).
    move NULL-RELATION to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 1).
*** Resolve CHILD-PARENT and CHILD-SPOUSE-PARENT relations
*** to SIBLING relations.
    perform FIND-SIBLINGS varying KEY-INDEX from 1 by 1
        until RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = NULL-RELATION.
*** Resolve CHILD-CHILD-... and PARENT-PARENT-... relations to
*** direct descendant or ancestor relations.
    perform FIND-ANCESTORS-OR-DESCENDANTS varying KEY-INDEX from 1 by 1
    until RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = NULL-RELATION.
*** Resolve CHILD-SIBLING-PARENT to COUSIN,
***
***
        CHILD-SIBLING to NEPHEW,
        SIBLING-PARENT to UNCLE.
    perform FIND-COUSINS-NEPHEWS-UNCLES varying KEY-INDEX from 1 by 1
        until RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = NULL-RELATION.
```


## 夫夫*

## ***

Loop below will pick out valid adjacent strings of elements
to be displayed. KEY-INDEX points to first element, LATER-KEY-INDEX to last element, and PRIMARY-INDEX to the element which determines the primary English word to be used. Associativity of adjacent elements in condensed table
is based on English usage.
set KEY-INDEX to 1.
display " Condensed path:".
perform CONSOLIDATE-ADJACENT-PERSONS
unti1 RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = NULL-RELATION set THIS-NODE to PERSON-INDEX of KEY-PERSON (KEY-INDEX).
display " ", NAME of PERSON (THIS-NODE).

```
TRAVERSE-SHORTEST-PATH.
    set PERSON-INDEX of KEY-PERSON (KEY-INDEX) to THIS-NODE.
    move FULL to PROXIMITY of KEY-PERSON (KEY-INDEX).
    move EDGE-TO-PREDECESSOR of PERSON (THIS-NODE)
            to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX).
    if EDGE-TO-PREDECESSOR of PERSON (THIS-NODE) = SPOUSE
        move zero to GENERATION-GAP of KEY-PERSON (KEY-INDEX)
        display " ", NAME of PERSON (THIS-NODE), " is spouse of"
    else
        move 1 to GENERATION-GAP of KEY-PERSON (KEY-INDEX)
        if EDGE-TO-PREDECESSOR of PERSON (THIS-NODE) = PARENT
            display " ", NAME of PERSON (THIS-NODE), " is parent of"
        else
            edge is child-type
            display " ", NAME of PERSON (THIS-NODE), " is child of".
    set THIS-NODE to PATH-PREDECESSOR of PERSON (THIS-NODE).
    FIND-SIBLINGS.
    if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = CHILD
        move RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 1)
                to LATER-KEY-RELATION
    if LATER-KEY-RELATION = PARENT
*** then found either full or half SIBLINGs
            perform SET-UP-FULL-HALF-SIBLING
        else
            if LATER-KEY-RELATION = SPOUSE and
                    RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 2) = PARENT
****
            then found step-siblings
                move zero to GENERATION-GAP of KEY-PERSON (KEY-INDEX)
                    move STEP to PROXIMITY of KEY-PERSON (KEY-INDEX)
                    move SIBLING to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX)
                    move 2 to GAP-SIZE
                    perform CONDENSE-KEY-PERSONS.
    SET-UP-FULL-HALF-SIBLING.
*** Determines whether two PERSONs are full siblings, i.e.,
*** have the same two parents.
    set INDEX1 to PERSON-INDEX of KEY-PERSON (KEY-INDEX).
    set INDEX2 to PERSON-INDEX of KEY-PERSON (KEY-INDEX + 2).
    if (NULL-IDENT not =
                    RELATIVE-IDENTIFIER of PERSON (INDEX1, FATHER-IDENT)
            and RELATIVE-IDENTIFIER of PERSON (INDEX1, MOTHER-IDENT))
            and (RELATIVE-IDENTIFIER of PERSON (INDEX1, FATHER-IDENT) =
                RELATIVE-IDENTIFIER of PERSON (INDEX2, FATHER-IDENT))
            and (RELATIVE-IDENTIFIER of PERSON (INDEX1, MOTHER-IDENT) =
                RELATIVE-IDENTIFIER of PERSON (INDEX2, MOTHER-IDENT))
        move FULL to PROXIMITY of KEY-PERSON (KEY-INDEX)
    else
            move HALF to PROXIMITY of KEY-PERSON (KEY-INDEX).
    move zero to GENERATION-GAP of KEY-PERSON (KEY-INDEX).
    move SIBLING to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX).
    move 1 to GAP-SIZE.
    perform CONDENSE-KEY-PERSONS.
```

```
FIND-ANCESTORS-OR-DESCENDANTS.
    if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = CHILD or PARENT
        perform NULL varying LATER-KEY-INDEX from KEY-INDEX by l
            until RELATION-TO-NEXT of KEY-PERSON (LATER-KEY-INDEX) not =
                    RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX)
    set GENERATION-COUNT to LATER-KEY-INDEX
    set TEMP-NUMBER to KEY-INDEX
    subtract TEMP-NUMBER from GENERATION-COUNT
    if GENERATION-COUNT > 1
            compress generations
            move GENERATION-COUNT to GENERATION-GAP of KEY-PERSON (KEY-INDEX)
            subtract l from GENERATION-COUNT giving GAP-SIZE
            perform CONDENSE-KEY-PERSONS.
    FIND-COUSINS-NE PHEWS-UNCLES.
    move RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 1)
        to LATER-KEY-RELATION
    if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = CHILD and
        LATER-KEY-RELATION = SIBLING
*** then COUSIN or NEPHEW
        if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 2) = PARENT
            perform FOUND-COUSIN
        else
            found NEPHEW
            move PROXIMITY of KEY-PERSON (KEY-INDEX + 1) to
                    PROXIMITY of KEY-PERSON (KEY-INDEX)
            move NEPHEW to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX)
            move 1 to GAP-SIZE
            perform CONDENSE-KEY-PERSONS
    else
        if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) = SIBLING and
            LATER-KEY-RELATION = PARENT
*** then found UNCLE
            move GENERATION-GAP of KEY-PERSON (KEY-INDEX + 1) to
                    GENERATION-GAP of KEY-PERSON (KEY-INDEX)
            move UNCLE to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX)
            move 1 to GAP-SIZE
            perform CONDENSE-KEY-PERSONS.
FOUND-COUSIN.
    if GENERATION-GAP of KEY-PERSON (KEY-INDEX)
            < GENERATION-GAP of KEY-PERSON (KEY-INDEX + 2)
        move GENERATION-GAP of KEY-PERSON (KEY-INDEX)
            to COUSIN-RANK of KEY-PERSON (KEY-INDEX)
    else
        move GENERATION-GAP of KEY-PERSON (KEY-INDEX + 2)
            to COUSIN-RANK of KEY-PERSON (KEY-INDEX).
*** subtract moves in absolute value since GENERATION-GAP is unsigned
    subtract GENERATION-GAP of KEY-PERSON (KEY-INDEX + 2)
        from GENERATION-GAP of KEY-PERSON (KEY-INDEX).
    move PROXIMITY of KEY-PERSON (KEY-INDEX + 1)
            to PROXIMITY of KEY-PERSON (KEY-INDEX).
    move COUSIN to RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX).
    move 2 to GAP-SIZE.
    perform CONDENSE-KEY-PERSONS.
NULL.
    exit.
```

```
    CONDENSE-KEY-PERSONS.
*** CONDENSE-KEY-PERSONS condenses superfluous entries from the
*** KEY-PERSON array, starting at KEY-INDEX.
    set RECEIVE-INDEX to KEY-INDEX.
    set RECEIVE-INDEX up by 1.
    set SEND-INDEX to RECEIVE-INDEX.
    set SEND-INDEX up by GAP-SIZE.
    perform SLIDE-IT-DOWN varying RECEIVE-INDEX from RECEIVE-INDEX by 1
        until RELATION-TO-NEXT of KEY-PERSON (RECEIVE-INDEX - 1)
            = NULL-RELATION.
SLIDE-IT-DOWN.
    move KEY-PERSON (SEND-INDEX) to KEY-PERSON (RECEIVE-INDEX).
    set SEND-INDEX up by 1.
CONSOLIDATE-ADJACENT-PERSONS.
    move RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX) to KEY-RELATION.
    set LATER-KEY-INDEX, PRIMARY-INDEX to KEY-INDEX.
    if RELATION-TO-NEXT of KEY-PERSON (KEY-INDEX + 1) not = NULL-RELATION
    perform SEEK-MULTI-ELEMENT-COMBINATION.
    set FIRST-INDEX to KEY-INDEX.
    set LAST-INDEX to LATER-KEY-INDEX.
    perform DISPLAY-RELATION.
    set KEY-INDEX to LATER-KEY-INDEX.
    set KEY-INDEX up by 1.
SEEK-MULTI-ELEMENT-COMBINATION.
    move IS-TRUE to ANOTHER-ELEMENT-POSSIBLE.
    if KEY-RELATION = SPOUSE
        set LATER-KEY-INDEX up by 1
        set PRIMARY-INDEX up by 1
        if RELATION-TO-NEXT of KEY-PERSON (LATER-KEY-INDEX)
            = SIBLING or COUSIN
    then nothing can follow spouse-sibling or spouse-cousin
        move IS-FALSE to ANOTHER-ELEMENT-POSSIBLE.
*** PRIMARY-INDEX is now correctly set. Next if-statement
***
    determines if a following SPOUSE relation should be
    appended to this combination or left for the next
    combination.
    if RELATION-TO-NEXT of KEY-PERSON (PRIMARY-INDEX + 1) = SPOUSE
        and ANOTHER-ELEMENT-IS-POSSIBLE
    Only a SPOUSE can follow a Primary
    check primary preceding and following SPOUSE.
    move RELATION-TO-NEXT of KEY-PERSON (PRIMARY-INDEX)
        to PRIMARY-RELATION
    move RELATION-TO-NEXT of KEY-PERSON (PRIMARY-INDEX + 2)
        to NEXT-PRIMARY-RELATION
    if (NEXT-PRIMARY-RELATION = NEPHEW or COUSIN or NULL-RELATION)
        or (PRIMARY-RELATION = NEPHEW)
        or ( (PRIMARY-RELATION = SIBLING or PARENT)
                    and NEXT-PRIMARY-RELATION not = UNCLE )
        then append following SPOUSE with this combination.
        set LATER-KEY-INDEX up by 1.
```

```
    DISPLAY-RELATION.
*** DISPLAY-RELATION takes 1, 2, or 3 adjacent elements in the
*** condensed table and generates the English description of
*** the relation between the first and last + 1 elements.
    move RELATION-TO-NEXT of KEY-PERSON (FIRST-INDEX)
    to FIRST-RELATION.
    move RELATION-TO-NEXT of KEY-PERSON (LAST-INDEX)
        to LAST-RELATION.
    move RELATION-TO-NEXT of KEY-PERSON (PRIMARY-INDEX)
        to PRIMARY-RELATION.
*** set THIS-PROXIMITY
    if (PRIMARY-RELATION = PARENT and FIRST-RELATION = SPOUSE) or
        (PRIMARY-RELATION = CHILD and LAST-RELATION = SPOUSE)
        move STEP to THIS-PROXIMITY
    else
        if PRIMARY-RELATION = SIBLING or UNCLE or NEPHEW or COUSIN
            move PROXIMITY of KEY-PERSON (PRIMARY-INDEX) to THIS-PROXIMITY
        else
            move FULL to THIS-PROXIMITY.
    set THIS-GENERATION-GAP
    if PRIMARY-RELATION = PARENT or CHILD or UNCLE or NEPHEW or COUSIN
        move GENERATION-GAP of KEY-PERSON (PRIMARY-INDEX)
            to THIS-GENERATION-GAP
    else
        move zero to THIS-GENERATION-GAP.
***
    set INLAW
    if (FIRST-RELATION = SPOUSE) and
        (PRIMARY-RELATION = SIBLING or CHILD or NEPHEW or COUSIN)
        move IS-TRUE to INLAW
    else
        if (LAST-RELATION = SPOUSE) and
            (PRIMARY-RELATION = SIBLING or PARENT or UNCLE or COUSIN)
            move IS-TRUE to INLAW
        else
            move IS-FALSE to INLAW.
*** set THIS-COUSIN-RANK
    if PRIMARY-RELATION = COUSIN
        move COUSIN-RANK of KEY-PERSON (PRIMARY-INDEX) to THIS-COUSIN-RANK
    else
        move zero to THIS-COUSIN-RANK.
```

```
*** parameters are set - now generate display.
set THIS-NODE to PERSON-INDEX of KEY-PERSON (FIRST-INDEX).
move spaces to DISPLAY-BUFFER.
move 1 to DISPLAY-POINTER.
string " ", NAME of PERSON (THIS-NODE), " is "
    delimited by size
    into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
if PRIMARY-RELATION = PARENT or CHILD or UNCLE or NEPHEW
    perform GENERATE-GENERATION-QUALIFIER
else
    if (PRIMARY-RELATION = COUSIN) and (THIS-COUSIN-RANK > 1)
        move THIS-COUSIN-RANK to TWO-DIGIT-FIELD
        string TWO-DIGIT-FIELD delimited by size into DISPLAY-BUFFER
                with pointer DISPLAY-POINTER
        divide THIS-COUSIN-RANK by 10 giving TEMP-NUMBER
                remainder SUFFIX-INDICATOR
        if SUFFIX-INDICATOR = 1
            string "st " delimited by size
                into DISPLAY-BUFFER with pointer DISPLAY-POINTER
        else if SUFFIX-INDICATOR = 2
            string "nd " delimited by size
                into DISPLAY-BUFFER with pointer DISPLAY-POINTER
        else if SUFFIX-INDICATOR = 3
            string "rd " delimited by size
                into DISPLAY-BUFFER with pointer DISPLAY-POINTER
        else
            string "th " delimited by size
                    into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
if THIS-PROXIMITY = STEP
    string "step-" delimited by size
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER
else
    if THIS-PROXIMITY = HALF
        string "half-" delimited by size
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
set THIS-NODE to PERSON-INDEX of KEY-PERSON (FIRST-INDEX).
move GENDER of PERSON (THIS-NODE) to THIS-GENDER.
set MALE-INDEX, FEMALE-INDEX to PRIMARY-RELATION.
if THIS-GENDER = MALE
    string PRIMARY-MALE-NAME (MALE-INDEX) delimited by space
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER
else
    string PRIMARY-FEMALE-NAME (FEMALE-INDEX) delimited by space
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
if RELATION-IS-INLAW
    string "-in-law" delimited by size
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
```

```
    if (PRIMARY-RELATION = COUSIN) and (THIS-GENERATION-GAP > 0)
    if THIS-GENERATION-GAP > 1
        move THIS-GENERATION-GAP to TWO-DIGIT-FIELD
        string " ", TWO-DIGIT-FIELD, " times removed"
            delimited by size
            into DISPLAY-BUFFER with pointer DISPLAY-POINTER
    else
        string " once removed" delimited by size
            into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
    string " of" delimited by size
    into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
display DISPLAY-BUFFER.
GENERATE-GENERATION-QUALIFIER.
    if THIS-GENERATION-GAP not < 3
    string "great" delimited by size
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER
    if THIS-GENERATION-GAP > 3
        subtract 2 from THIS-GENERATION-GAP giving TWO-DIGIT-FIELD
        string "*", TWO-DIGIT-FIELD, "-" delimited by size
            into DISPLAY-BUFFER with pointer DISPLAY-POINTER
    else
        string "-" delimited by size
            into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
    if THIS-GENERATION-GAP not < 2
    string "grand-" delimited by size
        into DISPLAY-BUFFER with pointer DISPLAY-POINTER.
```

```
* ---- Compilation unit number 3 ----
    identification division.
    program-id. COMGENES.
    COMGENES assumes that each ancestor contributes
    half of the genetic material to a PERSON. It finds common
    ancestors between two PERSONs and computes the expected
    value of the PROPORTION of common material.
    environment division.
    configuration section.
    source-computer. VAX-11.
    object-computer. VAX-11.
    data division.
    working-storage section.
    01 RELATION-TYPE.
\begin{tabular}{lllll}
05 & PARENT & pic 9 & value 1. \\
05 & CHILD & pic 9 & value 2. \\
05 & SPOUSE & pic 9 & value 3. \\
05 & SIBLING & pic 9 & value 4. \\
05 & UNCLE & pic 9 & value 5. \\
05 & NEPHEW & pic 9 & value 6. \\
05 & COUSIN & pic 9 & value 7. \\
05 & NULL-RELATION & pic 9 & value 8.
\end{tabular}
    01 AUXILIARY-VARIABLES.
    O5 COMMON-PROPORTION pic 9V 9999999999.
    05 MATCH-IDENTIFIER pic 999.
    05 TEN-DIGIT-FIELD
    pic 9.9999999999.
    01 STACKED-VARIABLES.
*** used to simulate recursion
    0 5 \text { STACK-ENTRY occurs 50 times indexed by STACK-INDEX.}
        10 PROPORTION pic 9V 9999999999.
        10 THIS-CONTRIBUTION pic 9V 9999999999.
        10 ALREADY-COUNTED pic 9V9999999999.
        10 PERSON-INDEX usage index.
        10 NEXT-NEIGHBOR pic 999.
```

linkage section.

