

DEPARTMENT OF THE INTERIOR

U.S. GEOLOGICAL SURVEY

Technical Manual and Program Listing for REFORM:  
A Rock-Sample Database Program in FORTRAN-77

by

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## INTRODUCTION

This manual documents REFORM version 2.1, and describes system requirements, files used or created by REFORM, the REFORM program and subroutines, and how to modify the REFORM software. The source code for REFORM is included at the end of this manual. Instructions for using REFORM, program specifications, and examples are contained in the companion report, User's Manual for REFORM: A Rock-Sample Database Program in FORTRAN-77, USGS Open File Report 87-141-A. It will be referred to hence as the User's Manual. The present manual assumes you are familiar with the material in the User's Manual.

REFORM is a program that allows you to create and use a database for certain kinds of rock sample data. It has sections for information on sample location, formation, rock type, characteristics (e.g., modes), major and trace element chemistry, and isotopes. It is especially suited to igneous rocks.

REFORM generates printouts of sample data, reformats chemical data for GNAIP and TOMGNAIP (normative calculation programs), and generates tables of selected data fields.

For portability of the database, data is stored in ASCII text files. For portability of the REFORM system, the software is written in FORTRAN-77. REFORM is currently installed on the U.S. Geological Survey Information Systems Division Digital Equipment Corporation VAX/VMS in Menlo Park. REFORM's hierarchical structure and input and output routines can be modified with relatively simple programming.

Data is stored in one or more text files. New data can be added with REFORM's interactive data entry routine, or via an editor or punched cards. Sorting of database records is done with one of REFORM's routines and/or with a text editor. Corrections to data are made with a text editor.

An initial version of REFORM was written in PL1 by Ming Ko using MRDS on a Honeywell Multics computer. Charlotte Allen assisted with program testing. The FORTRAN version described here is a new stand-alone revision with substantial extensions. It was developed in conjunction with the Pacific-to-Arizona-Crustal-Experiment (PACE) by Bob Simpson and Todd Fitzgibbon.

## PROGRAM DESCRIPTION

REFORM offers five options that are displayed on the main menu. They are:

- A=Make a new card file
- B=Make a printout (of a card file)
- C=Make a table
- D=Sort a card file
- E=Reformat chemistry data for TOMGNAP

If option E is chosen a second menu is displayed showing the TOMGNAP options:

- 1=Reformat data from the ANALYSIS 1 block
- 2=Reformat data from the ANALYSIS 2 block
- 3=Reformat data from the ANALYSIS 3 and 1 blocks

REFORM.FOR is the main loop. Option A is performed by the subroutine MAKECARDS.FOR, option B by OUTSAMP.FOR, option C by OUTTABLE.FOR, option D by SHUFFLE.FOR, option E1 by OUT1.FOR, E2 by OUT2.FOR, and E3 by OUTINAA.FOR. For options B, C, and E an existing card file must be read into memory, one sample at a time, prior to being output to a file, again one sample at a time. Nine "read" subroutines perform the input task.

See the User's Manual for details of what each option does.

## HARDWARE AND SOFTWARE REQUIREMENTS

REFORM version 2.1 currently runs on the VAX/VMS system under VMS version 4.2. (See "Porting REFORM to Other Systems" in this manual if you will not be running REFORM on a VAX/VMS system). REFORM was compiled using VAX-11 FORTRAN version 4.2. The Digital Equipment Corporation manuals for VAX FORTRAN are:

VAX-11 FORTRAN Language Reference Manual

VAX-11 FORTRAN User's Guide

REFORM has three options that format chemistry data from card files for TOMGNAP, a norm calculation and plotting program. GNAP stands for Graphic Normative Analysis Program. TOMGNAP is an extended version of GNAP by Tom Wright and Sam Priebe of the USGS. The reference manual for GNAP is:

Stuckless, J.S., and VanTrump, Jr., G., 1979, A revised version of Graphic Normative Analysis Program (GNAP) with examples of petrologic problem solving: U.S. Geological Survey Open-File Report 79-1237, 51 p.