```
77 PARM-INDE X1
77 PARM-INDEX2
pic 999.
pic 999.
01 PERSON-TABLE.
    05 NUMBER-OF-PERSONS usage index.
    05 PERSON occurs 300 times indexed by
            INDEX1, INDEX2, THIS-NODE.
*** static information - filled from PEOPLE file:
    10 NAME
    10 IDENTIFIER
    10 GENDER
        pic X(20).
        pic 999.
    pic X.
*** IDENTIFIERs of immediate relatives - father, mother, spouse
    10 IMMEDIATE-RELATIONS.
        15 RELATIVE-IDENTIFIER occurs 3 times indexed by RELATIONSHIP
                            pic 999.
*** pointers to immediate neighbors in graph
    10 NEIGHBOR-COUNT . pic 99.
    10 NEIGHBOR-RECORD occurs 20 times indexed by THIS-NEIGHBOR.
        15 NEIGHBOR-INDEX usage index.
        15 NEIGHBOR-EDGE pic 9.
*** data used when traversing graph to resolve user request:
    10 DISTANCE-FROM-SOURCE pic 99999V9.
    10 PATH-PREDECESSOR usage index.
    10 EDGE-TO-PREDECESSOR pic 9.
    10 REACHED-STATUS pic 9.
*** data used to compute common genetic material
    10 DESCENDANT-IDENTIFIER pic 999.
    10 DESCENDANT-GENES pic 9V99999999.
```

```
    procedure division using
                            PARM-INDEX1, PÁRM-INDEX2, PERSON-TABLE.
MAIN-LINE.
    set INDEX1 to PARM-INDEX1.
    set INDEX2 to PARM-INDEX2.
*** First zero out all ancestors to allow adding. This is necessary
*** because there might be two paths to an ancestor.
    set STACK-INDEX to 1.
    set PERSON-INDEX (STACK-INDEX) to INDEX1.
    move zero to NEXT-NEIGHBOR (STACK-INDEX).
    perform ZERO-PROPORTION until STACK-INDEX < 1.
*** now mark with shared PROPORTION
    move IDENTIFIER of PERSON (INDEXI) to MATCH-IDENTIFIER.
    set STACK-INDEX to 1.
    set PERSON-INDEX (STACK-INDEX) to INDEX1.
    move zero to NEXT-NEIGHBOR (STACK-INDEX).
    move 1.0 to PROPORTION (STACK-INDEX).
    perform MARK-PROPORTION until STACK-INDEX < 1.
*** traverse ancestor tree for INDEX2, summing overlap
*** with marked tree of INDEX1
    move zero to COMMON-PROPORTION
    set STACK-INDEX to 1.
    set PERSON-INDEX (STACK-INDEX) to INDEX2.
    move IDENTIFIER of PERSON (INDEX1) to MATCH-IDENTIFIER.
    move zero to NEXT-NEIGHBOR (STACK-INDEX).
    move 1.0 to PROPORTION (STACK-INDEX).
    move zero to ALREADY-COUNTED (STACK-INDEX).
    perform CHECK-COMMON-PROPORTION until STACK-INDEX < 1.
    move COMMON-PROPORTION to TEN-DIGIT-FIELD.
    display " Proportion of common genetic material = ", TEN-DIGIT-FIELD.
    END-OF-COMGENES.
    exit program.
    ZERO-PROPORTION.
*** ZERO-PROPORTION recursively seeks out all ancestors and
*** zeros them out.
    set THIS-NODE to PERSON-INDEX (STACK-INDEX).
    if NEXT-NEIGHBOR (STACK-INDEX) = zero
        move zero to DESCENDANT-GENES of PERSON (THIS-NODE)
        move 1 to NEXT-NEIGHBOR (STACK-INDEX).
    perform NULL
        varying THIS-NEIGHBOR from NEXT-NEIGHBOR (STACK-INDEX) by 1
        until THIS-NEIGHBOR > NEIGHBOR-COUNT (THIS-NODE)
            or NEIGHBOR-EDGE (THIS-NODE, THIS-NEIGHBOR) = PARENT.
    if THIS-NEIGHBOR > NEIGHBOR-COUNT (THIS-NODE)
    then no more ancestors
        set STACK-INDEX down by 1
    else
        set up for next ancestor
        set NEXT-NEIGHBOR (STACK-INDEX) to THIS-NEIGHBOR
        add 1 to NEXT-NEIGHBOR (STACK-INDEX)
        set STACK-INDEX up by 1
        set PERSON-INDEX (STACK-INDEX)
            to NEIGHBOR-INDEX (THIS-NODE, THIS-NEIGHBOR)
        move zero to NEXT-NEIGHBOR (STACK-INDEX).
```

```
    MARK-PROPORTION.
*** MARK-PROPORTION recursively seeks out all ancestors and
*** marks them with the sender's PROPORTION of shared
*** genetic material. This PROPORTION is diluted by one-half
*** for each generation.
set THIS-NODE to PERSON-INDEX (STACK-INDEX).
if NEXT-NEIGHBOR (STACK-INDEX) = zero
    move MATCH-IDENTIFIER
            to DESCENDANT-IDENTIFIER of PERSON (THIS-NODE)
    add PROPORTION (STACK-INDEX)
        to DESCENDANT-GENES of PERSON (THIS-NODE)
        move 1 to NEXT-NEIGHBOR (STACK-INDEX).
perform NULL
    varying THIS-NEIGHBOR from NEXT-NEIGHBOR (STACK-INDEX) by 1
    until THIS-NEIGHBOR > NEIGHBOR-COUNT (THIS-NODE)
            or NEIGHBOR-EDGE (THIS-NODE, THIS-NEIGHBOR) = PARENT.
if THIS-NEIGHBOR > NEIGHBOR-COUNT (THIS-NODE)
then no more ancestors
    set STACK-INDEX down by 1
else
    set up for next ancestor
    set NEXT-NEIGHBOR (STACK-INDEX) to THIS-NEIGHBOR
    add 1 to NEXT-NEIGHBOR (STACK-INDEX)
    set STACK-INDEX up by 1
    set PERSON-INDEX (STACK-INDEX)
        to NEIGHBOR-INDEX (THIS-NODE, THIS-NEIGHBOR)
    move zero to NEXT-NEIGHBOR (STACK-INDEX)
    divide PROPORTION (STACK-INDEX - 1) by 2 giving
        PROPORTION (STACK-INDEX).
```

CHECK-COMMON-PROPORTION.
*** CHECK-COMMON-PROPORTION searches all the ancestors of
*** CHECK-INDEX to see if any have been marked, and if so
*** adds the appropriate amount to COMMON-PROPORTION.
set THIS-NODE to PERSON-INDEX (STACK-INDEX).
if NEXT-NEIGHBOR (STACK-INDEX) = zero
move 1 to NEXT-NEIGHBOR (STACK-INDEX)
if DESCENDANT-IDENTIFIER of PERSON (THIS-NODE) = MATCH-IDENTIFIER
***
***
*** Increment COMMON-PROPORTION by the contribution of this common ancestor, but discount for the contribution of less remote ancestors already counted.
multiply DESCENDANT-GENES of PERSON (THIS-NODE)
by PROPORTION (STACK-INDEX)
giving THIS-CONTRIBUTION (STACK-INDEX)
compute COMMON-PROPORTION $=$ COMMON-PROPORTION

+ THIS-CONTRIBUTION (STACK-INDEX)
- ALREADY-COUNTED (STACK-INDEX) else
move zero to THIS-CONTRIBUTION (STACK-INDEX).
perform NULL
varying THIS-NEIGHBOR from NEXT-NEIGHBOR (STACK-INDEX) by 1 until THIS-NEIGHBOR $>$ NEIGHBOR-COUNT (THIS-NODE)
or NEIGHBOR-EDGE (THIS-NODE, THIS-NEIGHBOR) $=$ PARENT.
if THIS-NEIGHBOR > NEIGHBOR-COUNT (THIS-NODE)
then no more ancestors
set STACK-INDEX down by 1
else
set up for next ancestor
set NEXT-NEIGHBOR (STACK-INDEX) to THIS-NEIGHBOR
add 1 to NEXT-NEIGHBOR (STACK-INDEX)
set STACK-INDEX up by 1
set PERSON-INDEX (STACK-INDEX)
to NEIGHBOR-INDEX (THIS-NODE, THIS-NEIGHBOR)
move zero to NEXT-NEIGHBOR (STACK-INDEX)
divide PROPORTION (STACK-INDEX - 1) by 2 giving
PROPORTION (STACK-INDEX)
divide THIS-CONTRIBUTION (STACK-INDEX - 1) by 4 giving
ALREADY-COUNTED (STACK-INDEX).

NULL.
exit.

### 6.0 FORTRAN

In keeping with the general convention of the examples, language-supplied keywords and identifiers are written in lower case in the program. To conform strictly to the FORTRAN standard, however, programs must use only upper-case letters.

```
    program RELATE
c Establish global constants
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
character NULLID*(IDLEN)
parameter (NULLID = '000')
c Each PERSON's record in the file identifies at most three
c others directly related: father, mother, and spouse
    integer FATHID, MOTHID, SPOUID
    parameter (FATHID = 1, MOTHID = 2, SPOUID = 3)
    character REQOK*10, REQSTP*4
    parameter (REQOK = 'Request OK', REQSTP = 'stop`)
    character MALE*1, FEMALE*1
    parameter (MALE = 'M`, FEMALE = ' F')
    integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NE PHEW, COUSIN, NULLRL
    parameter (PARENT = 1, CHILD = 2, SPOUSE = 3, SIBLNG = 4,
1 UNCLE = 5, NEPHEW = 6, COUSIN = 7, NULLRL = 8)
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
```

c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file

| character*(NAMLEN) | NAME | (MAXPRS) |
| :--- | :--- | :--- |
| character*(IDLEN) | IDENT | (MAXPRS) |
| character*1 | GENDER | (MAXPRS) |

c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS) integer $\quad$ NBRDEX (MAXPRS, MAXNBR) integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request: real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material character*(IDLEN) DSCID (MAXPRS) real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c *** end of declarations for common data ***
c These variables are used when establishing the PERSON array
c from the PEOPLE file.
integer CURRNT, PRVDEX character* (IDLEN) PREVID, CURRID integer RELSHP
c These variables are used to accept and resolve requests for
c RELSHP information.
integer
character*(BUFLEN)
character* (NAMLEN)
integer
character*(MSGLEN) ERRMSG
integer P1DEX, P2DEX
character*7 PRNOUN

BUFDEX, SEMLOC
REQBUF
P1IDNT, P2IDNT
P1FND, P2FND

```
c &** execution of main sequence begins here ***
    open (unit=10, file=`PEOPLE.DAT`, status=`old`, form=`formatted`)
    This loop reads in the PEOPLE file and constructs the PERSON
    array from it (one PERSON = one record = one array entry).
    As records are read in, links are constructed to represent the
    PARENT-CHILD or SPOUSE relationship. The array then implements
    a directed graph which is used to satisfy subsequent user
    requests. The file is assumed to be correct - no validation
    is performed on it.
    do 110 CURRNT=1, MAXPRS
        copy direct information from file to array
        read (unit=10, fmt='(a20, a3, a1, 3a3)', end=111)
    1 NAME(CURRNT), IDENT(CURRNT), GENDER(CURRNT)
    2 ((RELID(CURRNT, ITEMP), ITEMP=FATHID, SPOUID))
        Location of adjacent persons as yet undetermined
        NBRCNT (CURRNT) = 0
        Descendants as yet undetermined
        DSCID (CURRNT) = NULLID
        Compare this PERSON against all previously entered PERSONs
        to search for relationships.
        CURRID = IDENT (CURRNT)
        do 120 PRVDEX = 1, CURRNT-1
        PREVID = IDENT (PRVDEX)
        Search for father, mother, or spouse relationship in
        either direction between this and previous PERSON.
        Assume at most one relationship exists.
        do }130\mathrm{ RELSHP = FATHID, SPOUID
            if (PREVID .eq. RELID (CURRNT, RELSHP)) then
                    ca11 LNKREL (CURRNT, RELSHP, PRVDEX)
                goto 131
            else if (CURRID .eq. RELID (PRVDEX, RELSHP)) then
                call LNKREL (PRVDEX, RELSHP, CURRNT)
                goto 131
            end if
        continue
        continue
        continue
    continue
    continue
    NUMPER = CURRNT - 1
    close (unit=10, status=' keep`)
c PERSON array is now loaded and edges between immediate relatives
c (PARENT-CHILD or SPOUSE-SPOUSE) are established.
```

c Loop accepts requests and finds relationship (if any)
c between pairs of PERSONs.
c
end if
if (P2FND .eq. 0) then

else if (P2FND .gt. 1) then write (unit=*,

1
2
end if
end if
end processing of syntactically legal request
else
write (unit=*, fmt=9004) ERRMSG
format (' Incorrect request format: ', a40)
end if
goto 200
201
continue
write (unit=*, fmt=-(-- End of relation-finder." ${ }^{\left.-)^{-}\right)}$
c End of main line of RELATE
end
c
procedures under RELATE
subroutine LNKREL (FRMDEX, RELSHP, TODEX)
C
establishes cross-indexing between immediately related PERSONs.
integer FRMDEX, TODEX, RELSHP
c Each PERSON's record in the file identifies at most three
c others directly related: father, mother, and spouse
integer FATHID, MOTHID, SPOUID
parameter $($ FATHID $=1$, MOTHID $=2, \operatorname{SPOUID}=3$ )
integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NEPHEW, COUSIN, NULLRL
parameter (PARENT $=1$, CHILD $=2$, SPOUSE $=3$, SIBLNG $=4$,
$1 \quad$ UNCLE $=5$, NEPHEW $=6$, COUSIN $=7$, NULLRL $=8$ )
if (RELSHP .eq. SPOUID) then
call LNKONE (FRMDEX, SPOUSE, TODEX)
call LNKONE (TODEX, SPOUSE, FRMDEX)
else
c

RELSHP is father or mother
call LNKONE (FRMDEX, PARENT, TODEX)
call LNKONE (TODEX, CHILD, FRMDEX)
end if
end

```
    subroutine LNKONE (FRMDEX, THSEDG, TODEX)
c Establishes the NBR pointers from one PERSON to another
    integer FRMDEX, TODEX, THSEDG
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
    1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
    1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
    character NULLID*(IDLEN)
    parameter (NULLID = '000')
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
character*(NAMLEN) NAME (MAXPRS)
character*(IDLEN) IDENT (MAXPRS)
character* GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS)
integer NBRDEX (MAXPRS, MAXNBR)
integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons
integer NUMPER
c *** end of declarations for common data ***
ITEMP \(=\) NBRCNT \((\) FRMDEX) +1
NBRCNT (FRMDEX) \(=\) ITEMP
NBRDEX (FRMDEX, ITEMP) \(=\) TODEX
NBREDG (FRMDEX, ITEMP) \(=\) THSEDG
end
```

```
    subroutine PROMPT (REQBUF)
c Issues prompt for user-request, reads in request,
c blank-fills buffer, and skips to next line of input.
    character*(*) REQBUF
    write (unit=*, fmt=9001)
9 0 0 1
format (/,', Enter two person-identifiers (name or number),'
    2 /...' separated by semicolon. Enter "stop" to stop.")
c *** NOTE THAT THIS IS NOT A STANDARD WAY TO READ A LINE FROM
c *** THE TERMINAL (see section 12.9.5.2.1). THE STANDARD
c *** PROVIDES NO SUCH CAPABILITY.
    read (unit=*, fmt=-(a60)-) REQBUF
    end
    subroutine CHKRQS (REQBUF, REQST, PIIDNT, P2IDNT)
c Performs syntactic check on request in buffer.
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
    1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
    1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
    character NULLID*(IDLEN)
    parameter (NULLID = '000`)
character REQOK*10, REQSTP*4
parameter (REQOK = 'Request OK', REQSTP = 'stop')
character REQBUF*(BUFLEN), REQST*(MSGLEN)
character*(NAMLEN) P1IDNT, P2IDNT, LTRIM
integer SEMLOC
SEMLOC = INDEX (REQBUF,`;`)
P2IDNT = REQBUF (SEMLOC+1 : BUFLEN)
c set REQST, based on results of scan of REQBUF, and
c fill in P1IDNT and P2IDNT.
```

```
    if (SEMLOC .eq. 0 .or. INDEX (P2IDNT, `;`) .ne. 0) then
    REQST = "must be exactly one semicolon."
    else
        if (SEMLOC .eq. 1) then
        P1IDNT = - -
    else
            P1IDNT = REQBUF (1 : SEMLOC-1)
    end if
    if (P1IDNT .eq. - ") then
        REQST = 'null field preceding semicolon.'
    else if (P2IDNT .eq. - ') then
        REQST = 'null field following semicolon.'
    else
            REQST = REQOK
            PIIDNT = LTRIM (P1IDNT)
            P2IDNT = LTRIM (P2IDNT)
        end if
    end if
    end
    character*(*) function LTRIM (STRING)
c
    LTRIM deletes leading spaces and returns the resulting value.
    character*(*) STRING
    do 100 ITEMP = 1, 1en(STRING)
    if (STRING (ITEMP : ITEMP) .ne. ' `) goto 101
    continue
    continue
    LTRIM = STRING (ITEMP : len(STRING))
    end
    subroutine SEEKPR (P1IDNT, P2IDNT, P1DEX, P2DEX,
1
                            P1FND, P2FND)
SEEKPR scans through the PERSON array, looking for the two
requested PERSONs. Match may be by NAME or unique IDENT-number.
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
    character NULLID*(IDLEN)
    parameter (NULLID = ' O00')
character*(NAMLEN) P1IDNT, P2IDNT
integer P1DEX, P2DEX, P1FND, P2FND
integer CURRNT
```

c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
character*(NAMLEN) NAME (MAXPRS)
character*(IDLEN) IDENT (MAXPRS)
character*1 GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS)
integer NBRDEX (MAXPRS, MAXNBR)
integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real
DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character* (IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons
integer NUMPER
c *** end of declarations for common data ***
P1DEX $=0$
P2DEX $=0$
P1FND $=0$
P2FND $=0$
do 100 CURRNT $=1$, NUMPER
c allow identification by name or number.
if (P1IDNT .eq. IDENT (CURRNT) .or.
1 P1IDNT .eq. NAME (CURRNT)) then
P1FND $=$ P1FND +1
P1DEX = CURRNT
end if
if (P2IDNT .eq. IDENT (CURRNT) .or.
1
P2IDNT .eq. NAME (CURRNT)) then
P2FND $=$ P2FND +1
P2DEX = CURRNT
end if
continue
end
subroutine FINDRL (TRGDEX, SRCDEX)
c Finds shortest path (if any) between two PERSONs and c determines their relationship based on immediate relations c traversed in path. PERSON array simulates a directed graph, c and algorithm finds shortest path, based on following $c$ weights: PARENT-CHILD edge $=1.0$
c $\quad$ SPOUSE-SPOUSE edge $=1.8$

```
    integer TRGDEX, SRCDEX
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
```

    character NULLID*(IDLEN)
    parameter ( $\mathrm{NULLID}=-\mathrm{OOO}^{\prime}$ )
c A node in the graph ( $=$ PERSON) has either already been reached,
c is immediately adjacent to those reached, or farther away.
integer REACHD, NEARBY, NOSEEN
parameter ( REACHD $=1$, NEARBY $=2$, NOSEEN $=3$ )
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
1
EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file

| character*(NAMLEN) | NAME | (MAXPRS) |
| :--- | :--- | :--- |
| character*(IDLEN) | IDENT | (MAXPRS) |
| character*1 | GENDER | (MAXPRS) |

c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph

| integer | NBRCNT (MAXPRS) |
| :--- | :--- |
| integer | NBRDEX (MAXPRS, MAXNBR) |
| integer | NBREDG (MAXPRS, MAXNBR) |

C data used when traversing graph to resolve user request:
real : DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c *** end of declarations for common data ***

```
        integer PERDEX, THSNOD, ADJNOD,
        1
        integer THSEDG, THSNBR
        integer RELSHP
        real
        MINDIS
    integer SRCHNG, SUCCES, FAILED
    parameter (SRCHNG = 1, SUCCES = 2, FAILED = 3)
    integer SRCHST
c begin execution of FINDRL
c initialize PERSON-array for processing -
c mark all nodes as not seen
    do }100\mathrm{ PERDEX = 1, NUMPER
        RCHST (PERDEX) = NOSEEN
100 continue
    THSNOD = SRCDEX
c mark source node as reached
    RCHST (THSNOD) = REACHD
    DSTSRC (THSNOD) = 0.0
c no NEARBY nodes exist yet
    LASTNR = 0
    if (THSNOD .eq. TRGDEX) then
        SRCHST = SUCCES
    else
        SRCHST = SRCHNG
    end if
```

c Loop keeps processing closest-to-source, unreached node c until target reached, or no more connected nodes.
c to translate path to English description of relationship.
if (SRCHST .eq. FAILED) then
write (unit=*, fmt=9001) NAME (TRGDEX), NAME (SRCDEX)
format (a22, - is not related to ${ }^{-}$, a20)
else
c success - parse path to find and display relationship
call RESOLV (SRCDEX, TRGDEX)
compute proportion of common genetic material
call CMPTGN (SRCDEX, TRGDEX)
end if
end
c procedures under FINDRL

```
    subroutine PROCAD (BASNOD, NXTNOD, NBEDGE, NEARND, LASTNR)
    NXTNOD is adjacent to last-reached node (= BASNOD).
    If NXTNOD already reached, do nothing.
    If previously seen, check whether path thru BASNOD is
    shorter than current path to NXTNOD, and if so re-link
    next to base.
    If not previously seen, link next to base node.
    integer NXTNOD, BASNOD, NEARND(*), LASTNR
    integer NBEDGE
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
    character NULLID*(IDLEN)
    parameter (NULLID = '000-)
```

c A node in the graph (= PERSON) has either already been reached, c is immediately adjacent to those reached, or farther away.
integer REACHD, NEARBY, NOSEEN
parameter ( REACHD $=1$, NEARBY $=2$, NOSEEN $=3$ )
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
1
EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
character*(NAMLEN) NAME (MAXPRS)
character*(IDLEN) IDENT (MAXPRS)
character*1 GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer $\quad$ NBRCNT (MAXPRS)
integer $\quad$ NBRDEX (MAXPRS, MAXNBR)
integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real
DSCGEN (MAXPRS)

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c

```
NUMPER keeps track of the actual number of persons
    integer NUMPER
*** end of declarations for common data ***
    real WGHTEG, DSTBAS
    begin execution of PROCAD
    if (RCHST (NXTNOD) .ne. REACHD) then
        if (NBEDGE .eq. SPOUSE) then
        WGHTEG = 1.8
    else
        WGHTEG = 1.0
    end if
    DSTBAS = WGHTEG + DSTSRC (BASNOD)
    if (RCHST (NXTNOD) .eq. NOSEEN) then
        change status of THSNOD from not-seen to NEARBY
        RCHST (NXTNOD) = NEARBY
        LASTNR = LASTNR + 1
        NEARND (LASTNR) = NXTNOD
        link next to base by re-setting its predecessor index to
        point to base, note type of edge, and re-set distance
        as it is through base node.
        DSTSRC (NXTNOD) = DSTBAS
        PATHPR (NXTNOD) = BASNOD
        EDGPRD (NXTNOD) = NBEDGE
    else
        RCHST is NEARBY
            if (DSTBAS .lt. DSTSRC (NXTNOD)) then
                    link next to base by re-setting its predecessor index to
                point to base, note type of edge, and re-set distance
                    as it is through base node.
                DSTSRC (NXTNOD) = DSTBAS
                    PATHPR (NXTNOD) = BASNOD
                    EDGPRD (NXTNOD) = NBEDGE
            end if
        end if
    end if
    end
```

subroutine RESOLV (SRCDEX, TRGDEX)
c RESOLV condenses the shortest path to a series of
c relationships for which there are English descriptions.
integer SRCDEX, TRGDEX
c Establish global constants

```
    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
    1 MSGLEN, MAXNBR, MAXGVN
    parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
    1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
    character NULLID*(IDLEN)
    parameter (NULLID = '000')
    character MALE*1, FEMALE*1
    parameter (MALE = 'M', FEMALE = ' F')
    integer PARENT, CHILD, SPOUSE, SIBLNG,
    1 UNCLE, NEPHEW, COUSIN, NULLRL
    parameter (PARENT = 1, CHILD = 2, SPOUSE = 3, SIBLNG = 4,
    1 UNCLE = 5, NEPHEW = 6, COUSIN = 7, NULLRL = 8)
```

c sibling proximity can have three values
integer STEP, HALF, FULL
parameter $(S T E P=1$, HALF $=2$, FULL $=3$ )
c These common blocks hold the PERSON array, which is global to
c the entire program. common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file

| character*(NAMLEN) | NAME | (MAXPRS) |
| :--- | :--- | :--- |
| character*(IDLEN) | IDENT | (MAXPRS) |
| character*1 | GENDER | (MAXPRS) |

c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS) integer NBRDEX (MAXPRS, MAXNBR) integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c *** end of declarations for common data ***
c these variables are used to generate key-person data integer GENCNT, THSCUZ integer THS PRX
c these variables are used to condense the path
common /KEYPER/ RELNXT, PERDEX, GENGAP, PRXMTY, CUZRNK
c Key persons are the ones in the relationship path which remain
c after the path is condensed.