Version 1.0 of REFORM was the Multics version. Version 2.0 was an early VAX/VMS version that included several preliminary routines that have, in the current version, been either combined or discarded.

## FILES

All files created and used by REFORM are sequential formatted (ASCII) files. That is, ACCESS=SEQUENTIAL, and FORM=FORMATTED. Each line is one record.

VAX-11 FORTRAN provides a CARRIAGE CONTROL parameter for OPEN statements. CARRIAGE CONTROL=FORTRAN specifies the normal FORTRAN interpretation of the first character of each line as a carriage control character. CARRIAGE CONTROL=LIST causes the first character of each line not to be interpreted as a carriage control character, and specifies single spacing between lines. REFORM uses both types as noted below.

## DATABASE FILE FORMAT

Data are stored in one or more ASCII text files, referred to as card files. CARRIAGE CONTROL=LIST. STATUS=NEW when opened by MAKECARDS and SHUFFLE, and OLD when opened by REFORM and SHUFFLE. Each sample is stored as a set of records (cards) which are subdivided into one or more data blocks (e.g., LOCATION) containing information of a particular type. Not all kinds of data blocks need be present for any one sample. See page 4 for a printout of the database format, with underlines showing field sizes. Only those fields for which information is available need be entered. Block names (e.g., LOCATION) and field names must be spelled exactly as shown and be positioned in the exact column shown.

Note that this database has a strict column-oriented format, a characteristic inherited from the original Multics program in which input was in the form of punched cards (and thus the term "card files").

For a given sample entry, at most one copy of each block may be present. That is, only one CHARACTERISTICS block (therefore only one mode per sample), only one CHEMISTRY block (therefore only one of each of the three types of analyses provided for), etc. Each block is terminated by COMMENT=\$\$\$ starting in column 11, and the ANALYSIS 1,2, and 3 sections of the CHEMISTRY block are terminated by \$\$\$ starting in column 19.

Each field, with the exception of the COMMENT, SOURCE, and TEXTURE fields, has an alphanumeric key that is used to specify output items in REFORM's table routine. In general, numeric data have numeric keys and character data have alphabetic keys. Some numeric fields, such as CPS, that have variable formats or multiple entries, have alphabetic keys. See the User's Manual for a list of these keys.

Numeric fields may contain the less-than sign (<). It will appear on printouts (option B) of the data, but when the data is output for tables (option C) or for TOMGNAP files (option E) the number without the less-than sign will be used.

Refer to the User's Manual for an example of a card file, and for a glossary of terminology and abbreviations.

# FORMAT SHOWING FIELD SIZES

SAMPLE NO=\_\_\_\_\_

LOCATION:

MOUNTAIN RANGE=\_\_\_\_\_

LATITUDE=\_\_\_\_\_ LONGITUDE=\_\_\_\_\_

QUADRANGLE=\_\_\_\_\_

TRS=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

SOURCE=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\*\*\*

FORMATION:

FORMATION SYMBOL=\_\_\_\_\_

FORMATION NAME=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\*\*\*

ROCK TYPE:

ROCK TYPE=\_\_\_\_\_

GRAIN SIZE=\_\_\_\_\_

TEXTURE=\_\_\_\_\_

TEXTURE=\_\_\_\_\_

TEXTURE=\_\_\_\_\_

TEXTURE=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\*\*\*

CHARACTERISTICS:

CPS=\_\_\_\_\_ SB=\_\_\_\_\_ CI=\_\_\_\_\_ MAG=\_\_\_\_\_

SN=\_\_\_\_\_ SQ=\_\_\_\_\_ SP=\_\_\_\_\_ SA=\_\_\_\_\_ SM=\_\_\_\_\_

TN=\_\_\_\_\_ TQPA=\_\_\_\_\_ TB=\_\_\_\_\_ TH=\_\_\_\_\_ TM=\_\_\_\_\_ TSP=\_\_\_\_\_ TMG=\_\_\_\_\_

%Q1=\_\_\_\_\_ %P1=\_\_\_\_\_ %A1=\_\_\_\_\_ %Q2=\_\_\_\_\_ %P2=\_\_\_\_\_ %A2=\_\_\_\_\_ %M2=\_\_\_\_\_

OTHER MINERALS=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_

COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\$\$\$

CHEMISTRY:

ANALYSIS 1 BY: \_\_\_\_\_  
SI02=\_\_\_\_\_ TI02=\_\_\_\_\_ AL203=\_\_\_\_\_ FE203=\_\_\_\_\_ FEO=\_\_\_\_\_  
MNO=\_\_\_\_\_ MGO=\_\_\_\_\_ CAO=\_\_\_\_\_ NA2O=\_\_\_\_\_ K2O=\_\_\_\_\_  
P2O5=\_\_\_\_\_ H2O+=\_\_\_\_\_ H2O-=\_\_\_\_\_ CO2=\_\_\_\_\_ TOTAL=\_\_\_\_\_  
Y=\_\_\_\_\_ SR=\_\_\_\_\_ ZR=\_\_\_\_\_ U=\_\_\_\_\_ RB=\_\_\_\_\_  
TH=\_\_\_\_\_ PB=\_\_\_\_\_ GA=\_\_\_\_\_ ZN=\_\_\_\_\_ CU=\_\_\_\_\_  
NI=\_\_\_\_\_ CR=\_\_\_\_\_ V=\_\_\_\_\_ BA=\_\_\_\_\_ NB=\_\_\_\_\_  
\$\$\$

ANALYSIS 2 BY: \_\_\_\_\_  
SI02=\_\_\_\_\_ TI02=\_\_\_\_\_ AL203=\_\_\_\_\_ FE203=\_\_\_\_\_ MNO=\_\_\_\_\_  
MGO=\_\_\_\_\_ CAO=\_\_\_\_\_ NA2O=\_\_\_\_\_ K2O=\_\_\_\_\_ P2O5=\_\_\_\_\_  
Y=\_\_\_\_\_ SR=\_\_\_\_\_ ZR=\_\_\_\_\_ RB=\_\_\_\_\_ BA=\_\_\_\_\_  
NB=\_\_\_\_\_ FEO=\_\_\_\_\_  
\$\$\$

ANALYSIS 3 BY: \_\_\_\_\_  
FE=\_\_\_\_\_ K=\_\_\_\_\_ NA=\_\_\_\_\_ BA=\_\_\_\_\_  
CO=\_\_\_\_\_ CR=\_\_\_\_\_ CS=\_\_\_\_\_ HF=\_\_\_\_\_  
MN=\_\_\_\_\_ RB=\_\_\_\_\_ SB=\_\_\_\_\_ SR=\_\_\_\_\_  
TA=\_\_\_\_\_ TH=\_\_\_\_\_ U=\_\_\_\_\_ ZR=\_\_\_\_\_  
SC=\_\_\_\_\_ LA=\_\_\_\_\_ CE=\_\_\_\_\_ ND=\_\_\_\_\_  
SM=\_\_\_\_\_ EU=\_\_\_\_\_ GD=\_\_\_\_\_ TB=\_\_\_\_\_  
DY=\_\_\_\_\_ TM=\_\_\_\_\_ YB=\_\_\_\_\_ LU=\_\_\_\_\_  
HO=\_\_\_\_\_  
\$\$\$

RATIOS: BA/RB=\_\_\_\_\_ BA/SR=\_\_\_\_\_ RB/SR=\_\_\_\_\_  
RATIOS: U/TH=\_\_\_\_\_ K/RB=\_\_\_\_\_ K/BA=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\$\$\$

ISOTOPES:

ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
ANALYSIS=\_\_\_\_\_ ANALYST=\_\_\_\_\_ MINERAL=\_\_\_\_\_  
AGE=\_\_\_\_\_ RATIO=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\_\_\_\_\_  
COMMENT=\$\$\$

### TABLE FILE FORMAT

Tables have lines up to 126 characters long with data for one sample on one line. Each line contains user-specified fields aligned in columns, separated by a space.