```
integer RELNXT (MAXPRS)
integer PERDEX (MAXPRS)
integer GENGAP (MAXPRS)
integer PRXMTY (MAXPRS)
integer CUZRNK (MAXPRS)
integer KEYREL, LATREL, PRIREL, NXTPRI
integer KEYDEX, LATDEX, PRIDEX, THSNOD
integer GAP1, GAP2
logical SEEKMR, FULSIB
```

c begin execution of RESOLV
write (unit=*,
1 fmt=-(-" Shortest path between identified persons: "-)")
c Display path and initialize key person arrays from path elements.
THSNOD $=$ TRGDEX
do 100 KEYDEX $=1$, MAXPRS
if (THSNOD .eq. SRCDEX) goto 101
PERDEX (KEYDEX) $=$ THSNOD
PRXMTY (KEYDEX) = FULL
RELNXT (KEYDEX) = EDGPRD (THSNOD)
if (EDGPRD (THSNOD) .eq. SPOUSE) then
write (unit=*, fmt=-(a22, ${ }^{--}$is spouse of $\left.\left.{ }^{-9}\right)^{-}\right)$NAME (THSNOD)
GENGAP (KEYDEX) $=0$ else

GENGAP (KEYDEX) $=1$
if (EDGPRD (THSNOD) .eq. PARENT) then write (unit=*, $\left.f m t={ }^{\prime}\left(a 22,{ }^{\prime \prime} \text { is parent of" }{ }^{\prime \prime}\right)^{\prime}\right)$

1 NAME (THSNOD)
else
write (unit=*, fmt='(a22, " is child of ${ }^{\left.\left.-{ }^{-}\right)^{-}\right)}$
1 NAME (THSNOD)
end if
end if
THSNOD $=$ PATHPR (THSNOD)
continue
continue
write (unit=*, fmt=-(a22)$\left.{ }^{-}\right)$NAME (THSNOD)
PERDEX (KEYDEX) = THSNOD
RELNXT (KEYDEX) = NULLRL
RELNXT (KEYDEX + 1) = NULLRL
c resolve CHILD-PARENT and CHILD-SPOUSE-PARENT relations
c to SIBLNG relations.
do 200 KEYDEX $=1$, MAXPRS
if (RELNXT (KEYDEX) .eq. NULLRL) goto 201
if (RELNXT (KEYDEX) .eq. CHILD) then
LATREL = RELNXT (KEYDEX + 1)
if (LATREL .eq. PARENT) then
c

1
c
c resolve CHILD-CHILD-... and PARENT-PARENT-... relations to
c direct descendant or ancestor relations.
do 300 KEYDEX $=1$, MAXPRS
if (RELNXT (KEYDEX) .eq. NULLRL) goto 301
if (RELNXT (KEYDEX) .eq. CHILD .or.
RELNXT (KE YDEX) .eq. PARENT) then do 310 LATDEX $=$ KEYDEX +1 , MAXPRS
if (RELNXT (LATDEX) . ne. RELNXT (KEYDEX)) goto 311
continue
continue
GENCNT = LATDEX - KEYDEX
if (GENCNT .gt. 1) then
compress generations
GENGAP (KEYDEX) $=$ GENCNT
call CONDNS (KEYDEX, GENCNT - 1)
end if
end if
c
c

```
resolve CHILD-SIBLNG-PARENT to COUSIN,
    CHILD-SIBLNG to NEPHEW,
    SIBLNG-PARENT
    to UNCLE.
    do 400 KEYDEX \(=1\), MAXPRS
    if (RELNXT (KEYDEX) .eq. NULLRL) goto 401
    LATREL = RELNXT (KEYDEX + 1)
    if (RELNXT (KEYDEX) .eq. CHILD .and. LATREL .eq. SIBLNG) then
        found COUSIN or NEPHEW
        \(\operatorname{PRXMTY}\) (KEYDEX) \(=\) PRXMTY (KE YDEX + 1)
        if (RELNXT (KEYDEX + 2) .eq. PARENT) then
            found COUSIN
            GAP1 = GENGAP (KEYDEX)
            GAP2 \(=\) GENGAP (KEYDEX + 2)
            GENGAP (KEYDEX) = abs (GAP1 - GAP2)
            CUZRNK (KEYDEX) \(=\) min (GAP1, GAP2)
            RELNXT (KEYDEX) \(=\) COUSIN
            call CONDNS (KEYDEX, 2)
        else
            found NEPHEW
            RELNXT (KEYDEX) = NEPHEW
            call CONDNS (KEYDEX, 1)
        end if
    else
            if (RELNXT (KEYDEX) .eq. SIBLNG .and.
1 LATREL .eq. PARENT) then
            found UNCLE
            GENGAP (KEYDEX) = GENGAP (KEYDEX + 1)
            RELNXT (KEYDEX) = UNCLE
            call CONDNS (KEYDEX, 1)
        end if
    end if
continue
continue
```

c Loop below will pick out valid adjacent strings of elements

C
c
C
c
C
call SHOWRE (KEYDEX, LATDEX, PRIDEX)
KEYDEX = LATDEX + 1
goto 500

```
    logical function FULSIB (INDEX1, INDEX2)
c Determines whether two PERSONs are full siblings, i.e.,
c have the same two parents.
```



```
    character NULLID*(IDLEN)
parameter (NULLID = '000')
integer FATHID, MOTHID, SPOUID
parameter (FATHID = 1, MOTHID = 2, SPOUID = 3)
c These common blocks hold the PERSON array, which is global to
c the entire program.
    common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
    1 EDGPRD, RCHST, DSCGEN, NUMPER
    common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
luracter*(NAMLEN) 
c IDENTs of immediate relatives - father, mother, spouse
character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph
integer NBRCNT (MAXPRS)
integer NBRDEX (MAXPRS, MAXNBR)
integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real . DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c *** end of declarations for common data ***
```

```
    FULSIB =
```

    FULSIB =
    1 RELID (INDEX1, FATHID) .ne. NULLID .and.
    2 RELID (INDEX1, MOTHID) . ne. NULLID . and.
    3 RELID (INDEX1, FATHID) .eq. RELID (INDEX2, FATHID) .and.
    4 RELID (INDEX1, MOTHID) .eq. RELID (INDEX2, MOTHID)
    end
    ```

Page
subroutine CONDNS (ATDEX, GAPSIZ)
c CONDNS condenses superfluous entries from the
key person arrays, starting at ATDEX.
integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
parameter (MAXPRS \(=300\), NAMLEN \(=20\), IDLEN \(=3\), BUFLEN \(=60\),
\(1 \quad\) MSGLEN \(=40\), MAXNBR \(=20\), MAXGVN \(=3\) )
character NULLID*(IDLEN)
parameter (NULLID \(={ }^{-} 000^{-}\))
integer PARENT, CHILD, SPOUSE, SIBLNG, 1 UNCLE, NEPHEW, COUSIN, NULLRL
parameter (PARENT \(=1\), CHILD \(=2\), SPOUSE \(=3, \operatorname{SIBLNG~}=4\),
\(1 \quad\) UNCLE \(=5\), NEPHEW \(=6\), COUSIN \(=7\), NULLRL \(=8\) )
common /KEYPER/ RELNXT, PERDEX, GENGAP, PRXMTY, CUZRNK
c Key persons are the ones in the relationship path which remain
\begin{tabular}{ll} 
integer & RELNXT (MAXPRS) \\
integer & PERDEX (MAXPRS) \\
integer & GENGAP (MAXPRS) \\
integer & PRXMTY (MAXPRS) \\
integer & CUZRNK (MAXPRS) \\
integer & \\
&
\end{tabular}

RCVDEX \(=\) ATDEX
100
continue
RCVDEX \(=\) RCVDEX +1
SENDEX = RCVDEX + GAPSIZ
RELNXT (RCVDEX) = RELNXT (SENDEX)
PERDEX (RCVDEX) = PERDEX (SENDEX)
GENGAP (RCVDEX) \(=\) GENGAP (SENDEX)
PRXMTY (RCVDEX) = PRXMTY (SENDEX)
CUZRNK (RCVDEX) = CUZRNK (SENDEX)
if (RELNXT (SENDEX) .ne. NULLRL) goto 100
end
c procedures under RESOLV
subroutine SHOWRE (FSTDEX, LSTDEX, PRIDEX)
c SHOWRE takes 1,2 , or 3 adjacent elements in the
c condensed table and generates the English description of
c
c Establish global constants
integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
parameter \((\) MAXPRS \(=300\), NAMLEN \(=20\), IDLEN \(=3\), BUFLEN \(=60\),
1 MSGLEN \(=40\), MAXNBR \(=20\), MAXGVN \(=3\) )
character NULLID* (IDLEN)
parameter (NULLID \(={ }^{-} \mathrm{OOO}^{\circ}\) )
character MALE*1, FEMALE*1
parameter (MALE \(={ }^{-} M^{-}\), FEMALE \(=-F^{\prime}\) )
integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NEPHEW, COUSIN, NULLRL
parameter (PARENT \(=1\), CHILD \(=2\), SPOUSE \(=3\), SIBLNG \(=4\),
1 UNCLE \(=5\), NEPHEW \(=6\), COUSIN \(=7\), NULLRL \(=8\) )
c sibling proximity can have three values
integer STEP, HALF, FULL
parameter \((\operatorname{STEP}=1\), HALF \(=2\), FULL \(=3\) )
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR,
1
EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
\begin{tabular}{lll} 
character*(NAMLEN) & NAME & (MAXPRS) \\
character*(IDLEN) & IDENT & (MAXPRS) \\
character*1 & GENDER & (MAXPRS)
\end{tabular}
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph \(\begin{array}{lll}\text { integer } & \text { NBRCNT } & \text { (MAXPRS) } \\ \text { integer } & \text { NBRDEX } & \text { (MAXPRS, MAXNBR) } \\ \text { integer } & \text { NBREDG } & \text { (MAXPRS, MAXNBR) }\end{array}\)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER common /KEYPER/ RELNXT, PERDEX, GENGAP, PRXMTY, CUZRNK
c Key persons are the ones in the relationship path which remain
c after the path is condensed.
```

integer RELNXT (MAXPRS)
integer PERDEX (MAXPRS)
integer GENGAP (MAXPRS)
integer PRXMTY (MAXPRS)
integer CUZRNK (MAXPRS)

```
c *** end of declarations for common data ***
```

logical
integer
character
integer
character
integer
integer
character*75 OUTBUF
integer OUTPTR

```
    set THSGAP
    if (PRIREL .eq. PARENT .or. PRIREL .eq. CHILD .or.
    1 PRIREL .eq. UNCLE .or. PRIREL .eq. NEPHEW .or.
    2 PRIREL .eq. COUSIN) then
        THSGAP = GENGAP (PRIDEX)
    else
        THSGAP \(=0\)
    end if
    if (FSTREL .eq. SPOUSE .and.
    1 (PRIREL .eq. SIBLNG .or. PRIREL .eq. CHILD .or.
    2 PRIREL .eq. NEPHEW .or. PRIREL .eq. COUSIN)) then
        INLAW \(=\).true.
    else
        if (LSTREL .eq. SPOUSE .and.
1 (PRIREL .eq. SIBLNG .or. PRIREL .eq. PARENT .or.
2 PRIREL .eq. UNCLE .or. PRIREL .eq. COUSIN)) then
INLAW \(=\).true.
        else
            INLAW \(=. f a l s e\).
        end if
    end if
c set THSCUZ
    if (PRIREL .eq. COUSIN) then
        THSCUZ = CUZRNK (PRIDEX)
    else
        THSCUZ \(=0\)
    end if
c
c
```

    parameters are set - now generate display.
    OUTBUF = NAME (PERDEX (FSTDEX)) // ' is `
    OUTPTR = NAMLEN + 5
    if (PRIREL .eq. PARENT .or. PRIREL .eq. CHILD .or.
    1 PRIREL .eq. UNCLE .or. PRIREL .eq. NEPHEW) then
display generation-qualifier
if (THSGAP .ge. 3) then
cal1 APPEND (OUTBUF, OUTPTR, 'great')
if (THSGAP .gt. 3) then
write (unit=TWODIG, fmt='(i2)') THSGAP - 2
call APPEND (OUTBUF, OUTPTR, "*' // TWODIG)
end if
cal1 APPEND (OUTBUF, OUTPTR, "-`)         end if         if (THSGAP .ge. 2) then             call APPEND (OUTBUF, OUTPTR, 'grand-`)
end if
else
if (PRIREL .eq. COUSIN .and. THSCUZ .gt. 1) then
display cousin-degree
write (unit=TWODIG, fmt='(i2)') THSCUZ
ca11 APPEND (OUTBUF, OUTPTR, TWODIG)
SUFPTR = mod (THSCUZ, 10)
if (SUFPTR .gt. 3) SUFPTR = 0
SUFPTR = 3 * SUFPTR + 1
SUFCHR = "th st nd rd
cal1 APPEND (OUTBUF, OUTPTR, SUFCHR (SUFPTR : SUFPTR + 2))
end if
end if
if (THSPRX .eq. STEP) then
ca11 APPEND (OUTBUF, OUTPTR, `step-`)
else
if (THSPRX .eq. HALF) then
ca11 APPEND (OUTBUF, OUTPTR, `half-`)
end if
end if

```
if (GENDER (PERDEX (FSTDEX)) .eq. MALE) then
    goto ( \(201,202,203,204,205,206,297,298)\), PRIREL
    continue
        call APPEND (OUTBUF, OUTPTR, "father")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, 'son')
        goto 300
    continue
        cal1 APPEND (OUTBUF, OUTPTR, "husband")
        goto 300
    continue
        ca11 APPEND (OUTBUF, OUTPTR, "brother")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, "uncle")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, 'nephew")
        goto 300
    else
    gender is FEMALE
    goto ( \(251,252,253,254,255,256,297,298\) ), PRIREL
    continue
        call APPEND (OUTBUF, OUTPTR, "mother")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, "daughter")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, "wife")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, 'sister")
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, 'aunt")
        goto 300
    continue
        cal1 APPEND (OUTBUF, OUTPTR, 'niece")
        goto 300
    end if
    continue
        call APPEND (OUTBUF, OUTPTR, "cousin')
        goto 300
    continue
        call APPEND (OUTBUF, OUTPTR, 'nul1')
        goto 300
    continue
```

    if (INLAW) cal1 APPEND (OUTBUF, OUTPTR, "-in-1aw`)
    if (PRIREL .eq. COUSIN .and. THSGAP .gt. 0) then
        if (THSGAP .gt. 1) then
            write (unit=TWODIG, fmt=`(i2)`) THSGAP
            call APPEND (OUTBUF, OUTPTR, ' '//TWODIG//' times removed")
        else
            call APPEND (OUTBUF, OUTPTR, ' once removed")
        end if
    end if
cal1 APPEND (OUTBUF, OUTPTR, ` of`)
write (unit=*, fmt='(a77)') OUTBUF
end
subroutine APPEND (STRING, PTR, ADDEND)
c APPEND appends the contents of ADDEND to STRING in the position
c indicated by PTR, and increments PTR

```
```

character STRING*(*), ADDEND*(*)

```
character STRING*(*), ADDEND*(*)
integer PTR, ADDLEN
integer PTR, ADDLEN
ADDLEN = 1en (ADDEND)
ADDLEN = 1en (ADDEND)
STRING (PTR : PTR + ADDLEN - 1) = ADDEND
STRING (PTR : PTR + ADDLEN - 1) = ADDEND
PTR = PTR + ADDLEN
PTR = PTR + ADDLEN
end
```

end

```
c procedures under FINDRL
subroutine CMPTGN (INDEX1, INDEX2)
c CMPTGN assumes that each ancestor contributes
c half of the genetic material to a PERSON. It finds common
c ancestors between two PERSONs and computes the expected
c value of the proportion of common material.
integer INDEX1, INDEX2
integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
parameter (MAXPRS \(=300\), NAMLEN \(=20\), IDLEN \(=3\), BUFLEN \(=60\),
1 MSGLEN \(=40\), MAXNBR \(=20\), MAXGVN \(=3\) )
character NULLID*(IDLEN)
parameter ( NULLID \(^{\prime}={ }^{-} 000^{\circ}\) )
c These common blocks hold the PERSON array, which is global to
\(=\) the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file character*(NAMLEN) NAME (MAXPRS) character*(IDLEN) IDENT (MAXPRS) character*1 GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS) integer NBRDEX (MAXPRS, MAXNBR) integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request: real DSTSRC (MAXPRS) integer PATHPR (MAXPRS) integer EDGPRD (MAXPRS) integer RCHST (MAXPRS)
c data used to compute common genetic material character*(IDLEN) DSCID (MAXPRS) rea1 DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c STACK is common to the routines which calculate genetic overlap.
c It is used to implement recursive traversal of the ancestor trees.
integer STKSIZ
parameter \(\left(S^{\prime} T K S I Z=50\right)\)
common /STACK/ PROPTN, CONTRB, COUNTD, PERDEX, NXTNBR, 1 STKPTR
\begin{tabular}{lll} 
real & PROPTN & (STKSIZ) \\
real & CONTRB & (STKSIZ) \\
real & COUNTD & (STKSIZ) \\
integer & PERDEX & (STKSIZ) \\
integer & NXTNBR & (STKSIZ) \\
integer & STKPTR &
\end{tabular}
c *** end of declarations for common data ***
real COMPRP
c First zero out all ancestors to allow adding. This is necessary
c because there might be two paths to an ancestor.
STKPTR = 1
PERDEX (STKPTR) = INDEX1
NXTNBR \((\) STKPTR) \(=0\)
continue
call ZERPRO
if (STKPTR .ge. 1) goto 100
continue
now mark with shared PROPTN
STKPTR = 1
PERDEX (STKPTR) = INDEX1
NXTNBR \((\) STKPTR) \(=0\)
PROPTN \((S T K P T R)=1.0\)
continue
call MRKPRO (IDENT (INDEX1))
if (STKPTR .ge. 1) goto 200
continue
traverse ancestor tree for INDEX2. summing overlap with
marked tree of INDEX1
COMPRP \(=0.0\)
STKPTR \(=1\)
PERDEX (STKPTR) = INDEX2
NXTNBR \((\) STKPTR \()=0\)
PROPTN \((S T K P T R)=1.0\)
COUNTD \((S T K P T R)=0.0\)
continue
call CHKCOM (COMPRP, IDENT (INDEX1))
if (STKPTR .ge. 1) goto 300
continue
write (unit=*, fmt=9001) COMPRP
9001 format( \({ }^{( }\)Proportion of common genetic material \(=\)' , lp, el2.5e2)
end
subroutine ZERPRO
c ZERPRO recursively seeks out all ancestors and
c zeros them out.
integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
parameter (MAXPRS \(=300\), NAMLEN \(=20\), IDLEN \(=3\), BUFLEN \(=60\),
1 MSGLEN \(=40\), MAXNBR \(=20\), MAXGVN \(=3\) )
character NULLID*(IDLEN)
parameter ( \(\mathrm{NULLID}={ }^{-} \mathrm{OOO}^{\prime}\) )
integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NE PHEW, COUSIN, NULLRL
parameter (PARENT \(=1\), CHILD \(=2\), SPOUSE \(=3\), SIBLNG \(=4\),
1 UNCLE \(=5\), NEPHEW \(=6\), COUSIN \(=7\), NULLRL \(=8\) )
c These common blocks hold the PERSON array, which is global to
c the entire program. common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
character* (NAMLEN) NAME (MAXPRS)
character*(IDLEN) IDENT (MAXPRS)
character*1 GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS) integer NBRDEX (MAXPRS, MAXNBR) integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character*(IDLEN) DSCID (MAXPRS)
real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer

NUMPER
c STACK is common to the routines which calculate genetic overlap.
c It is used to implement recursive traversal of the ancestor trees.
integer STKSIZ
parameter \((S T K S I Z=50)\)
common /STACK/ PROPTN, CONTRB, COUNTD, PERDEX, NXTNBR, 1 STKPTR
\begin{tabular}{lll} 
real & PROPTN & (STKSIZ) \\
real & CONTRB & (STKSIZ) \\
real & COUNTD & (STKSIZ) \\
integer & PERDEX & (STKSIZ) \\
integer & NXTNBR & (STKSIZ) \\
integer & STKPTR &
\end{tabular}
c *** end of declarations for common data ***

\section*{Page 130}
integer ZERDEX, THSNBR
ZERDEX = PERDEX (STKPTR)
if (NXTNBR (STKPTR) .eq. 0) then
DSCGEN (ZERDEX) \(=0.0\)
NXTNBR \((S T K P T R)=1\)
end if
do 100 THSNBR = NXTNBR (STKPTR), NBRCNT (ZERDEX)
if (NBREDG (ZERDEX, THSNBR) .eq. PARENT) goto 101
100 continue
101 continue
if (THSNBR .gt. NBRCNT (ZERDEX)) then
c no more ancestors from this person STKPTR = STKPTR - 1
else
c
set up for next ancestor
NXTNBR (STKPTR) \(=\) THSNBR +1
STKPTR \(=\) STKPTR +1
PERDEX (STKPTR) \(=\) NBRDEX (ZERDEX, THSNBR)
NXTNBR \((S T K P T R)=0\)
end if
end
subroutine MRKPRO (MARKER)
c MRKPRO recursively seeks out all ancestors and
c marks them with the sender's proportion of shared
c genetic material. This proportion is diluted by one-half
c for each generation.
```

    integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
    1 MSGLEN, MAXNBR, MAXGVN
parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
character NULLID*(IDLEN)
parameter (NULLID = - 000')
integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NE PHEW, COUSIN, NULLRL
parameter (PARENT = 1, CHILD = 2, SPOUSE = 3, SIBLNG = 4,
1 UNCLE = 5, NEPHEW = 6, COUSIN = 7, NULLRL = 8)

```
c These common blocks hold the PERSON array, which is global to
c the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file
\begin{tabular}{lll} 
character*(NAMLEN) & NAME & (MAXPRS) \\
character*(IDLEN) & IDENT & (MAXPRS) \\
character*1 & GENDER & (MAXPRS)
\end{tabular}
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph integer NBRCNT (MAXPRS)
integer NBRDEX (MAXPRS, MAXNBR) integer NBREDG (MAXPRS, MAXNBR)
c data used when traversing graph to resolve user request:
real DSTSRC (MAXPRS)
integer PATHPR (MAXPRS)
integer EDGPRD (MAXPRS)
integer RCHST (MAXPRS)
c data used to compute common genetic material
character* (IDLEN) DSCID (MAXPRS)
real
DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons
integer NUMPER
c STACK is common to the routines which calculate genetic overlap.
c It is used to implement recursive traversal of the ancestor trees.
integer STKSIZ
parameter (STKSIZ = 50)
common /STACK/ PROPTN, CONTRB, COUNTD, PERDEX, NXTNBR,
1 STKPTR
\begin{tabular}{lll} 
real & PROPTN & (STKSIZ) \\
real & CONTRB & (STKSIZ) \\
real & COUNTD & (STKSIZ) \\
integer & PERDEX & (STKSIZ) \\
integer & NXTNBR & (STKSIZ) \\
integer & STKPTR &
\end{tabular}
c \(* * *\) end of declarations for common data \(* * *\)
character*(IDLEN) MARKER
integer MRKDEX, THSNBR
```

    MRKDEX = PERDEX (STKPTR)
    if (NXTNBR (STKPTR) .eq. 0) then
        DSCID (MRKDEX) = MARKER
        DSCGEN (MRKDEX) = DSCGEN (MRKDEX) + PROPTN (STKPTR)
        NXTNBR (STKPTR) = 1
    end if
    do 100 THSNBR = NXTNBR (STKPTR), NBRCNT (MRKDEX)
        if (NBREDG (MRKDEX, THSNBR) .eq. PARENT) goto 101
    continue
    continue
    if (THSNBR .gt. NBRCNT (MRKDEX)) then
    no more ancestors from this person
    STKPTR = STKPTR - 1
    else
    c set up for next ancestor
NXTNBR (STKPTR) = THSNBR + 1
STKPTR = STKPTR + 1
PERDEX (STKPTR) = NBRDEX (MRKDEX, THSNBR)
NXTNBR (STKPTR) = 0
PROPTN (STKPTR) = PROPTN (STKPTR - 1) / 2.0
end if
end
subroutine CHKCOM (COMPRP, MTCHID)
CHKCOM searches all the ancestors of CHKDEX to see if any have
been marked, and if so adds the appropriate amount to COMPRP.
integer MAXPRS, NAMLEN, IDLEN, BUFLEN,
1 MSGLEN, MAXNBR, MAXGVN
parameter (MAXPRS = 300, NAMLEN = 20, IDLEN = 3, BUFLEN = 60,
1 MSGLEN = 40, MAXNBR = 20, MAXGVN = 3)
character NULLID*(IDLEN)
parameter (NULLID = '000-)
integer PARENT, CHILD, SPOUSE, SIBLNG,
1 UNCLE, NEPHEW, COUSIN, NULLRL
parameter (PARENT = 1, CHILD = 2, SPOUSE = 3, SIBLNG = 4,
1 UNCLE = 5, NEPHEW = 6, COUSIN = 7, NULLRL = 8)
c These common blocks hold the PERSON array, which is global to
$c$ the entire program.
common /PERNUM/ NBRCNT, NBRDEX, NBREDG, DSTSRC, PATHPR, 1 EDGPRD, RCHST, DSCGEN, NUMPER
common /PERCHR/ NAME, IDENT, GENDER, RELID, DSCID

```
c The following data items constitute the PERSON array, which
c is the central repository of information about inter-relationships.
c static information - filled from PEOPLE file character*(NAMLEN) NAME (MAXPRS) character*(IDLEN) IDENT (MAXPRS) character*1 GENDER (MAXPRS)
c IDENTs of immediate relatives - father, mother, spouse character*(IDLEN) RELID (MAXPRS, MAXGVN)
c pointers to immediate neighbors in graph
\begin{tabular}{lll} 
integer & NBRCNT & (MAXPRS) \\
integer & NBRDEX & (MAXPRS, MAXNBR) \\
integer & NBREDG & (MAXPRS, MAXNBR)
\end{tabular}
c data used when traversing graph to resolve user request: real DSTSRC (MAXPRS) integer PATHPR (MAXPRS) integer EDGPRD (MAXPRS) integer RCHST (MAXPRS)
c data used to compute common genetic material character*(IDLEN) DSCID (MAXPRS) real DSCGEN (MAXPRS)
c NUMPER keeps track of the actual number of persons integer NUMPER
c STACK is common to the routines which calculate genetic overlap.
c It is used to implement recursive traversal of the ancestor trees.
integer STKSIZ
parameter (STKSIZ = 50)
common /STACK/ PROPTN, CONTRB, COUNTD, PERDEX, NXTNBR, 1 STKPTR
```

real PROPTN (STKSIZ)
real CONTRB (STKSIZ)
real COUNTD (STKSIZ)
integer PERDEX (STKSIZ)
integer NXTNBR (STKSIZ)
integer STKPTR

```
c *** end of declarations for common data ***
\begin{tabular}{ll} 
real & COMPRP \\
character*(IDLEN) & MTCHID \\
integer & CHKDEX
\end{tabular}
```

    CHKDEX = PERDEX (STKPTR)
    if (NXTNBR (STKPTR) .eq. 0) then
        NXTNBR (STKPTR) = 1
    if (DSCID (CHKDEX) .eq. MTCHID) then
    c Increment COMPRP by the contribution of this
C
c
100
c
common ancestor, but discount for the contribution
of less remote ancestors already counted.
CONTRB (STKPTR) = DSCGEN (CHKDEX) * PROPTN (STKPTR)
COMPRP = COMPRP + CONTRB (STKPTR) - COUNTD (STKPTR)
else
CONTRB (STKPTR) = 0.0
end if
end if
do 100 THSNBR = NXTNBR (STKPTR), NBRCNT (CHKDEX)
if (NBREDG (CHKDEX, THSNBR) .eq. PARENT) goto 101
continue
continue
if (THSNBR .gt. NBRCNT (CHKDEX)) then
no more ancestors from this person
STKPTR = STKPTR - 1
else
set up for next ancestor
NXTNBR (STKPTR) = THSNBR + 1
STKPTR = STKPTR + 1
PERDEX (STKPTR) = NBRDEX (CHKDEX, THSNBR)
NXTNBR (STKPTR) = 0
PROPTN (STKPTR) = PROPTN (STKPTR - 1) / 2.0
COUNTD (STKPTR) = CONTRB (STKPTR - 1) / 4.0
end if
end

```

\subsection*{7.0 PASCAL}

User-defined identifiers are written in mixed upper and lower case, rather than all upper-case, because Pascal provides no separator character, such as "-" or " " for identifiers. Therefore, upper-case letters are used for readability, e.g., EdgeToPredecessor is used in Pascal where EDGE_TO PREDECESSOR is used in most of the other languages.
```

program Relate (input, output, People);
const
MaxPersons = 300;
NameLength = 20;
{ every Person has a unique 3-digit Identifier }
IdentifierLength = 3;
BufferLength = 60;
RequestOk =
Request OK
RequestToStop =
stop
type
IdentifierRange = 1..IdentifierLength;
BufferRange = 1..BufferLength;
NameRange = 1..NameLength;
DigitType = '0'..'`';
NameType = packed array [NameRange] of char;
BufferType = packed array [BufferRange] of char;
MessageType = packed array [1..40] of char;
IdentifierType = array [IdentifierRange] of DigitType;
{ each Person's record in the file identifies at most three
others directly related: father, mother, and spouse }
GivenIdentifiers = (FatherIdent, MotherIdent, SpouseIdent);
RelativeArray = array [GivenIdentifiers] of IdentifierType;
Counter = 0..maxint;
{ this is the format of records in the file to be read in }
FilePersonRecord = record
Name : NameType;
Identifier : IdentifierType;
{ 'M' for Male and 'F' for Female }
Gender : char;
RelativeIdentifier : RelativeArray
end;

```
```

    IndexType = 0..MaxPersons;
    GenderType = (Male, Female);
    RelationType = (Parent, Child, Spouse, Sibling, Uncle,
Nephew, Cousin, NullRelation);
{ directed edges in the graph are of a given type }
EdgeType = Parent..Spouse;
{ A node in the graph (= Person) has either already been reached,
is immediately adjacent to those reached, or farther away. }
ReachedType = (Reached, Nearby, NotSeen);
{ each Person has a linked list of adjacent nodes, called neighbors }
NeighborPointer = `NeighborRecord;
NeighborRecord = record
NeighborIndex : IndexType;
NeighborEdge : EdgeType;
NextNeighbor : NeighborPointer
end;
{ All Relationships are captured in the directed graph of which
each record is a node. }
PersonRecord = record
{ static information - filled from People file: }
Name : NameType;
Identifier : IdentifierType;
Gender : GenderType;
{ Identifiers of immediate relatives - father, mother, spouse }
RelativeIdentifier : RelativeArray;
{ head of linked list of adjacent nodes }
NeighborListHeader : NeighborPointer;
{ data used when traversing graph to resolve user request: }
DistanceFromSource : real;
PathPredecessor : IndexType;
EdgeToPredecessor : EdgeType;
ReachedStatus : ReachedType;
{ data used to compute common genetic material }
DescendantIdentifier : IdentifierType;
DescendantGenes : real
end;
var
{ The Person array is the central repository of information
about inter-relationships. }
Person : array [IndexType] of PersonRecord;
{ These variables are used when establishing the Person array
from the People file.}
People : file of FilePersonRecord;
Current, Previous, NumberOf Persons
: IndexType;
IdentifierIndex : IdentifierRange;
PreviousIdent, CurrentIdent, NullIdent
: IdentifierType;
Relationship : GivenIdentifiers;
RelationLoopDone : boolean;

```
```

    { These variables are used to accept and resolve requests for
        Relationship information. }
    BufferIndex, SemicolonLocation
                            : BufferRange;
    Request Buffer : BufferType;
    PersonlIdent, Person2Ident
                            : NameType;
    Person1Found, Person2Found
                            : Counter;
    ErrorMessage : MessageType;
PersonlIndex, Person2Index
: IndexType;
function IdentsEqual (Identa, Identb: IdentifierType) : boolean;
{ Determines whether two numeric Person-Identifiers are equal.
A function is necessary because the "=- operator does not
work for arrays of anything but char. }
var
Index : 1..IdentifierLength;
begin
IdentsEqual := true;
for Index := 1 to IdentifierLength do
if Identa [Index] <> Identb [Index] then
IdentsEqual := false
end; { IdentsEqual }

```
```

procedure LinkRelatives (FromIndex : IndexType;
Relationship : GivenIdentifiers;
ToIndex : IndexType);
{ establishes cross-indexing between immediately related Persons. }
procedure LinkOneWay (FromIndex : IndexType;
ThisEdge : EdgeType;
ToIndex : IndexType);
{ Establishes the NeighborRecord from one Person to another }
var
NewNeighbor : NeighborPointer;
begin
new (NewNeighbor);
with NewNeighbor^ do
begin
NeighborIndex := ToIndex;
NeighborEdge := ThisEdge;
NextNeighbor := Person [FromIndex] . NeighborListHeader
end;
Person [FromIndex] . NeighborListHeader := NewNeighbor
end;
begin { execution of LinkRelatives }
if Relationship = SpouseIdent then
begin
LinkOneWay (FromIndex, Spouse, ToIndex);
LinkOneWay (Toindex, Spouse, FromIndex)
end
else { Relationship is Mother or Father }
begin
LinkOneWay (FromIndex, Parent, Toindex);
LinkOneWay (ToIndex, Child, FromIndex)
end
end; { LinkRelatives }
procedure PromptAndRead;
{ Issues prompt for user-request, reads in request,
blank-fills buffer, and skips to next line of input. }
var
BufferIndex : BufferRange;
begin
writeln (' ");
writeln (` ---------------------------------------------------------
writeln (' Enter two person-identifiers (name or number), ');
writeln (" separated by semicolon. Enter "stop" to stop.");
for BufferIndex := 1 to BufferLength do
if eoln(input) then
RequestBuffer [BufferIndex] := - -
else
read (input, RequestBuffer [BufferIndex] );
readln(input)
end; { PromptAndRead }

```
```

procedure CheckRequest (var RequestStatus : MessageType;
var SemicolonLocation : BufferRange);
{ Performs syntactic check on request in buffer. }
var
BufferIndex : BufferRange;
SemicolonCount : Counter;
Person1FieldExists, Person2FieldExists
: boolean;
begin
RequestStatus := RequestOk;
PersonlFieldExists := false;
Person2FieldExists := false;
SemicolonCount := 0;
for BufferIndex := 1 to BufferLength do
if RequestBuffer [BufferIndex] <> ' " then
if RequestBuffer [BufferIndex] = ';' then
begin
SemicolonLocation := BufferIndex;
SemicolonCount := SemicolonCount + 1
end
else { Check for non-blanks before/after semicolon. }
if SemicolonCount < 1 then
Person1FieldExists := true
else
Person2FieldExists := true;
{ set RequestStatus, based on results of scan of RequestBuffer. }
if SemicolonCount <> 1 then
RequestStatus := "must be exactly one semicolon.
else
if not Person1FieldExists then
RequestStatus := 'null field preceding semicolon.
else
if not Person2FieldExists then
RequestStatus := 'null field following semicolon.
end; { CheckRequest }
procedure BufferToPerson (var PersonId : NameType;
StartLocation, StopLocation : BufferRange);
{ fills in the PersonId from the designated portion
of the RequestBuffer. }
var
BufferIndex : 1..61; { cannot say "BufferLength + 1" }
PersonIndex : NameRange;
begin
BufferIndex := StartLocation;
while RequestBuffer [BufferIndex] = - do
BufferIndex := BufferIndex + 1;
for PersonIndex := 1 to NameLength do
if BufferIndex > StopLocation then
PersonId [PersonIndex] := " -
else
begin
PersonId [PersonIndex] := RequestBuffer [BufferIndex];
BufferIndex := BufferIndex + 1
end
end; { BufferToPerson }

```
```

procedure SearchForRequestedPersons (Person1Ident, Person2Ident : NameType;
var Person1Index, Person2Index : IndexType;
var Person1Found, Person2Found : Counter);
{ SearchForRequestedPersons scans through the Person array,
looking for the two requested persons. Match may be by name
or unique identifier-number. }
var
Current : IndexType;
ThisIdent : NameType;
IdentifierIndex : IdentifierRange;
begin
Person1Found := 0;
Person2Found := 0;
ThisIdent := - ';
for Current := 1 to NumberOfPersons do
with Person [Current] do
begin
{ ThisIdent contains Current Person's numeric Identifier
left-justified, padded with blanks. }
for IdentifierIndex := 1 to IdentifierLength do
ThisIdent [IdentifierIndex] := Identifier [IdentifierIndex];
{ allow identification by name or number. }
if (Person1Ident = ThisIdent) or (Person1Ident = Name) then
begin
Person1Found := Person1Found + 1;
Person1Index := Current
end;
if (Person2Ident = ThisIdent) or (Person2Ident = Name) then
begin
Person2Found := Person2Found + 1;
Person2Index := Current
end
end { with Person [Current] }
end; { SearchForRequestedPersons }
procedure FindRelationship (TargetIndex, SourceIndex : IndexType);
{ Finds shortest path (if any) between two Persons and
determines their Relationship based on immediate relations
traversed in path. Person array simulates a directed graph,
and algorithm finds shortest path, based on following
weights: Parent-Child edge = 1.0
Spouse-Spouse edge = 1.8 }
var
SearchStatus : (Searching, Succeeded, Failed);
PersonIndex, ThisNode, AdjacentNode, BestNearby Index, LastNearby Index
: IndexType;
NearbyNode : array [IndexType] of IndexType;
ThisEdge : EdgeType;
ThisNeighbor : NeighborPointer;
Relationship : GivenIdentifiers;
MinimalDistance : real;

```
```