CARRIAGE CONTROL=LIST. STATUS=NEW when opened by REFORM, and OLD when opened by SHUFFLE.

### PRINTOUT FILE FORMAT

Printout files contain lines with up to 126 characters per line. They put information for one sample on one or two pages, formatted on the page for easier reading.

CARRIAGE CONTROL=FORTRAN. STATUS=NEW, opened by REFORM.

### TOMGNAP FILE FORMAT

TOMGNAP files have 80-character lines. Chemistry data for one sample is contained on up to three lines, called analysis cards in the TOMGNAP manual. The format for TOMGNAP analysis cards is:

column 1		73	80
NRMX	...		Y

where X is the plotting symbol for the sample  
... is the chemistry data in percent with implied decimal points (as specified in the TOMGNAP command file. See the User's Manual and the GMAP manual).  
Y is the sample identification, up to eight characters starting in column 73.

CARRIAGE CONTROL=LIST, STATUS=NEW, called by REFORM.



## ERROR HANDLING

There are three classes of errors detected by REFORM. The first are caused by unacceptable user input and are detected during an interactive terminal session. The program's action is to display an error message and prompt the user again for input. The second are caused by card file format errors and are detected during reading of the card file or when converting strings to real numbers. The program displays an error message and tries to continue (which may cause more error messages for subsequent lines). The third class are fatal errors. The program displays an error message and stops.

A brief table of error messages follows which includes the class of error as described above (1, 2 or 3). The User's Manual contains a section on error handling and a more detailed description of error messages and the conditions for which they occur. The last error listed in the table below (error...inout variable...) is not described in the User's Manual because it results from incorrect use of the subroutine GETFILE and would be detected during program testing.

TABLE OF ERROR MESSAGES

ERROR CLASS	GENERATED WHEN RUNNING OPTION	MESSAGE (ABBREVIATED)	SUBROUTINE GENERATING MESSAGE
1	C (Tables)	invalid choice ...	GETENTRY
1	"	invalid choice (too long) ...	"
1	"	you've used XX columns ...	"
	E (TOMGNAP)		
2	All of	error trying to convert string...	RLNU
2	above	line reads ... error trying to read in line ...	BADCARD & CARDERROR
	B (Printout)		
2	All of	unrecognized line in XXXX block..	READXXXX & BADCARD
	above		
2	"	unknown datatype at line ...	REFORM
2	"	XX block not properly ended ...	READXXX
2	"	too many analysis lines ...	READISOT
2	"	" comment "	NEWCMT
2	"	" source "	NEWSOURC
2	"	" texture "	READROCK
1	A (Makecards)	answer too long ...	ASK
3	D (Shuffle)	error...array not dimensioned large enough ...	SHUFFLE
1	All of	error in opening ...	GETFILE
1	above	try again ...	REFORM
3	"	error...inout variable in sub getfile not recognized ...	GETFILE