procedure ProcessAdjacentNode (BaseNode, NextNode : IndexType;
NextBaseEdge : EdgeType);
{ NextNode is adjacent to last-reached node (= BaseNode).
if NextNode already Reached, do nothing.
If previously seen, check whether path thru base node is
shorter than current path to NextNode, and if so re-link
next to base.
If not previously seen, link next to base node. }
var
WeightThisEdge, DistanceThruBaseNode
: real;
procedure LinkNextNodeToBaseNode;
{ link next to base by re-setting its predecessor Index to
point to base, note type of edge, and re-set distance
as it is through base node. }
begin { execution of LinkNextNodeTo BaseNode }
with Person [NextNode] do
begin
DistanceFromSource := DistanceThruBaseNode;
PathPredecessor := BaseNode;
EdgeToPredecessor := NextBaseEdge
end
end; { LinkNextNodeToBaseNode }
begin { execution of ProcessAdjacentNode }
with Person [NextNode] do
if ReachedStatus <> Reached then
begin
if NextBaseEdge = Spouse then
WeightThisEdge := 1.8
else
WeightThisEdge := 1.0;
DistanceThruBaseNode := WeightThisEdge +
Person [BaseNode] . DistanceFromSource;
if ReachedStatus = NotSeen then
begin
ReachedStatus := Nearby;
LastNearbyIndex := LastNearbyIndex + 1;
NearbyNode [LastNearbyIndex] := NextNode;
LinkNext NodeToBaseNode
end
else { ReachedStatus = Nearby }
if DistanceThruBaseNode < DistanceFromSource then
LinkNextNodeToBaseNode;
end { if ReachedStatus <> Reached }
end; { ProcessAd jacentNode }

```
```

procedure ResolvePathToEnglish;
\{ ResolvePathToEnglish condenses the shortest path to a
series of Relationships for which there are English
descriptions. \}
type
\{ Key Persons are the ones in the Relationship path which remain
after the path is condensed. \}
SiblingType $=$ (Step, Half, Full);
KeyPersonRecord $=$ record
PersonIndex : IndexType;
GenerationGap : Counter;
Proximity : SiblingType;
case RelationToNext : RelationType of
Parent, Child, Spouse, Sibling, Uncle, Nephew, NullRelation
: ();
Cousin : (CousinRank : Counter)
end;
var
\{ these variables are used to condense the path \}
KeyPerson : array [IndexType] of KeyPersonRecord;
KeyRelation, LaterKeyRelation, PrimaryRelation, NextPrimaryRelation
: RelationType;
GenerationCount : Counter;
KeyIndex, LaterKeyIndex, PrimaryIndex
: IndexType;
AnotherElementPossible : boolean;
function FullSibling (Index1, Index2 : IndexType) : boolean;
\{ Determines whether two Persons are full siblings, i.e.,
have the same two Parents. \}
var
IdentIndex : 1..IdentifierLength;
begin
with Person [Index1] do
FullSibling :=
(not IdentsEqual (RelativeIdentifier [FatherIdent], NullIdent)) and
(not IdentsEqual (RelativeIdentifier [MotherIdent], Nullídent)) and
(IdentsEqual (RelativeIdentifier [FatherIdent],
Person [Index2] . RelativeIdentifier [FatherIdent] )) and
(IdentsEqual (RelativeIdentifier [MotherIdent],
Person [Index2] . RelativeIdentifier [MotherIdent] ))
end; \{ FullSibling \}
procedure CondenseKeyPersons (AtIndex : IndexType; GapSize : Counter);
\{ CondenseKeyPersons condenses superfluous entries from the
KeyPerson array, starting at AtIndex. \}
var
ReceiveIndex, SendIndex : IndexType;
begin
ReceiveIndex := AtIndex;
repeat
ReceiveIndex := ReceiveIndex +1 ;
SendIndex $:=$ ReceiveIndex + GapSize;
KeyPerson [ReceiveIndex] := KeyPerson [SendIndex];
until KeyPerson [SendIndex] . RelationToNext = NullRelation
end; \{ CondenseKeyPersons \}

```
```

procedure DisplayRelation (FirstIndex, LastIndex, PrimaryIndex
: IndexType);
{ DisplayRelation takes 1, 2, or 3 adjacent elements in the
condensed table and generates the English description of
the relation between the first and last + l elements. }
var
Inlaw : boolean;
ThisProximity : SiblingType;
ThisGender : GenderType;
SuffixIndicator : 0..9;
FirstRelation, LastRelation, PrimaryRelation
: RelationType;
ThisGenerationGap, ThisCousinRank
: Counter;
begin { execution of DisplayRelation }
FirstRelation := KeyPerson [FirstIndex] . RelationToNext;
LastRelation := KeyPerson [LastIndex] . RelationToNext;
PrimaryRelation := KeyPerson [PrimaryIndex] . RelationToNext;
{ set ThisProximity }
if ((PrimaryRelation = Parent) and (FirstRelation = Spouse)) or
((PrimaryRelation = Child) and (LastRelation = Spouse))
then
ThisProximity := Step
else
if PrimaryRelation in
[Sibling, Uncle, Nephew, Cousin]
then
ThisProximity := KeyPerson [PrimaryIndex] . Proximity
else
ThisProximity := Full;
{ set ThisGenerationGap }
if PrimaryRelation in [Parent, Child, Uncle, Nephew, Cousin]
then
ThisGenerationGap := KeyPerson [PrimaryIndex] . GenerationGap
else
ThisGenerationGap := 0;
{ set Inlaw }
Inlaw := false;
if (FirstRelation = Spouse) and
(PrimaryRelation in [Sibling, Child, Nephew, Cousin] )
then
Inlaw := true;
if (LastRelation = Spouse) and
(PrimaryRelation in [Sibling, Parent, Uncle, Cousin] )
then
Inlaw := true;
{ set ThisCousinRank }
if PrimaryRelation = Cousin then
ThisCousinRank := KeyPerson [PrimaryIndex] . CousinRank
else
ThisCousinRank := 0;

```
```

\{ parameters are set - now generate display. \}
write (- -, Person [KeyPerson [FirstIndex] • PersonIndex] . Name,
- is ${ }^{-}$);
if PrimaryRelation in [Parent, Child, Uncle, Nephew] then
begin \{ write generation-qualifier \}
if ThisGenerationGap $>=3$ then
begin
write (-great');
if ThisGenerationGap > 3 then
write ( ${ }^{*-}$, ThisGenerationGap - 2 : 1);
write (--)
end;
if ThisGenerationGap $>=2$ then
write ('grand-")
end
else
if (PrimaryRelation $=$ Cousin) and (ThisCousinRank $>1$ ) then
begin
write (ThisCousinRank : 1);
SuffixIndicator := ThisCousinRank mod 10;
case SuffixIndicator of
1 : write ( ${ }^{\prime}$ st -);
2 : write ('nd ");
3 : write ('rd ");
$0,4,5,6,7,8,9$
: write ( ${ }^{-}$th ${ }^{-}$)
end
end;
if ThisProximity $=$ Step then
write ('step-')
else
if ThisProximity $=$ Half then
write ( ${ }^{\text {half }}{ }^{-}$);
ThisGender := Person [KeyPerson [FirstIndex] . PersonIndex] • Gender;
case PrimaryRelation of
Parent : if ThisGender = Male then write ('father")
else write ('mother");
Child : if ThisGender = Male then write ( ${ }^{\prime}$ son ${ }^{-}$)
else write ('daughter");
Spouse $\quad:$ if ThisGender $=$ Male then write ( ${ }^{\prime}$ husband ${ }^{-}$)
else write ("wife");
Sibling : if ThisGender = Male then write ('brother-)
else write ('sister-);
Uncle : if ThisGender = Male then write ('uncle-)
else write ('aunt');
Nephew : if ThisGender = Male then write ('nephew')
else write ('niece");
Cousin : write ('cousin-);
NullRe1ation : write ('null')
end; \{ case \}

```
```

    if Inlaw then
    write (`-in-law');
    if (PrimaryRelation = Cousin) and (ThisGenerationGap > 0) then
    if ThisGenerationGap > 1 then
                write (` ', ThisGenerationGap : 1, ' times removed")
            else
                write (` once removed`);
        writeln (` of`)
    end; { DisplayRelation }
    begin { execution of ResolvePathToEnglish }
writeln (` Shor*est path between identified persons: ');     ThisNode := Target Index;     KeyIndex := 1;     { Display path and initialize KeyPerson array from path elements. }     while ThisNode <> SourceIndex do         with Person [ThisNode] do         begin         write (` `, Name, ` is `);         case EdgeToPredecessor of             Parent : writeln ('parent of`);
Child : writeln (`child of }\mp@subsup{}{}{-}\mathrm{ );             Spouse : writeln (`spouse of`)         end;             KeyPerson [KeyIndex] . PersonIndex := ThisNode;             KeyPerson [KeyIndex] . RelationToNext := EdgeToPredecessor;             if EdgeToPredecessor = Spouse then                     KeyPerson [KeyIndex] . GenerationGap := 0             else { Parent or Child }                         KeyPerson [KeyIndex] . GenerationGap := 1;             KeyIndex := KeyIndex + 1;             ThisNode := PathPredecessor             end;     writeln(` ", Person [ThisNode] . Name);
KeyPerson [KeyIndex] . PersonIndex := ThisNode;
KeyPerson [KeyIndex] . RelationToNext := Nul1Relation;
KeyPerson [KeyIndex + 1] . RelationToNext := NullRelation;

```
```

\{ Resolve Child-Parent and Child-Spouse-Parent relations
to Sibling relations. \}
KeyIndex := 1;
while KeyPerson [KeyIndex] . RelationToNext 〈〉 NullRelation do
with KeyPerson [KeyIndex] do
begin
if RelationToNext $=$ Child then
begin
LaterKeyRelation := KeyPerson [KeyIndex + 1] . RelationToNext;
if LaterKeyRelation = Parent then
\{ found either full or half siblings \}
begin
RelationToNext := Sibling;
if FullSibling (PersonIndex,
KeyPerson [KeyIndex + 2] . PersonIndex)
then
Proximity := Full
else
Proximity := Half;
CondenseKeyPersons (Key Index, 1)
end \{ processing of full/half siblings \}
else
if (LaterKeyRelation $=$ Spouse) and
(KeyPerson [KeyIndex + 2] . RelationToNext = Parent)
then \{ found step-siblings \}
begin
RelationToNext := Sibling;
Proximity := Step;
CondenseKeyPersons (KeyIndex, 2)
end \{ processing of step-siblings \}
end; $\quad$ if RelationToNext $=$ Child \}
KeyIndex := Key Index +1
end; \{ with KeyPerson [KeyIndex] \}
\{ Resolve Child-Child-... and Parent-Parent-... relations to
direct descendant or ancestor relations. \}
KeyIndex := 1;
while KeyPerson [KeyIndex] . RelationToNext 〈> NullRelation do
with KeyPerson [KeyIndex] do
begin
if (RelationToNext $=$ Child) or (RelationToNext $=$ Parent) then
begin
LaterKeyIndex := KeyIndex + 1;
while KeyPerson [LaterKeyIndex] . RelationToNext =
RelationToNext do
LaterKeyIndex := LaterKeyIndex + 1;
GenerationCount := LaterKeyIndex - KeyIndex;
if GenerationCount > 1 then
begin \{ compress generations \}
GenerationGap := GenerationCount;
CondenseKeyPersons (KeyIndex, GenerationCount - 1)
end
end; \{ if RelationToNext $=$ Child or Parent $\}$
KeyIndex := KeyIndex + 1
end; \{ with KeyPerson [KeyIndex] \}

```
```

{ Resolve Child-Sibling-Parent to Cousin,
Child-Sibling to Nephew,
Sibling-Parent to Uncle.}
KeyIndex := 1;
while KeyPerson [KeyIndex] . RelationToNext <> NullRelation do
with KeyPerson [KeyIndex] do
begin
LaterKeyRelation := KeyPerson [KeyIndex + 1] . RelationToNext;
if (RelationToNext = Child) and
(LaterKeyRelation = Sibling)
then { Cousin or Nephew }
if KeyPerson [KeyIndex + 2] . RelationToNext = Parent then
{ found Cousin }
begin
RelationToNext := Cousin;
Proximity := KeyPerson [KeyIndex + l] . Proximity;
if GenerationGap < KeyPerson [KeyIndex + 2] . GenerationGap
then
CousinRank := GenerationGap
else
CousinRank := KeyPerson [KeyIndex + 2] . GenerationGap;
GenerationGap := abs (GenerationGap -
KeyPerson [KeyIndex + 2] . GenerationGap);
CondenseKeyPersons (KeyIndex, 2)
end
else { found Nephew }
begin
RelationToNext := Nephew;
Proximity := KeyPerson [KeyIndex + 1] . Proximity;
CondenseKeyPersons (KeyIndex, 1)
end
else { not Cousin or Nephew }
if (RelationToNext = Sibling) and (LaterKeyRelation = Parent)
then { found Uncle }
begin
RelationToNext := Uncle;
GenerationGap := KeyPerson [KeyIndex + 1] . GenerationGap;
CondenseKeyPersons (KeyIndex, 1)
end;
KeyIndex := KeyIndex + 1
end; { with KeyPerson [KeyIndex] }

```
```

    { Loop below will pick out valid adjacent strings of elements
    to be displayed. KeyIndex points to first element,
    LaterKeyIndex to last element, and PrimaryIndex to the
    element which determines the primary English word to be used.
    Associativity of adjacent elements in condensed table
    is based on English usage. }
    KeyIndex := 1;
    writeln (` Condensed path:`);
    while KeyPerson [KeyIndex] . RelationToNext <> NullRelation do
    begin
    KeyRelation := KeyPerson [KeyIndex] . RelationToNext;
    LaterKeyIndex := KeyIndex;
    PrimaryIndex := KeyIndex;
    if KeyPerson [KeyIndex + 1] . RelationToNext <> Nul1Relation then
        begin { seek multi-element combination }
        AnotherElementPossible := true;
        if KeyRelation = Spouse then
            begin
            LaterKeyIndex := LaterKeyIndex + 1;
            PrimaryIndex := LaterKeyIndex;
            if (KeyPerson [LaterKeyIndex] . RelationToNext = Sibling) or
                    (KeyPerson [LaterKeyIndex] . RelationToNext = Cousin)
            then { Nothing can follow Spouse-Sibling or Spouse-Cousin }
            AnotherElementPossible := false
            end;
        { PrimaryIndex is now correctly set. Next if-statement
                determines if a following Spouse relation should be
                appended to this combination or left for the next
            combination. }
        if AnotherElementPossible and
            (KeyPerson [PrimaryIndex + 1] . RelationToNext = Spouse)
            { Only a Spouse can follow a Primary }
        then
            begin { check primary preceding and following Spouse. }
            PrimaryRelation :=
                KeyPerson [PrimaryIndex] . RelationToNext;
            NextPrimaryRelation :=
                KeyPerson [PrimaryIndex + 2] . RelationToNext;
            if (NextPrimaryRelation in [Nephew, Cousin, NullRelation] )
                    or (PrimaryRelation = Nephew)
                    or ( ( PrimaryRelation in [Sibling, Parent] )
                    and (NextPrimaryRelation <> Uncle ) )
            then { append following Spouse with this combination. }
                    LaterKeyIndex := LaterKeyIndex + 1
            end { check primary preceding and following Spouse }
        end; { multi-element combination }
    DisplayRelation (KeyIndex, LaterKeyIndex, PrimaryIndex);
    KeyIndex := LaterKeyIndex + 1
    end; { while }
    writeln (' ', Person [KeyPerson [KeyIndex] . PersonIndex] . Name)
    end; , { ResolvePathToEnglish }

```
```

procedure ComputeCommonGenes (Index1, Index2 : IndexType);
{ ComputeCommonGenes assumes that each ancestor contributes
half of the genetic material to a Person. It finds common
ancestors between two Persons and computes the expected
value of the Proportion of common material. }
var
CommonProportion : real;
procedure ZeroProportion (ZeroIndex : IndexType);
{ ZeroProportion recursively seeks out all ancestors and
zeros them out. }
var
ThisNeighbor : NeighborPointer;
begin
with Person [ZeroIndex] do
begin
DescendantGenes := 0.0;
ThisNeighbor := NeighborListHeader
end;
while ThisNeighbor <> nil do
with ThisNeighbor^ do
begin
if NeighborEdge = Parent then
ZeroProportion (NeighborIndex);
ThisNeighbor := NextNeighbor
end { with }
end; { ZeroProportion }
procedure MarkProportion (Marker : IdentifierType;
Proportion : real; Marked Index : IndexType);
{ MarkProportion recursively seeks out all ancestors and
marks them with the sender's Proportion of shared
genetic material. This Proportion is diluted by one-half
for each generation. }
var
ThisNeighbor : NeighborPointer;
begin
with Person [MarkedIndex] do
begin
DescendantIdentifier := Marker;
DescendantGenes := DescendantGenes + Proportion;
ThisNeighbor := NeighborListHeader
end;
while ThisNeighbor <> nil do
with ThisNeighbor^ do
begin
if NeighborEdge = Parent then
MarkProportion (Marker, Proportion / 2.0,
NeighborIndex );
ThisNeighbor := NextNeighbor
end
end; { MarkProportion }

```
```

    procedure CheckCommonProportion
            (var CommonProportion : real;
                            MatchIdentifier : IdentifierType;
                            Proportion : real;
                            AlreadyCounted : real;
                    CheckIndex : IndexType);
    { CheckCommonProportion searches all the ancestors of
        CheckIndex to see if any have been marked, and if so
        adds the appropriate amount to CommonProportion. }
    var
    ThisNeighbor : NeighborPointer;
    ThisContribution : real;
    begin
        with Person [CheckIndex] do
        begin
        if IdentsEqual (DescendantIdentifier, MatchIdentifier) then
            begin
            { Increment CommonProportion by the contribution of
                this common ancestor, but discount for the contribution
                    of less remote ancestors already counted. }
            ThisContribution := DescendantGenes * Proportion;
            CommonProportion := CommonProportion +
                        ThisContribution - AlreadyCounted
            end
        else
            ThisContribution := 0.0;
        ThisNeighbor := NeighborListHeader
        end; { with Person [CheckIndex] }
    while ThisNeighbor <> nil do
        with ThisNeighbor^ do
            begin
            if NeighborEdge = Parent then
                    CheckCommonProportion (CommonProportion,
                    MatchIdentifier, Proportion / 2.0,
                    ThisContribution / 4.0,
                    NeighborIndex );
            ThisNeighbor := NextNeighbor
            end
    end; { CheckCommonProportion }
    begin { ComputeCommonGenes }
{ First zero out all ancestors to allow adding. This is necessary
because there might be two paths to an ancestor. }
ZeroProportion (Indexl);
{ now mark with shared Proportion }
MarkProportion ( Person [Indexl] . Identifier, 1.0, Indexl);
CommonProportion := 0.0;
CheckCommonProportion (CommonProportion,
Person [Index1] . Identifier, 1.0, 0.0, Index2);
writeln (' Proportion of common genetic material = ',
CommonProportion : 12)
end; { ComputeCommonGenes }

```
```

begin { execution of FindRelationship }
{ initialize Person-array for processing -
mark all nodes as not seen }
for PersonIndex := 1 to NumberOfPersons do
Person [PersonIndex] . ReachedStatus := NotSeen;
{ mark source node as Reached }
ThisNode := SourceIndex;
with Person [ThisNode] do
begin
ReachedStatus := Reached;
DistanceFromSource := 0.0
end;
{ no Nearby nodes exist yet }
LastNearbyIndex := 0;
if ThisNode = TargetIndex then
SearchStatus := Succeeded
else
SearchStatus := Searching;
{ Loop keeps processing closest-to-source, unreached node
until target Reached, or no more connected nodes. }
while SearchStatus = Searching do
begin
{ Process all nodes adjacent to ThisNode }
ThisNeighbor := Person [ThisNode] . NeighborListHeader;
while ThisNeighbor <> nil do
with ThisNeighbor^ do
begin
ProcessAdjacentNode (ThisNode, NeighborIndex, NeighborEdge);
ThisNeighbor := NextNeighbor
end;
{ All nodes adjacent to ThisNode are set. Now search for
shortest-distance unreached (but Nearby) node to process next. }
if LastNearbyIndex = 0 then
SearchStatus := Failed
else
begin
MinimalDistance := 1.0e+18;
for PersonIndex := 1 to LastNearbyIndex do
with Person [NearbyNode [PersonIndex]] do
if DistanceFromSource < MinimalDistance then
begin
BestNearbyIndex := PersonIndex;
MinimalDistance := DistanceFromSource
end;
{ Establish new ThisNode }
ThisNode := NearbyNode [BestNearbyIndex];
{ change ThisNode from being Nearby to Reached }
Person [ThisNode] . ReachedStatus := Reached;
{ remove ThisNode from Nearby list }
NearbyNode [BestNearbyIndex] := NearbyNode [LastNearbyIndex];
LastNearbyIndex := LastNearbyIndex - 1;
if ThisNode = TargetIndex then
SearchStatus := Succeeded
end { determination of next node to process }
end; { while SearchStatus = Searching }

```
\{ Shortest path between Persons now established. Next task is to translate path to English description of Relationship. \}
if SearchStatus = Failed then
writeln ( \({ }^{-}\), Person [TargetIndex] . Name, \({ }^{-}\)is not related to \({ }^{-}\), Person [SourceIndex] . Name)
else \{ success - parse path to find and display Relationship \} begin ResolvePathToEng1ish; ComputeCommonGenes (SourceIndex, TargetIndex) end
end; \{ FindRelationship \}
```