# LIST OF SUBROUTINES

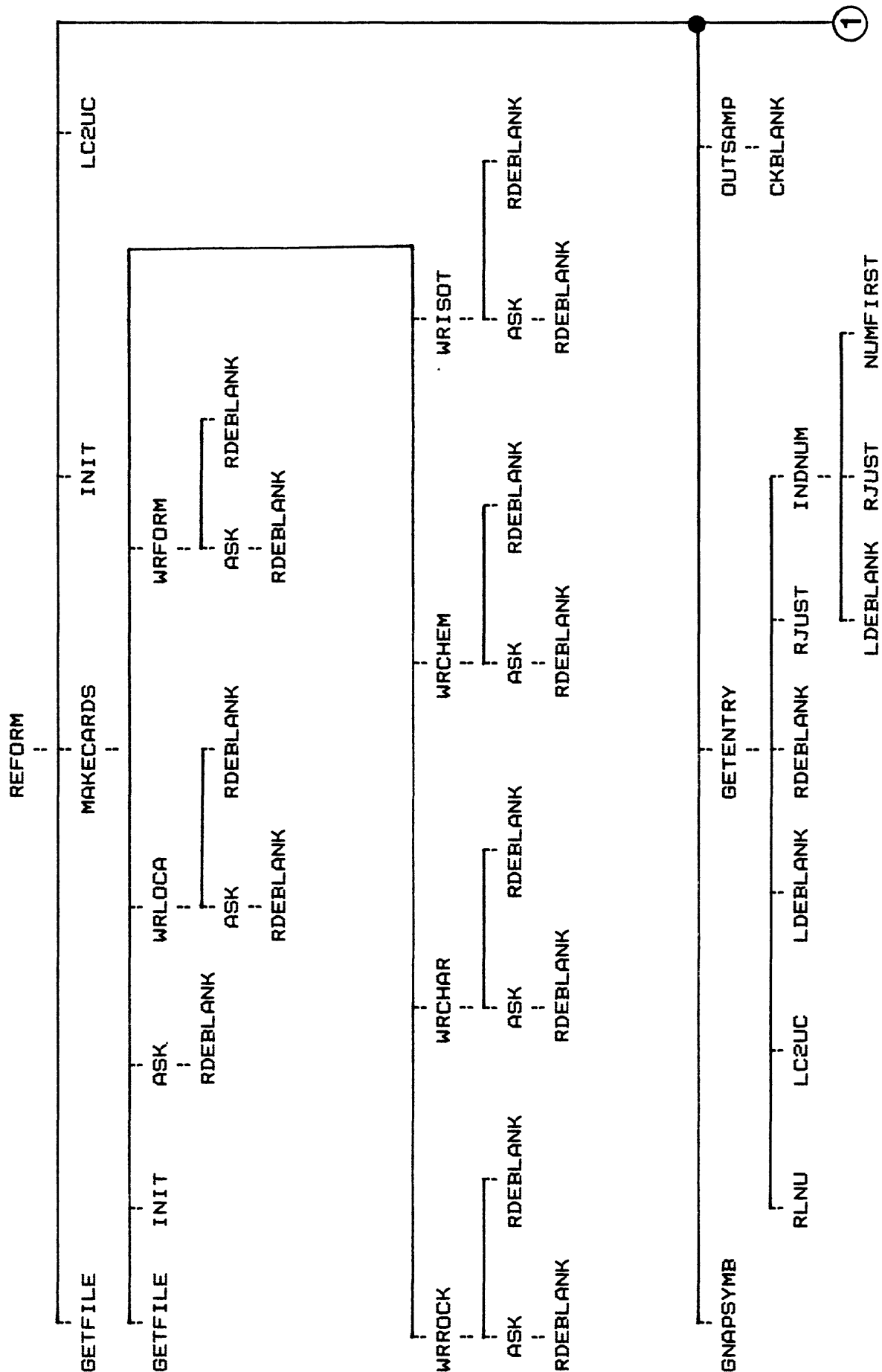
SUBROUTINE		SIZE OF SOURCE CODE FILE IN BYTES	FUNCTION
REFORM.FOR	X	4454	Main loop, displays menus.
VAR.INC	X	2175	Variable include module.
COMMON.INC	X	1329	Common include module.
BLDATA.FOR		4129	Block data.
ASK.FOR	X	944	Prompts user for data fields.
BADCARD.FOR		306	Error message, displays line with the error and the sample number.
BREAKSN.FOR	X	822	Breaks Needles-Project format sample number into components.
CARDERROR.FOR		275	Error message, displays line no.
CHRFIRST.FOR		421	Function, returns position of first alphabetic character in string.
CKBLANK.FOR		368	Counts how many entries in a character array are not blank.
FILLARRY.FOR	X	2200	Assigns data strings to character array or converts numeric strings to reals and assigns to real array.
GETENTRY.FOR	X	2567	Prompts user for fields to include in table.
GETFILE.FOR	X	1886	Opens files.
GNAPSYMB.FOR	X	1535	Prompts for plotting symbol.
INDNUM.FOR	X	1431	Converts alphabetic and numeric keys for a table to integer array.
INIT.FOR		1694	Initializes variables prior to reading each sample's data from card file.
LC2UC.FOR		409	Converts lower case letters to upper case.
LDEBLANK.FOR		530	Deletes blanks at the left end of a string.
MAKECARDS.FOR	X	9894*	Main loop of interactive data entry routine.
NEWCMT.FOR		428	Adds comment to comment array.
NEWSOURC.FOR		459	Adds source (bibliographic reference) to source array.
NUMFIRST.FOR		410	Function, finds position of first number in a string.
OUT1.FOR	X	2376	Outputs 2 cards of data from ANALYSIS 1 formatted for TOMGNAP norm program.
OUT2.FOR	X	1943	As above, but from ANALYSIS 2.
OUTINAA.FOR	X	3283	Outputs 3 cards of data from ANALYSES 1 and 3 for TOMGNAP.
OUTSAMP.FOR	X	5324	Produces printout file of sample data.