{ *** execution of main sequence begins here *** }

```
begin
    for Identifier Index \(:=1\) to IdentifierLength do
        NullIdent [IdentifierIndex] : = \({ }^{-} 0^{-}\);
    reset (People);
    \{ Current location in array being filled \}
    Current : = 0;
    \{ This loop reads in the People file and constructs the Person
        array from it (one Person \(=\) one record \(=\) one array entry).
        As records are read in, links are constructed to represent the
        Parent-Child or Spouse relationship. The array then implements
        a directed graph which is used to satisfy subsequent user
        requests. The file is assumed to be correct - no validation
        is performed on it. \}
    while not eof(People) do
        begin
        Current := Current+1;
        with Person [Current] do
            begin
            \{ copy direct information from file to array \}
            Name \(\quad:=\) People^ . Name;
            Identifier \(:=\) People^. Identifier;
            if People^. Gender \(={ }^{\prime} M^{\prime}\) then
                Gender := Male
            else
                Gender := Female;
            RelativeIdentifier := People^ . RelativeIdentifier;
            \{ Location of adjacent persons as yet undetermined \}
            NeighborListHeader \(:=\) nil;
            \{ Descendants as yet undetermined. \}
            DescendantIdentifier \(:=\) NullIdent;
            CurrentIdent := Identifier;
```

    \{ Compare this Person against all previously entered Persons
    to search for Relationships. \}
    for Previous := 1 to (Current-1) do
        begin
        PreviousIdent := Person [Previous] . Identifier;
        RelationLoopDone \(:=\) false;
        Relationship := FatherIdent;
        \{ Search for father, mother, or spouse Relationship in
        either direction between this and previous Person.
        Assume at most one Relationship exists. \}
        repeat
            if IdentsEqual (RelativeIdentifier [Relationship],
                                    PreviousIdent) then
            begin
            LinkRelatives (Current, Relationship, Previous);
            RelationLoopDone := true
            end
        else
            if IdentsEqual (CurrentIdent,
                    Person [Previous] . RelativeIdentifier [Relationship])
            then
                    begin
                    LinkRelatives (Previous, Relationship, Current);
                    RelationLoopDone := true
                    end;
        if Relationship < SpouseIdent then
            Relationship := succ(Relationship)
        else
            RelationLoopDone := true;
        until RelationLoopDone
        end; \{ for Previous \}
        get(People)
        end \{ with Person [Current] \}
    end; \{ while not eof(People) \}
NumberOfPersons := Current;
\{ Person array is now loaded and edges between immediate relatives
(Parent-Child or Spouse-Spouse) are established.
While-loop accepts requests and finds Relationship (if any)
between pairs of Persons. \}

```
```

reset(input);
PromptAndRead;
while RequestBuffer <> RequestToStop do
{ The following code retrieves and validates a user request
for the Relationship between two identified Persons. }
begin
CheckRequest (ErrorMessage, SemicolonLocation);
{ Syntax check of request completed. Now either display error
message or search for the two Persons. }
if ErrorMessage = RequestOk then
begin { Request syntactically correct -
search for requested Persons. }
BufferToPerson (Person1Ident, 1, SemicolonLocation - 1);
BufferToPerson (Person2Ident, SemicolonLocation + 1, BufferLength);
SearchForRequestedPersons (Person1Ident, Person2Ident,
Person1Index, Person2Index,
Person1Found, Person2Found);
if (Person1Found = 1) and (Person2Found = 1) then
{ Exactly one match for each Person - proceed to
determine Relationship, if any.}
if PersonlIndex = Person2Index then
begin
write (- -, Person [Person1Index] . Name,
is identical to -);
if Person [Person1Index] . Gender = Male then
writeln(`himself.')                 else                     writeln('herself.')                 end             else                 FindRe1ationship (Person1Index, Person2Index)         else { either not found or more than one found }             begin                 if Person1Found = 0 then                 writeln (` First person not found.')
else
if Person1Found > 1 then
writeln (` Duplicate names for first person - use',                         - numeric identifier.');                 if Person2Found = 0 then                 writeln (' Second person not found.`)
else
if Person2Found > 1 then
writeln (- Duplicate names for second person - use`,                             - numeric identifier.')                 end         end { processing of syntactically legal request }     else         writeln (` Incorrect request format: `, ErrorMessage);     PromptAndRead     end; { while RequestBuffer } writeln (` End of relation-finder.');

```
end.

\subsection*{8.0 PL/I}

In keeping with the general convention of the examples, language-supplied keywords and identifiers are written in lower case in the program. To conform strictly to the PL/I standard, however, programs must use only upper-case letters. In the following program, the logical "Not" operator is represented by the graphic character "~".

RELATE: procedure options (main);
```

/* Begin declaration of global data */
declare
/* Used to index relative array, pointing to immediate relatives */
( FATHER IDENT initial (1),
MOTHER IDENT initial (2),
SPOUSE IDENT initial (3),
/* Use\overline{d}}\mathrm{ as mnemonics to represent basic English-word relationships. */
PARENT initial (1),
CHILD initial (2),
SPOUSE initial (3),
SIBLING initial (4),
UNCLE initial (5),
NEPHEW initial (6),
COUSIN initial (7),
NULL_RELATION initial (8),
/* Used as mnemonics to represent status of nodes during search
for shortest path thru graph. */
REACHED initial (1),
NEARBY initial (2),
NOT SEEN initial (3) )
fixed binary (4,0),
/* Used as mnemonics to represent truth-values */
( TRUE initial ('1`b),         FALSE initial (`0`b))     bit (1),     /* Used to control user requests. */     ( REQUEST_OK character (10) initial (`Request OK`),
REQUEST_TO_STOP character (4) initial ('stop')),
/* Used as mnemonics to represent GENDER */
( MALE initial ('M'),
FEMALE initial (}\mp@subsup{}{}{-}\mp@subsup{F}{}{\prime})\mathrm{ )
character (1);

```
```

declare
/* the PERSON array is the central repository of information
about inter-relationships. */
/* All relationships are captured in the directed graph of which
each record is a node. */
01 PERSON dimension (1:300),
/* static information - filled from PEOPLE file: */
05 NAME character (20),
05 IDENTIFIER picture '999',
0 5 ~ G E N D E R ~ c h a r a c t e r ~ ( 1 ) ,
/* IDENTIFIERs of immediate relatives - father, mother, spouse */
05 RELATIVE_IDENTIFIER (1:3)
picture '999',
/* head of linked list of adjacent nodes */
05 NEIGHBOR_LIST_HEADER pointer,
/* data used when traversing graph to resolve user request: */
0 5 DISTANCE FROM SOURCE float decimal (6),
05 PATH_PRE\overline{DECES}\overline{S}OR fixed binary (10,0),
05 EDGE TO PREDECESSOR fixed binary (4,0),
05 REAC\overline{HED STATUS fixed binary (4,0),}
/* data use\overline{d}}\mathrm{ to compute common genetic material */
05 DESCENDANT IDENTIFIER picture '999^,
0 5 DESCENDANT_GENES float decimal (6);
declare
/* each PERSON has a linked list of adjacent nodes, called neighbors */
O1 NEIGHBOR_RECORD based (NEW_NEIGHBOR),
05 NEIGHBOR_INDEX fixed binary (10,0),
05 NEIGHBOR EDGE fixed binary (4,0),
O5 NEXT_NEIGGHOR pointer;
/* End declaration of global data. */
declare
/* This is the format of records in the file to be read in. */
Ol PEOPLE RECORD,
O5 NAME character (20),
05 IDENTIFIER picture '999`,                 /* 'M' for MALE and 'F' for FEMALE */             0 5 ~ G E N D E R ~ c h a r a c t e r ~ ( 1 ) ,             O5 RELATIVE_IDENTIFIER (1:3) picture '999`;
declare
/* These variables are used when establishing the PERSON array
from the PEOPLE file. */
PEOPLE file record sequential input,
(CURRENT, PREVIOUS, NUMBER OF PERSONS)
fixed binary (10,0),
(PREVIOUS_IDENT, CURRENT_IDENT)
picture '999',
NULL_IDENT picture '999' static initial (000),
RELATIONSHIP fixed binary (4,0),
RELATION_LOOP_DONE bit (1),
END_OF_PEOPLE- bit (1);

```
```

declare
/* These variables are used to accept and resolve requests for
RELATIONSHIP information. */
sysin file record input environment (AREAD),
(BUFFER_INDEX, SEMICOLON LOCATION)
fixed binary (10,0),
REQUEST BUFFER character (60) varying,
(PERSON1_IDENT, PERSON2_IDENT)
character (20),
(PERSON1 FOUND, PERSON2 FOUND)
fixed binary (10,0),
ERROR MESSAGE character (40),
(PERSONN1_INDEX, PERSON2 INDEX)
fixed binary (10,0);
/* This on-block captures exceptions from the following code */
on endfile (PEOPLE)
begin;
END_OF_PEOPLE = TRUE;
end;

```
```

/* *** begin execution of main sequence RELATE *** */
open file (PEOPLE) title (`PEOPLE.DAT`);
END OF PEOPLE = FALSE;
/* This
array from it (one PERSON = one record = one array entry).
As records are read in, links are constructed to represent the
PARENT-CHILD or SPOUSE RELATIONSHIP. The array then implements
a directed graph which is used to satisfy subsequent user
requests. The file is assumed to be correct - no validation
is performed on it. */
read file (PEOPLE) into (PEOPLE RECORD);
READ IN PEOPLE:
do - CURRENT = 1 to 300 while (~ END OF PEOPLE);
/* copy direct information from \overline{file to array */}
PERSON (CURRENT) = PEOPLE RECORD, by name;
/* Location of adjacent persons as yet undetermined. */
PERSON (CURRENT) . NEIGHBOR_LIST HEADER = null();
/* Descendants as yet undetermined */
PERSON (CURRENT) . DESCENDANT IDENTIFIER = NULL_IDENT;
CURRENT_IDENT = PERSON (CURRENTT) . IDENTIFIER;
/* Compare this PERSON against all previously entered PERSONs
to search for RELATIONSHIPs. */
COMPARE TO PREVIOUS:
do \overline{PREVTIOUS = 1 to (CURRENT-1);}
PREVIOUS IDENT = PERSON (PREVIOUS) . IDENTIFIER;
RELATION LOOP DONE = FALSE;
/* Search for father, mother, or spouse relationship in
either direction between this and PREVIOUS PERSON.
Assume at most one RELATIONSHIP exists. */
TRY_ALL_RELATIONSHIPS:
do RELATIONSHIP = FATHER_IDENT to SPOUSE_IDENT
while (~ RELATION_LODOP_DONE);
if PERSON (CURRENT) - RELATTIVE_IDENTIFIER (RELATIONSHIP) =
PREVIOUS_IDENT then
do;
call LINK RELATIVES (CURRENT, RELATIONSHIP, PREVIOUS);
RELATION_LOOP_DONE = TRUE;
end;
else
if CURRENT IDENT =
PERSON (PREVIOUS) . RELATIVE_IDENTIFIER (RELATIONSHIP)
then
do;
call LINK RELATIVES (PREVIOUS, RELATIONSHIP, CURRENT);
RELATION_LOOP_DONE = TRUE;
end;
end TRY ALL RELATIONSHIPS;
end COMPA\overline{RE TO PREVIOUS;}
read file (\overline{PEOPLE) into (PEOPLE RECORD);}
end READ IN PEOPLE;
NUMBER O\overline{F}}\mathrm{ PERSONS = CURRENT - 1;
close \overline{file (PEOPLE);}

```
    /* PERSON array is now loaded and edges between immediate relatives
        (PARENT-CHILD or SPOUSE-SPOUSE) are established.
```

    While-loop accepts requests and finds RELATIONSHIP (if any)
    between pairs of PERSONs. */
    call PROMPT AND READ();
    READ AND PROCESS REQUEST:
do while (REQUEST BUFFER ~ = REQUEST TO STOP);
/* The following code retrieves and validates a user request
for the RELATIONSHIP between two identified PERSONs. */
call CHECK_REQUEST (ERROR_MESSTAGE, SEMICOLON_LOCATION);
/* Syntax check of request completed. Now either display error
message or search for the two PERSONs. */
if ERROR MESSAGE = REQUEST OK then
do; -/* Request syntactically correct -
search for requested PERSONs. */
ca11 BUFFER TO PERSON (PERSON1 IDENT, 1, SEMICOLON LOCATION - 1);
call BUFFER_TO_PERSON (PERSON2_IDENT, SEMICOLON LOCATION + 1,
length (REQUEST BUFFER));
call SEARCH_FOR_REQUESTED_PERSONS (PERSON1_IDENT, PERSON2_IDENT,
PERSON1 INDEX, PERSON2-INDEX,
PERSON1-FOUND, PERSON2_FOUND);
if (PERSON1 FOUND = 1) \& (PERSON2 FOUND = \overline{1}) then
/* Exact\overline{ly} one match for each \overline{PERSON - proceed to}
determine RELATIONSHIP, if any. */
if PERSON1 INDEX = PERSON2 INDEX then
if PERS\overline{ON (PERSON1 INDE\overline{X}) . GENDER = MALE then}
put skip list (`` - || PERSON (PERSON1 INDEX) . NAME ||
- is identical to himself.');
else
put skip list (- - || PERSON (PERSON1_INDEX) . NAME ||
- is identical to herself.`);                 else                         cal1 FIND_RELATIONSHIP (PERSON1_INDEX, PERSON2_INDEX);             else /* either not found or more than one found */                 do;                 if PERSON1_FOUND = 0 then                 put ski\overline{p}}\mathrm{ list (` First person not found.`);                     else                         if PERSONI FOUND > l then                         put ski\overline{p}}\mathrm{ list (` Duplicate names for first person - use` ||                             numeric identifier.');                         if PERSON2_FOUND = 0 then                 put skip list (` Second person not found.`);             else                                     if PERSON2_FOUND > 1 then                             put ski\overline{p}}\mathrm{ list (' Duplicate names for second person - use- ||                                     - numeric identifier.`);
end;
end; /* processing of syntactically legal request */
else
put skip list (- Incorrect request format: - || ERROR_MESSAGE);
call PROMPT AND READ();
end READ_AND \overline{PROCESS_REQUEST;}
put skip list (' End of relation-finder.');
/* End execution of main sequence RELATE

```
procedures under RELATE begin here */
LINK_RELATIVES: procedure (FROM_INDEX, RELATIONSHIP, TO_INDEX)
declare
FROM INDEX fixed binary ( 10,0 ),
RELATIONSHIP fixed binary ( 4,0 ),
TO INDEX fixed binary \((10,0)\);
/* begin execution of LINK RELATIVES */
if RELATIONSHIP = SPOUSE IDENT then
do;
cal1 LINK ONE WAY (FROM INDEX, SPOUSE, TO_INDEX);
ca11 LINK_ONE WAY (TO_INDDEX, SPOUSE, FROM-INDEX);
end;
else /* RELATIONSHIP is mother or father */
do;
call LINK ONE WAY (FROM INDEX, PARENT, TO INDEX);
call LINK_ONE_WAY (TO_INDDEX, CHILD, FROM_INDEX);
end;

LINK ONE WAY: procedure (FROM_INDEX, THIS_EDGE, TO_INDEX);
declare
FROM INDEX fixed binary ( 10,0 ), THIS EDGE fixed binary ( 4,0 ), TO_INDEX fixed binary (10,0);

\section*{declare}

NEW_NEIGHBOR pointer;
/* begin execution of LINK ONE WAY */
allocate NEIGHBOR RECORD set (NEW NEIGHBOR);
NEW NEIGHBOR \(\rightarrow\) NEIGHBOR INDEX = TO INDEX;
NEW NEIGHBOR \(\rightarrow\) NEIGHBOR EDGE \(=\) THĪS EDGE;
NEW_NEIGHBOR \(\rightarrow\) NEXT NEIGBBOR
PERSON (FROM INDEX) • NEIGHBOR LIST HEADER;
PERSON (FROM_INDEX) - NEIGHBOR_LIST_HEADER \(=\) NEW_NEIGHBOR; end LINK ONE WĀY;
end LINK RELATIVES;
PROMPT AND READ: procedure;
/* Issues prompt for user-request, reads in request, blank-fills buffer, and skips to next line of input. */
declare BUFFER INDEX fixed binary ( 10,0 ), SEMICOLON_COUNT fixed binary (4,0);
```

/* begin execution of PROMPT AND READ */
put skip (2) list (' -------------------------------------------------------
put skip list (' Enter two person-identifiers (name or number),');
put skip list (' separated by semicolon. Enter "stop" to stop.");
put skip list (' ');
/* The use of sysin for record-oriented, rather than stream-oriented,
input may not be considered to be standard usage. It is done here
because stream input cannot recognize line boundaries, so as to
read an entire line from the terminal. */
read file (sysin) into (REQUEST BUFFER);
end PROMPT AND READ;
CHECK REQUEST: procedure (REQUEST STATUS, SEMICOLON LOCATION);
/* Performs syntactic check on \overline{request in buffer.**/}
declare
REQUEST STATUS character (40),
SEMICOLONN_LOCATION fixed binary (10,0);
/* begin execution of CHECK REQUEST */
SEMICOLON LOCATION = inde\overline{x (REQUEST BUFFER, ';}\mp@subsup{}{}{`});     if SEMICOL         index (sub̄str (REQUEST_BUFFER, SEMICOLON_LOCATION + 1), `; ') > 0
then
REQUEST STATUS = 'must be exactly one semicolon.`;     else         if before (REQUEST BUFFER, ';`) = ' - then
REQUEST_STATUS = 'null field preceding semicolon.';
else
if after (REQUEST_BUFFER, ';`) = ' ' then
REQUEST_STATUS = 'null field following semicolon.';
else
REQUEST_STATUS = REQUEST_OK;
end CHECK_REQUEST;
BUFFER TO PERSON: procedure (PERSON ID, START LOCATION, STOP_LOCATION);
/* f\overline{i}11\overline{s}}\mathrm{ in the PERSON ID from the designated portion
of the REQUEST BUFFER. */
declare
PERSON_ID character (20),
(START_LOCATION, STOP LOCATION)
fixed biñary (10,0);
declare
FIRST_NON BLANK fixed binary (10,0);
/* begin execution of BUFFER TO PERSON */
do FIRST NON BLANK = START LOC
while - (sub̄str (REQUEST B
end;
PERSON_ID = substr (REQUEST BUFFER, FIRST_NON BLANK,
STOP_LOCATION - FIRST_NON_BLANK + 1);
end BUFFER_TO_PERSON;

```
```

SEARCH FOR REQUESTED_PERSONS: procedure (PERSON1_IDENT, PERSON2_IDENT,
PERSON1 INDEX, PERSON2 INDEX,
PERSON1 FOUND, PERSON2 FOUND);
/* SEARCH FOR REQUESTED PERSONS scans through the PERSON array,
looking for the two $\bar{r}$ requested PERSONs. Match may be by NAME
or unique IDENTIFIER-number . */
declare
(PERSON1_IDENT, PERSON2_IDENT) character (20),
(PERSON1_INDEX, PERSON2_INDEX) fixed binary ( 10,0 ),
(PERSON1_FOUND, PERSON2_FOUND) fixed binary (10,0);
declare
THIS IDENT character (20),
CURRENT fixed binary (10,0);
/* begin execution of SEARCH_FOR_REQUESTED_PERSONS */
PERSON1 FOUND $=0$;
PERSON2_FOUND $=0$;
SCAN ALL PERSONS:
do CUR $\overline{R E N T}=1$ to NUMBER OF PERSONS;
/* THIS IDENT contains $\bar{C} U R \bar{R} E N T$ PERSON's numeric IDENTIFIER
left-justified, padded with blanks. */
THIS_IDENT $=$ PERSON (CURRENT) - IDENTIFIER;
/* allow identification by name or number. */
if (PERSON1 IDENT $=$ THIS IDENT) |
(PERSON1-IDENT $=$ PERSON (CURRENT) $\cdot$ NAME)
then
do;
PERSON1 FOUND $=$ PERSON1 FOUND +1 ;
PERSON1_INDEX = CURRENT;
end;
if (PERSON2 IDENT = THIS IDENT) |
(PERSON2_IDENT $=$ PERSON (CURRENT) $\cdot$ NAME)
then
do;
PERSON2 FOUND $=$ PERSON2 FOUND +1 ;
PERSON2_INDEX = CURRENT;
end;
end SCAN ALL PERSONS;
end SEARCH_FOR_REQUESTED_PERSONS;
/* End of utility procedures under RELATE.

```
```

    FIND RELATIONSHIP does major work of program: determines
    relationship between any two people in PERSON array. */
    FIND RELATIONSHIP: procedure (TARGET INDEX, SOURCE INDEX);
/* Finds shortest path (if any) between two PERSONs and
determines their RELATIONSHIP based on immediate relations
traversed in path. PERSON array simulates a directed graph,
and algorithm finds shortest path, based on following
weights: PARENT-CHILD edge = 1.0
SPOUSE-SPOUSE edge = 1.8 */
declare
(TARGET_INDEX, SOURCE INDEX) fixed binary (10,0);
declare
SEARCH STATUS character (1),
/* values for SEARCH_STATUS */
(SEARCHING initial (`?`),
SUCCEEDED initial (`!`),
FAILED initial (' X')) character (1),
(PERSON_INDEX, THIS NODE, ADJACENT NODE, BEST_NEARBY INDEX,
LAST NEARBY_INDEX) fixed binary (10,0),
NEARB\overline{Y}_NODE - dimension (1:300) fixed binary (10,0),
THIS EDGE fixed binary (4,0),
THIS NEIGHBOR pointer,
RELATIONSHIP fixed binary (4,0),
MINIMAL_DISTANCE float decimal (6);
/* begin execution of FIND RELATIONSHIP */
/* initialize PERSON-array for processing -
mark all nodes as not seen */
PERSON . REACHED STATUS = NOT SEEN;
/* mark source nōde as REACHED */
THIS NODE = SOURCE INDEX;
PERS\overline{ON (THIS NODE) - REACHED STATUS = REACHED;}
PERSON (THIS_NODE) . DISTANCE FROM_SOURCE = 0.0;
/* no NEARBY nodes exist yet */
LAST NEARBY INDEX = 0;
if T\overline{HIS NODE = TARGET INDEX then}
SEAR\overline{CH}_STATUS = SUC\overline{CEEDED;}
else
SEARCH_STATUS = SEARCHING;

```
/* Loop keeps processing closest-to-source, unREACHED node until target REACHED, or no more connected nodes. */

\section*{SEARCH FOR TARGET:}
do while - (SEARCH STATUS = SEARCHING);
/* Process all nodes adjacent to THIS NODE */
THIS_NEIGHBOR \(=\) PERSON (THIS_NODE) - NEIGHBOR_LIST_HEADER;
do whille (THIS_NEIGHBOR \({ }^{\sim}=\) nū1l());
call PROCESS_ADJACENT NODE (THIS NODE,
THIS NEIGHBOR \(\rightarrow\) NEIGHBOR INDEX,
THIS_NEIGHBOR \(\rightarrow\) NEIGHBOR_EDGE);
THIS_NEIGHBOR \(=\) THIS_NEIGHBOR \(\rightarrow\) - NEXT_NEIGHBOR;
end;
/* All nodes adjacent to THIS NODE are set. Now search for shortest-distance unREACHED (but NEARBY) node to process next. */
if LAST NEARBY INDEX \(=0\) then SEARCH_STATUS = FAILED;
else
do;
MINIMAL DISTANCE \(=1.0 \mathrm{e}+18\);
do PERSŌN INDEX = 1 to LAST NEARBY INDEX;
if PERSON (NEARBY NODE (PERSON_INDEX)) . DISTANCE_FROM_SOURCE
< MINIMAL_DİSTANCE then
do;
BEST NEARBY INDEX = PERSON INDEX;
MINIMAL DISTANCE =
PERSON (NEARBY_NODE (PERSON_INDEX)) . DISTANCE_FROM_SOURCE; end;
end; /* PERSON_INDEX loop */
/* establish new THIS_NODE */
THIS NODE \(=\) NEARBY NODE (BEST_NEARBY_INDEX);
/* change tHIS_NOD \(\bar{E}\) from being NEARB \(\bar{Y}\) to REACHED */
PERSON (THIS NODE) . REACHED_STATUS = REACHED;
/* remove THĪS NODE from NEARBY list */
NEARBY NODE (BEST_NEARBY_INDEX) = NEARBY_NODE (LAST_NEARBY_INDEX);
LAST_NEARBY INDEX \(=\) LAST NEARBY INDEX - \(\overline{1}\);
if THIS NODE \(=\) TARGET INDEX then
SEARC̄_STATUS = SUC̄CEEDED;
end; /*- determination of next node to process */
end SEARCH_FOR_TARGET;
/* Shortest path between PERSONs now established. Next task is
to translate path to English description of RELATIONSHIP. */
if SEARCH_STATUS = FAILED then
put skip list ( \({ }^{-}\)', PERSON (TARGET INDEX) . NAME, - is not related to PERSON (SOURCE_INDE \(\overline{\mathrm{X}}\) ) - NAME);
else /* success - parse path to find and display RELATIONSHIP */ do;
call RESOLVE PATH TO ENGLISH;
call COMPUTE_COMMŌN_GENES (SOURCE_INDEX, TARGET_INDEX); end;
/* End execution of FIND_RELATIONSHIP.

Utility procedures begin here. */
PROCESS_ADJACENT NODE: procedure (BASE NODE, NEXT_NODE, NEXT BASE EDGE);
/* NEXTT_NODE is adjacent to last-REAC̄HED node ( \(\bar{s}\) BASE_NODE).
if NEXT NODE already REACHED, do nothing.
If previously seen, check whether path thru BASE NODE is
shorter than current path to NEXT_NODE, and if so re-link
next to base.
If not previously seen, link next to base node. */
declare
(BASE NODE, NEXT NODE) fixed binary ( 10,0 ),
NEXT_BASE_EDGE fixed binary ( 4,0 );
declare
(WEIGHT_THIS_EDGE, DISTANCE THRU_BASE NODE)
float decīmal (6);
/* begin execution of PROCESS ADJACENT NODE */
if PERSON (NEXT_NODE) . REACHED_STATŪS \(\sim\) REACHED then
do;
if NEXT BASE EDGE \(=\) SPOUSE then WEIGHT_THĪS_EDGE = 1.8;
else
WEIGHT THIS EDGE = 1.0;
DISTANCE THRU BASE NODE = WEIGHT THIS EDGE +
PERSŌN (BAASE NODE) . DISTANCE FROM SOURCE;
if PERSON (NEXT_NNODE) - REACHED_STATUS \(=\) NOT_SEEN then do;
PERSON (NEXT NODE) . REACHED STATUS = NEARBY;
LAST_NEARBY_-̄NDEX = LAST_NEARBY_INDEX + 1;
NEAR \(\bar{B} Y\) NODE \({ }^{-}\)(LAST_NEARBY_INDEX) \(=\)NEXT_NODE;
call LĪNK_NEXT_NODE_TO_BASE_NODE;
end;
else /* REACHED_STATUS = NEARBY */
if DISTANCE THRU BASE NODE <
PERSON (NEXT NODE) . DISTANCE FROM SOURCE then
call LINK NEXT NŌDE TO BASE_NODE;
end; /* if REĀCHED_STATUS not = REACHED */
LINK NEXT_NODE_TO_BASE NODE: procedure;
/* link next to base by re-setting its predecessor index to point to base, note type of edge, and re-set distance as it is through base node. */
/* begin execution of LINK_NEXT_NODE_TO BASE_NODE */
PERSON (NEXT_NODE) . DISTANCE FROM_SOURCE = DISTANCE_THRU_BASE_NODE;
PERSON (NEXT NODE) . PATH PREDECES \(\bar{S} O R=\) BASE_NODE;
PERSON (NEXT NODE) . EDGE TO PREDECESSOR = NEXTBBASE EDGE;
end LINK_NEXT_NTODE_TO_BASE_ \(\bar{N} O D \bar{E}\);
end PROCESS_ADJACENT_NODE ;
/* End utility procedures under FIND_RELATIONSHIP.

Begin two major procedures: RESOLVE_PATH_TO_ENGLISH and COMPUTE_COMMON GENES */

RESOLVE PATH TO ENGLISH: procedure;
/* RESOLVE PATH TO ENGLISH condenses the shortest path to a
series of RELATIONSHIPs for which there are English
descriptions. */
/* Key persons are the ones in the RELATIONSHIP path which remain after the path is condensed. */
declare
/* values for sibling proximity */
(STEP initial ('S'),
HALF initial ( \({ }^{-} \mathrm{H}^{-}\)),
FULL initial ( \({ }^{-} \mathrm{F}^{-}\))) character (1);
declare
01 KEY_PERSON dimension (1:300),
05 PERSON INDEX fixed binary ( 10,0 ),
05 GENERATION GAP fixed binary \((10,0)\),
05 PROXIMITY - character (1),
05 RELATION TO NEXT fixed binary ( 4,0 ),
05 COUSIN RĀNK fixed binary \((10,0)\);
declare
/* these variables are used to condense the path */
(KEY RELATION, LATER KEY RELATION, PRIMARY RELATION,
NEXT PRIMARY RELATION) - fixed binary \((\overline{4}, 0)\),
GENERATION COÜNT fixed binary (10,0),
(KEY_INDEX, LATER_KEY_INDEX, PRIMARY_INDEX)
fixed binary \((10,0)\),
ANOTHER_ELEMENT POSSIBLE bit (1);
/* begin execution of RESOLVE PATH TO ENGLISH */

THIS NODE = TARGET INDEX;
/* Dīsplay path and initialize KEY PERSON array from path elements. */
TRAVERSE SHORTEST PATH:
do KEY_INDEX \(=1\) to 300 while (THIS_NODE \(\sim=\) SOURCE INDEX);
begin;
declare
EDGE TYPE dimension (1:3) character (9) static
initial (-parent of", "child of", "spouse of");
put skip list ( \({ }^{-}\)- \| PERSON (THIS_NODE) • NAME \| \(\|^{-}\)is ||
EDGE TYPE (PERSON (THIS NODE) . EDGE TO PREDECESSOR));
end;
KEY PERSON (KEY INDEX) . PERSON INDEX = THIS NODE;
KEY PERSON (KEY INDEX) . RELATION TO NEXT =
PERSON (THIS NODE) . EDGE TO P \(\overline{R E D E C E S S O R ; ~}\)
if PERSON (THIS NODE) - EDGE TO PREDECESSOR \(=\) SPOUSE then KEY_PERSON (KEY INDEX) . GENERATION_GAP \(=0\);
else
KEY PERSON (KEY INDEX) . GENERATION GAP = 1;
THIS NODE \(=\) PERSON \({ }^{-}(T H I S ~ N O D E) ~ \cdot ~ P A T H ~ P R E D E C E S S O R ; ~\)
end TRĀVERSE SHORTEST PATH;
put skip list( \({ }^{-}\)- || \(\overline{\text { PERSON (THIS_NODE) }}\). NAME);
```

    KEY PERSON (KEY INDEX) . PERSON_INDEX = THIS NODE;
    KEY - PERSON (KEY INDEX) . RELATIONN TO NEXT = NULL - RELATION;
    KEY-PERSON (KEY INDEX + 1) . RELATION-TO-NEXT = NULL_RELATION;
    /* Resolve CHILD}-PARENT and CHILD-SPOUSSE-PARENT relation
    to SIBLING relations. */
    FIND_SIBLINGS:
do KEY INDEX = 1 to 300
while (KEY PERSON (KEY INDEX) . RELATION_TO NEXT ~= NULL_RELATION);
if KEY_PERSO\overline{N} (KEY_INDE\overline{X}) . RELATION_TO_NEXT = CHILD then
do;
LATER KEY RELATION = KEY PERSON (KEY_INDEX + 1) . RELATION_TO_NEXT;
if LATER_KEY_RELATION = \overline{PARENT then}
/* found either full or half SIBLINGs */
do;
KEY PERSON (KEY INDEX) . RELATION TO NEXT = SIBLING;
if \overline{FULL SIblING}\mp@subsup{}{}{-}(\mathrm{ KEY PERSON (KEY I}
KEY_PERSON (KEY_INDEX + 2) . PERSON_INDEX)
then
KEY PERSON (KEY_INDEX) . PROXIMITY = FULL;
else
KEY PERSON (KEY_INDEX) . PROXIMITY = HALF;
call CONDENSE_KEY_\overline{PERSONS (KEY_INDEX, 1);}
end; /* processing of full/half SIBLINGs */
else
if (LATER_KEY RELATION = SPOUSE) \&
(KEY PERSON
then /\overline{*}}\mathrm{ found step-SIBLINGs */
do;
KEY_PERSON (KEY INDEX) . RELATION_TO_NEXT = SIBLING;
KEY PERSON (KEY_INDEX) . PROXIMITY - = STEP;
cal\overline{1} CONDENSE K\overline{EY PERSONS (KEY INDEX, 2);}
end; /* processing of step-S\overline{I}BLINGs */
end; /* if RELATION_TO_NEXT = CHILD */
end FIND_SIBLINGS;
/* Resolve CHILD-CHILD-... and PARENT-PARENT-... relations to
direct descendant or ancestor relations. */
FIND_ANCESTORS_OR_DESCENDANTS:
do KEY INDE \overline{X }=-1 to 300
whi\overline{le (KEY PERSON (KEY_INDEX) . RELATION_TO NEXT ~= NULL_RELATION);}
if (KEY_PER\overline{SON (KEY_INDEX) . RELATION_TO_}\overline{NEXT}=\mathrm{ CHILD) |}
(KEY_PERSON (KEY_INDEX) . RELATIONTO_NEXT = PARENT)
then
do;
do LATER KEY INDEX = KEY INDEX + 1 to 300
while (KEY PERSON (LATER_KEY_INDEX) . RELATION_TO_NEXT =
KEY_PERSON (KEY_INDEX) . RELATION_TO_NEXT);
end;
GENERATION_COUNT = LATER_KEY_INDEX - KEY_INDEX;
if GENERATION_COUNT > 1 then
do; /* compress generations */
KEY_PERSON (KEY_INDEX) . GENERATION_GAP = GENERATION_COUNT;
cal\overline{1}}\mathrm{ CONDENSE_KEYY_PERSONS (KEY_INDE合, GENERATION_COUNT - 1);
end;
end; /* if RELATION_TO NEXT = CHILD or PARENT */
end FIND_ANCESTORS_OR_DESCEENDA\overline{NTS;}

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```

    /* Resolve CHILD-SIBLING-PARENT to COUSIN,
                        CHILD-SIBLING to NEPHEW,
        SIBLING-PARENT to UNCLE. */
    FIND COUSINS NEPHEWS UNCLES:
do KEY INDEX = 1 to 300
while (KEY PERSON (KEY INDEX) . RELATION TO NEXT ~= NULL RELATION);
LATER KEY RELATION = KE\overline{Y} PERSON (KEY INDE\overline{X + I) . RELATION TO NEXT;}
if (KEY PERSON (KEY INDE\overline{X}) . RELATION
(LATER_KEY_RELAT\overline{I}ON = SIBLING)
then /* COUSIN or NEPHEW */
if KEY PERSON (KEY INDEX + 2) . RELATION TO NEXT = PARENT then
/* \overline{found COUSIN **}
do;
KEY PERSON (KEY INDEX) . RELATION TO NEXT = COUSIN;
KEY PERSON (KEY INDEX) . PROXIMIT\overline{Y }
KEY PERSON (KEY INDEX + 1) . PROXIMITY;
KEY PERSON (KEY INDE\overline{X}) . COUSIN RANK =
min (KEY PERSON (KEY INDEX) - . GENERATION GAP,
KEY_PERSON (KEY_INDEX + 2) . GENERATION_GAP);
KEY PERSON (KEY INDEX) - GENERATION GAP =
abs (KEY_PER\overline{SON (KEY_INDEX) - GENERATION_GAP -}
KEY_PERSON (KEY_INDEX + 2) . GENERATION_GAP);
call CONDENSE_KEY_PERSONS (KEY_INDEX, 2);
end;
else /* found NEPHEW */
do;
KEY PERSON (KEY INDEX) . RELATION TO NEXT = NEPHEW;
KEY PERSON (KEY INDEX) . PROXIMITY =-
KEY PERSON (KEY INDEX + 1) . PROXIMITY;
call CONDDENSE_KEY_PERRONS (KEY_INDEX, 1);
end;
else /* not COUSIN or NEPHEW */
if (KEY_PERSON (KEY INDEX) . RELATION_TO_NEXT = SIBLING) \&
(LATER KEY RELATION = PARENT)
then /* found UNCLE */
do;
KEY PERSON (KEY INDEX) . RELATION TO NEXT = UNCLE;
KEY - PERSON (KEY INDEX) . GENERATIONO G
KEY PERSON' (KEY INDEX + 1) . GENERATION GAP;
cal1 CONDENSE_KEY PERSONS (KEY_INDEX, 1);
end;
end FIND COUSINS NEPHEWS UNCLES;