OUTTABLE.FOR	X	3096	Produces a table of selected fields of the database.
RDEBLANK.FOR		528	Deletes blanks at the right end of a string.
READCHAR.FOR	X	1728	-
READCHEM.FOR		2026	These routines read cards from
READFORM.FOR		1141	the blocks of a cardfile and
READISOT.FOR		1614	assign the data to string
READLOCA.FOR		1464	variables. E.g., READCHAR
READNIAA.FOR		1770	reads data from the
READROCK.FOR		1473	CHARACTERISTICS block.
READSHAW.FOR		1595	
READXRF.FOR		1388	-
REMCHAR.FOR		511	Removes character from string and returns length of new string.
RJUST.FOR		485	Right justifies string in its length.
RLNU.FOR		892	Function, reads character string containing real number.
SHUFFLE.FOR	X	941	Sorts a card file according to index fields in a table.
WRLOCA		*	-
WRFORM			These routines prompt user
WRROCK			for data, delete trailing
WRCHAR			blanks, and write it to a
WRCHEM			new card file.
WRISOT			-
TOTAL		95535 bytes	

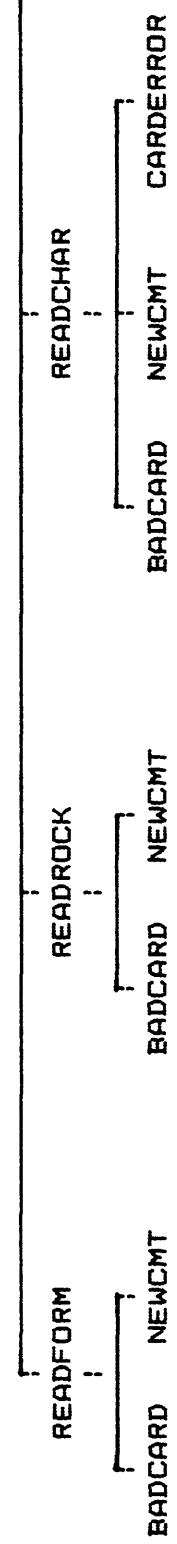
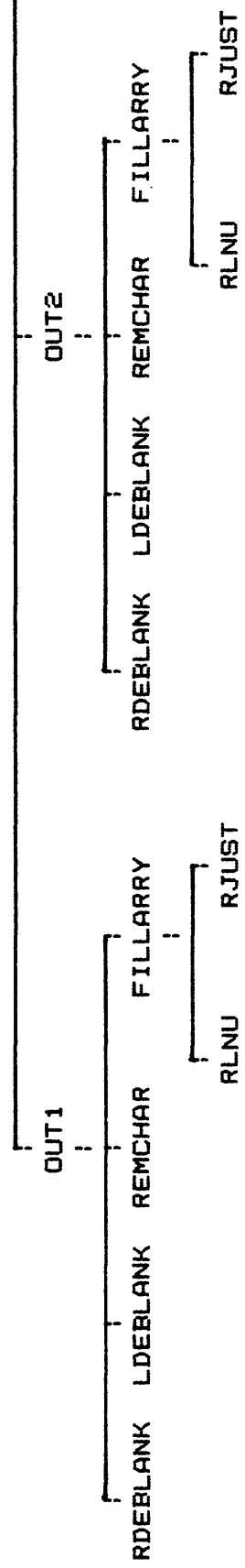
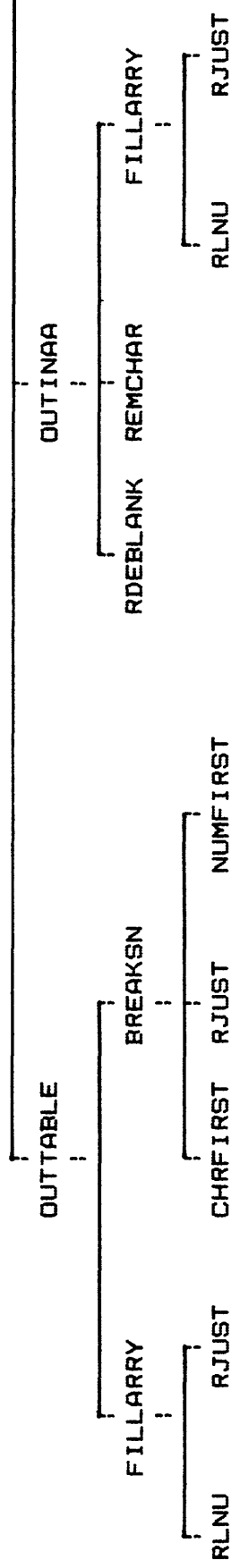
X indicates an extended description is given below.

\* the size of MAKECARDS includes the write routines (WRLOCA, etc.) They are described with MAKECARDS below.