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    /* Loop below will pick out valid adjacent strings of elements
        to be displayed. KEY_INDEX points to first element,
        LATER_KEY_INDEX to last element, and PRIMARY_INDEX to the
        element which determines the primary English word to be used.
        Associativity of adjacent elements in condensed table
        is based on English usage. */
    KEY INDEX = 1;
    put skip list (` Condensed path:`);
    CONSOLIDATE ADJACENT PERSONS:
do while (KEY PER\overline{SON (KEY INDEX) . RELATION TO NEXT ~= NULL RELATION);}
KEY_RELATIO\overline{N}}=\mathrm{ KEY_PERSON (KEY_INDEX) - RELATION_TO_NEXT;
LATER KEY INDEX = KEY_INDEX;
PRIMARYY INDEX = KEY INDEX;
if KEY PERSON (KEY INDEX + 1) . RELATION TO NEXT ~ = NULL RELATION then
do;- /* seek mūlti-element combination \overline{*/}
ANOTHER ELEMENT_POSSIBLE = TRUE;
if KEY RELATION = SPOUSE then
do;
LATER KEY INDEX = LATER KEY INDEX + 1;
PRIMA\overline{RY IN}DEX = LATER KEY - INDEX;
if (KEY PERSON (LATER KE\overline{Y INNDEX) . RELATION TO NEXT = SIBLING) |}
(KEY PERSON (LATER KEY - INDEX) . RELATION ' TO- NEXT = COUSIN)
then T* Nothing can fol\overline{low SPOUSE-SIBLING or SPOUSE-COUSIN */}
ANOTHER_ELEMENT_POSSIBLE = FALSE;
end;
/* PRIMARY INDEX is now correctly set. Next if-statement
determines if a following SPOUSE relation should be
appended to this combination or left for the next
combination. */
if ANOTHER ELEMENT POSSIBLE \&
(KEY_PERSON (PRTMARY_INDEX + 1) . RELATION_TO_NEXT = SPOUSE)
/* O\overline{nly a SPOUSE can-follow a Primary */}
then
do; /* check primary preceding and following SPOUSE. */
PRIMARY RELATION
KEY \overline{PERSON (PRIMARY INDEX) . RELATION TO NEXT;}
NEXT P\overline{RIMARY RELATION \#}
KEYY PERSON (PRIMARY INDEX + 2) . RELATION_TO_NEXT;
if (NE\overline{XT PRIMARY RELATITON = NEPHEW}
NEXT PRIMARY RELATION = COUSIN |
NEXT PRIMARY RELATION = NULL_RELATION)
| (PRIMMARY RELIATION = NEPHEW)
| ( ( PRIM\overline{ARY_RELATION = SIBLING |}
PRIMARY_RELATION = PARENT)
\& (NEXT_\overline{PRIMARY_RELATION ~= UNCLE ) )}
then /* append following SPOUSE with this combination. */
LATER KEY_INDEX = LATER_KEY_INDEX + 1;
end; /*- check primary preceding and following SPOUSE */
end; /* multi-element combination */
call DISPLAY_RELATION (KEY INDEX, LATER_KEY_INDEX, PRIMARY_INDEX);
KEY INDEX = LATER KEY INDEX + 1;
end CŌNSOLIDATE ADJA}\overline{CENT
put skip list (' ' | || PE\overline{RSON (KEY_PERSON (KEY_INDEX) . PERSON_INDEX) . NAME);}
/* End execution of RESOLVE_PATH_TO_ENGLISH.

```

Begin utility procedures for RESOLVE_PATH_TO_ENGLISH. */
FULL_SIBLING: procedure (INDEX1, INDEX2)
returns (bit(1));
/* Determines whether two PERSONs are full siblings, i.e., have the same two parents. */
declare
(INDEX1, INDEX2) fixed binary \((10,0)\);

\section*{return}
((PERSON (INDEX1) • RELATIVE IDENTIFIER (FATHER IDENT) ~ \(=\) NULL IDENT) \(\&\)
(PERSON (INDEX1) • RELATIVE IDENTIFIER (MOTHER IDENT) \(\sim=\) NULL IDENT) \&
(PERSON (INDEXI) • RELATIVE IDENTIFIER (FATHER IDENT) =
PERSON (INDEX2) • RELATIVE IDENTIFIER (FATHER IDENT) ) \&
(PERSON (INDEX1) • RELATIVE IDENTIFIER (MOTHER IDENT) =
PERSON (INDEX2) • RELATTVE IDENTIFIER (MOTHER_IDENT) ) ;
end FULL_SIBLING;
CONDENSE KEY PERSONS: procedure (AT_INDEX, GAP_SIZE);
/* CONDENSE KEY PERSONS condenses superfluous entries from the KEY PERSON array, starting at AT INDEX. */
declare
AT INDEX fixed binary \((10,0)\),
GAP SIZE fixed binary ( 10,0 );
declare
(RECEIVE INDEX, SEND INDEX) fixed binary (10,0);
/* begin execution of CONDENSE_KEY_PERSONS */
RECEIVE INDEX \(=A T\) INDEX +1 ;
SEND_INDEX = RECEIVE_INDEX + GAP SIZE;
KEY PERSON (RECEIVE INDEX) = KEY PERSON (SEND INDEX);
do \(\overline{\text { while (KEY PERSON }}\) (SEND_INDEX) - RELATION_TO_NEXT \(\sim=\) NULL_RELATION);
RECEIVE INDEX = RECEIVE \(\bar{I} N D E X+1\);
SEND_IN \(\overline{D E X}=\) RECEIVE_INDEX + GAP SIZE;
KEY_ \(\bar{P} E R S O N\) (RECEIVE_INDEX) \(=\) KEY_PERSON (SEND_INDEX);
end;
end CONDENSE KEY PERSONS;
/* End utility procedures.

Begin DISPLAY RELATION, which does major work of displaying under RESOLVE PATH_TO ENGLISH. */

DISPLAY RELATION: procedure (FIRST INDEX, LAST INDEX, PRIMARY INDEX);
/* DISPLAY RELATION takes 1, 2, or 3 adjacent elements in the condensed table and generates the English description of the relation between the first and last +1 elements. */
declare
(FIRST INDEX, LAST INDEX, PRIMARY_INDEX) fixed binary (10,0);
declare
DISPLAY_BUFFER character (80) varying,
INLAW bit (1),
THIS PROXIMITY character (1),
THIS GENDER character (1),
SUFFIX INDICATOR fixed binary ( 6,0 ),
(FIRST RELATION, LAST RELATION, PRIMARY RELATION)
fixed binary \((4,0)\),
(THIS_GENERATION_GAP, THIS_COUSIN RANK)
fixed biñary ( \(1 \overline{0}, 0\) );
```

/* begin execution of DISPLAY RELATION */
FIRST RELATION $=$ KEY PERSON (FIRST INDEX) - RELATION_TO NEXT;
LAST $\bar{R} E L A T I O N=K E Y$ PERSON (LAST $\overline{\text { INDEX) }}$ ) RELATION TO-NEXT;
PRIMĀRY RELATION = KEY PERSON (PRIMARY INDEX) - RELATION TO-NEXT;
/* set THIS PROXIMITY *T
if ((PRIMAR $\bar{Y}$ RELATION $=$ PARENT) $\&($ FIRST RELATION $=$ SPOUSE $)) \mid$
$(($ PRIMARY_RELATION $=$ CHILD) \& (LAST_RELATION = SPOUSE))
then
THIS PROXIMITY = STEP;
else
if PRIMARY RELATION = SIBLING
PRIMARY RELATION = UNCLE
PRIMARY RELATION = NEPHEW
PRIMARY-RELATION $=$ COUSIN
then
THIS_PROXIMITY = KEY PERSON (PRIMARY_INDEX) • PROXIMITY;
else
THIS PROXIMITY = FULL;
/* set THIS GENERATION GAP */
if PRIMARY RELATION = ̄̄ARENT
PRIMARY RELATION = CHILD
PRIMARY RELATION = UNCLE
PRIMARY RELATION = NEPHEW
PRIMARY_RELATION $=$ COUSIN
then
THIS_GENERATION GAP $=$ KEY_PERSON (PRIMARY INDEX) • GENERATION_GAP;
else
THIS_GENERATION_GAP $=0$;

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/* set INLAW */
INLAW = FALSE;
if (FIRST RELATION = SPOUSE) \&
(PRIMARTY RELATION = SIBLING
PRIMARY RELATION = CHILD
PRIMARY RELATION = NEPHEW |
PRIMARY_RELATION = COUSIN)
then
INLAW = TRUE;
if (LAST_RELATION = SPOUSE) \&
(PRIMĀRY RELATION = SIBLING |
PRIMARY - RELATION = PARENT
PRIMARY RELATION = UNCLE
PRIMARY_RELATION = COUSIN)
then
INLAW = TRUE;
/* set THIS COUSIN RANK */
if PRIMARY \overline{RELATIO}\overline{N}= COUSIN then
THIS_COUSIN_RANK = KEY_PERSON (PRIMARY_INDEX) . COUSIN_RANK;
else
THIS_COUSIN_RANK = 0;
/* parameters are set - now generate display. */
DISPLAY BUFFER =
- - TI PERSON (KEY_PERSON (FIRST_INDEX) . PERSON_INDEX) . NAME || is `; if PRIMARY RELATION = PARENT     PRIMARY RELATION = CHILD     PRIMARY RELATION = UNCLE     PRIMARY_RELATION = NEPHEW then     do; /* write generation-qualifier */     if THIS_GENERATION_GAP >= 3 then         do;         DISPLAY BUFFER = DISPLAY BUFFER || 'great`;
if THIS_GENERATION_GAP > 3 then
DISPLAY_BUFFER = DISPLAY BUFFER || -*- |
TRIM (THIS_GENERATION \overline{GAP - 2);}
DISPLAY_BUFFER = DISPLAY_BÜFFER || --';
end;
if THIS GENERATION GAP >= 2 then
DISPİAY BUFFER = DISPLAY_BUFFER || 'grand-`;         end; else         if (PRIMARY_RELATION = COUSIN) & (THIS_COUSIN_RANK > 1) then         do;         DISPLAY BUFFER = DISPLAY BUFFER || TRIM (THIS_COUSIN_RANK);         SUFFIX İNDICATOR = mod (THIS_COUSIN_RANK, 10);         if SUFFIX_INDICATOR > 3 then             SUFFIX_INDICATOR = 0;         DISPLAY_BÜFFER = DISPLAY BUFFER |             subs\overline{tr}(`}\mathrm{ (h st nd rd `', 3 * SUFFIX_INDICATOR + 1, 3);
end;

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if THIS PROXIMITY = STEP then
DISPĪAY_BUFFER = DISPLAY_BUFFER || 'step-';
else
if THIS PROXIMITY = HALF then
DISPİAY BUFFER = DISPLAY BUFFER || 'half-`; THIS_GENDER = PERSON (KEY_PERSON (FIRST_INDEX) . PERSON_INDEX) . GENDER; if PRIMARY RELATION = PARENT then     if THIS_GENDER = MALE then DISPLAY_BUFFER = DISPLAY BUFFER || -father`;
else DISPLAY BUFFER = DISPLAY_BUFFER | 'mother';
else if PRIMARY RELATION = CHILD then
if THIS GENDER = MALE then DISPLAY_BUFFER = DISPLAY_BUFFER || 'son`;     else - DISPLAY_BUFFER = DISPLAY_BUFFER | -daughter`;
else if PRIMARY RELATION = SPOUSE then
if THIS GENDER = MALE then DISPLAY BUFFER = DISPLAY BUFFER || 'husband';
else - DISPLAY BUFFER = DISPLAY BUFFER || 'wife`; else if PRIMARY RELATION = SIBLING then     if THIS_GENDER = MALE then DISPLAY BUFFER = DISPLAY BUFFER || 'brother`;
else DISPLAY_BUFFER = DISPLAY_BUFFER || 'sister`; else if PRIMARY RELATION = UNCLE then     if THIS GENDER = MALE then DISPLAY_BUFFER = DISPLAY_BUFFER || 'uncle`;
else - DISPLAY_BUFFER = DISPLAY_BUFFER || 'aunt`; else if PRIMARY RELATION = NEPHEW then     if THIS GENDER = MALE then DISPLAY_BUFFER = DISPLAY_BUFFER || 'nephew';     else - DISPLAY_BUFFER = DISPLAY_BUFFER || 'niece'; else if PRIMARY RELATION = COUSIN then                                     DISPLAY_BUFFER = DISPLAY_BUFFER || 'cousin'; else                             DISPLAY_BUFFER = DISPLAY_BUFFER || 'nul1'; if INLAW then     DISPLAY BUFFER = DISPLAY_BUFFER || '-in-law'; if (PRIMARY RELATION = COUSIN) & (THIS GENERATION_GAP > 0) then     if THIS GENERATION GAP > 1 then         DISPIAY BUFFER \equiv DISPLAY BUFFER || - ||             TRIM (THIS_GENERATION_GAP) || - times removed`;
else
DISPLAY BUFFER = DISPLAY_BUFFER || - once removed`; DISPLAY BUFFER = DISPLAY BUFFER | - of`;
put ski\overline{p}}\mathrm{ list (DISPLAY_BŪFFER);

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/* Begin utility procedure for DISPLAY RELATION */
TRIM: procedure (NUMERIC VALUE) returns (character (20) varying);
/* Returns character representation of numeric values
with no leading or trailing spaces. */
declare
NUMERIC VALUE fixed binary (10,0);
declare
STRING REPRESENTATION character (20),
(START_LOCATION, STOP_LOCATION)
fixed binary (10,0);
/* Begin execution of TRIM */
STRING REPRESENTATION = NUMERIC VALUE;
do STA\overline{RT}}\mathrm{ LOCATION = 1 to 20
while-}\mathrm{ (substr (STRING_REPRESENTATION, START LOCATION, 1) = - `);
end;
do STOP LOCATION = 20 to 1 by -1
whil\overline{e}}\mathrm{ (substr (STRING REPRESENTATION, STOP LOCATION, 1) = - ');
end;
return (substr (STRING REPRESENTATION, START LOCATION,
STOP_LŌCATION - START_LOCATIÖN + 1));
end TRIM;
end DISPLAY_RELATION;
end RESOLVE PATH TO ENGLISH;
/* COMPUTE COMMON GENES is second major procedure (after
RESOLVE PATH TO}\mathrm{ ENGLISH) under FIND RELATIONSHIP. */
COMPUTE COMMON GENES: procedure (INDEX1, INDEX2);
/* COMPUTE COMMON GENES assumes that each ancestor contributes
half of the genetic material to a PERSON. It finds common
ancestors between two PERSONs and computes the expected
value of the PROPORTION of common material. */
declare
(INDEX1, INDEX2) fixed binary (10,0);
declare
COMMON PROPORTION float decimal (6);
/* begin execution of COMPUTE COMMON GENES */
/* First zero out all ancestors to allow adding. This is necessary
because there might be two paths to an ancestor. */
call ZERO PROPORTION (INDEXI);
/* now mark with shared PROPORTION */
call MARK PROPORTION (PERSON (INDEXI) . IDENTIFIER, 1.0, INDEX1);
COMMON PROPORTION = 0.0;
call CHECK COMMON PROPORTION (COMMON_PROPORTION,
PERSON (INDEX1). IDENTIFIER, 1.0, 0.0, INDEX2);
put skip list (' Proportion of common genetic material = ');
put edit (COMMON_PROPORTION) (e(13,5,6));
/* End execution of COMPUTE_COMMON_GENES.

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    Begin utility procedures. */
    ZERO_PROPORTION: procedure (ZERO_INDEX) recursive;
/* ZERO_PROPORTION recursively seeks out all ancestors and
zeros them out. */
declare
ZERO_INDEX fixed binary (10,0),
THIS_NEIGHBOR pointer;
/* begin execution of ZERO PROPORTION */
PERSON (ZERO INDEX) . DESCENDANT GENES = 0.0;
THIS NEIGHBO\overline{R}= PERSON (ZERO_INDEX) . NEIGHBOR LIST HEADER;
do while (THIS NEIGHBOR ~= null());
if THIS NEIGHBOR }->\mathrm{ NEIGHBOR EDGE = PARENT then
call ZERO_PROPORTION (THIS_NEIGHBOR ->> NEIGHBOR_INDEX);
THIS_NEIGHBOR = THIS_NEIGHBOR -> NEXT_NEIGHBOR;
end;
end ZERO_PROPORTION;
MARK PROPORTION: procedure (MARKER, PROPORTION, MARKED_INDEX) recursive;
/*- MARK PROPORTION recursively seeks out all ancestors and
marks
genetic material. This PROPORTION is diluted by one-half
for each generation. */
declare
MARKER picture '999`,
PROPORTION float decimal (6),
MARKED INDEX fixed binary (10,0),
THIS_NEIGHBOR pointer;
/* begin execution of MARK PROPORTION */
PERSON (MARKED_INDEX) . DESCENDANT_IDENTIFIER = MARKER;
PERSON (MARKED INDEX) . DESCENDANT GENES =
PERSON (MAR\overline{RED INDEX) . DESCENDĀNT GENES + PROPORTION;}
THIS NEIGHBOR = PERSON (MARKED INDEX) - . NEIGHBOR LIST HEADER;
do while (THIS NEIGHBOR ~= nul\overline{1}());
if THIS NEIGHBOR -> NEIGHBOR EDGE = PARENT then
call MARK_PROPORTION (MARKER, PROPORTION / 2.0,
THIS_NEIGHBOR -> NEIGHBOR_INDEX);
THIS_NEIGHBOR = THIS_NEIG
end;
end MARK_PROPORTION;

```
```

    CHECK COMMON PROPORTION: procedure
                (COMMON PROPORTION, MATCH IDENTIFIER, PROPORTION,
                ALREAD\overline{Y}}\mathrm{ COUNTED, CHECK IN}DEX) recursive;
        /* CHECK COMMON PROPORTION searches all the ancestors of
            CHECK INDEX to see if any have been marked, and if so
            adds the appropriate amount to COMMON PROPORTION. */
        declare
        COMMON PROPORTION float decimal (6),
        MATCH_IDENTIFIER picture '999`,
        PROPORTION float decimal (6),
        ALREADY COUNTED float decimal (6),
        CHECK INDDEX fixed binary ( }10,0)\mathrm{ ,
        THIS NEIGHBOR pointer,
        THIS_CONTRIBUTION float decimal (6);
    /* begin execution of CHECK_COMMON PROPORTION */
        if PERSON (CHECK INDEX) . DESCENDANT IDENTIFIER = MATCH IDENTIFIER then
            /* Increment \overline{COMMON PROPORTION by the contribution of}
                this common ancestor, but discount for the contribution
                of less remote ancestors already counted. */
        do;
        THIS_CONTRIBUTION = PERSON (CHECK_INDEX) . DESCENDANT_GENES
                            * PROPORTION;
        COMMON PROPORTION = COMMON PROPORTION
            + T\overline{HIS_CONTRIBUTION - ALRREADY COUNTED;}
        end;
        else
            THIS CONTRIBUTION = 0.0;
        THIS NEIGHBOR = PERSON (CHECK INDEX) - NEIGHBOR LIST HEADER;
        do wh}ile (THIS NEIGHBOR ~= nu\overline{l}1())
            if THIS NEIGHBOR -> NEIGHBOR EDGE = PARENT then
                Ca11 CHECK COMMON PROPORTITON (COMMON PROPORTION,
                    MATC\overline{H}}\mathrm{ IDENTIFIER, PROPORTION / 2.0,
                THIS_\overline{CONTRIBUTION / 4.0,}
                THIS NEIGHBOR -> NEIGHBOR INDEX);
            THIS_NEIGHBOR = THIS_NEIGHBOR -> NEXT_NEIGHBOR;
        end;
    end CHECK_COMMON_PROPORTION;
    end COMPUTE COMMON GENES;
end FIND RELATIONSHIP;
end RELATE;

```
4. TITLE AND SUBTITLE

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5. AUTHOR(S)

John V. Cugini
6. PERFORMING ORGANIZATION (If joint or other than NBS, see instructions) 7. Contract/Grant No.

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10. SUPPLEMENTARY NOTES

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\(\square\) Document describes a computer program; SF-185, FIPS Software Summary, is attached.
11. ABSTRACT (A 200-word or less factual summary of most significant information. If document includes a significant bibliography or literature survey. mention it here)

Programming languages have been and will continue to be an important instrument for the automation of a wide variety of functions within industry and the Federal Government. Other instruments, such as program generators, application packages, query languages, and the like, are also available and their use is preferable in some circumstances.

Given that conventional programming is the appropriate technique for a particular application, the choice among the various languages becomes an important issue. There are a great number of selection criteria, not all of which depend directly on the language itself. Broadly speaking, the criteria are based on 1) the language and its implementation, 2) the application to be progranmed, and 3) the user's existing facilities and software.

This study presents a survey of selection factors for the major general-purpose languages: Ada, BASIC, C, COBOL, FORTRAN, Pascal, and PL/I. The factors covered include not only the logical operations within each language, but also the advantages and disadvantages stemming from the current computing environment, e.g., software packages, microcomputers, and standards. The criteria associated with the application and the user's facilities are explained. Finally, there is a set of program examples to illustrate the features of the various languages.

This volume includes the program examples. Volume 1 contains the discussion of language selection criteria.
12. KEY WORDS (Six to twelve entries; alphabetical order; capitalize only proper names; and separate key words by semicolons) Ada; alternatives to programming; BASIC; C; COBOL; FORTRAN; Pascal; PL/I; programming language features; programming languages; selection of programming language.
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