# SUBROUTINE HIERARCHY

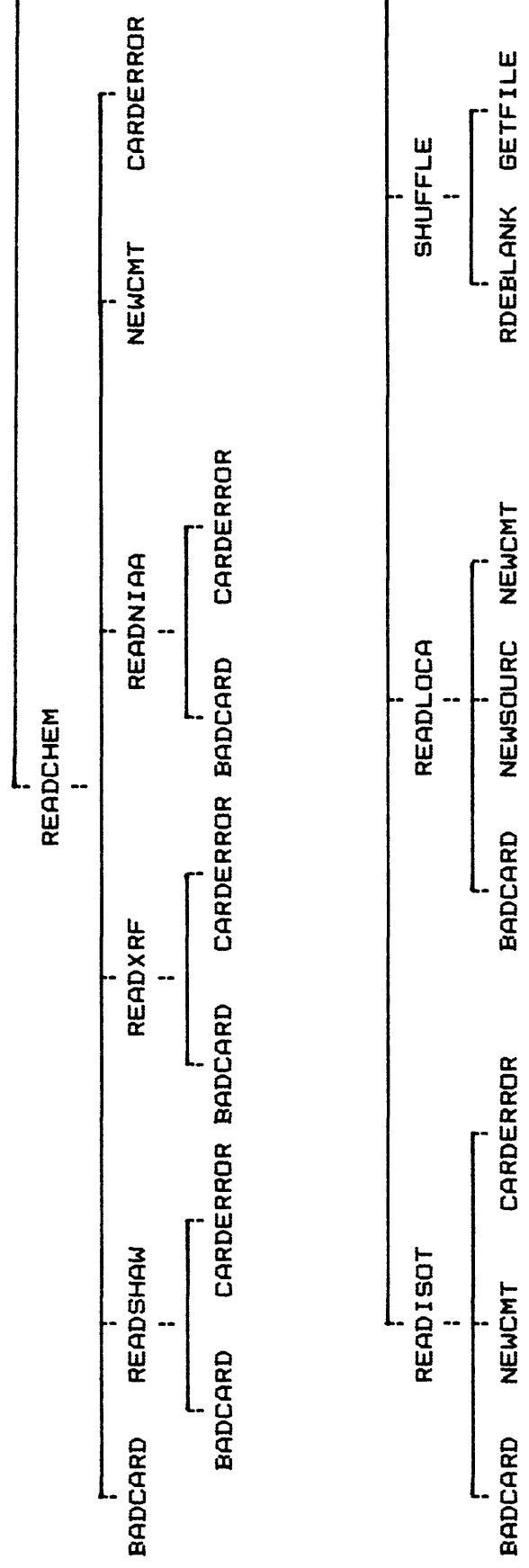


1



2

2



LIST OF SUBROUTINES AND THE SUBROUTINES THAT CALL THEM

SUBROUTINE	CALLED BY
VAR.INC, COMMON.INC	REFORM, FILLARRY, GETENTRY, GNAPSYMB, INDNUM, INIT, MAKECARDS, OUT1, OUT2, OUTINAA, OUTSAMP, OUTTABLE, READCHAR, READCHEM, READFORM, READISOT, READLOCA, READNIAA, READROCK, READSHAW, READXRF
ASK	MAKECARDS, WRLOCA, WRFORM, WRROCK, WRCHAR, WRCHEM, WRISOT
BADCARD	READCHAR, READCHEM, READFORM, READISOT, READLOCA, READNIAA, READROCK, READSHAW, READXRF
BREAKSN	OUTTABLE
CARDERROR	READCHAR, READSHAW, READXRF, READNIAA, READCHEM, READISOT
CHRFIRST	BREAKSN
CKBLANK	OUTSAMP
FILLARRY	OUT1, OUT2, OUTINAA, OUTTABLE
GETENTRY	REFORM
GETFILE	REFORM, MAKECARDS, SHUFFLE
GNAPSYMB	REFORM
INDNUM	GETENTRY
INIT	REFORM, MAKECARDS, SHUFFLE
LC2UC	REFORM, GETENTRY
LDEBLANK	GETENTRY, INDNUM, OUT1, OUT2
MAKECARDS	REFORM
NEWCMT	READLOCA, READFORM, READROCK, READCHAR, READCHEM, READISOT
NEWSOURC	READLOCA
NUMFIRST	INDNUM, BREAKSN
OUT1	REFORM
OUT2	REFORM
OUTINAA	REFORM
OUTSAMP	REFORM
OUTTABLE	REFORM

RDEBLANK	ASK, WRLOCA, WRFORM, WRROCK, WRCHAR, WRCHEM, WRISOT, GETENTRY, OUTINAA, OUT1, OUT2, SHUFFLE
READCHAR	REFORM
READCHEM	REFORM
READFORM	REFORM
READISOT	REFORM
READLOCA	REFORM
READINAA	READCHEM
READROCK	REFORM
READSHAW	READCHEM
READXRF	READCHEM
REMCHAR	OUTINAA, OUT1, OUT2
RJUST	WRCHEM, GETENTRY, INDNUM, FILLARRY, BREAKSN
RLNU	GETENTRY, FILLARRY
SHUFFLE	REFORM
WRCHAR	MAKECARDS
WRCHEM	MAKECARDS
WRFORM	MAKECARDS
WRISOT	MAKECARDS
WRLOCA	MAKECARDS
WRROCK	MAKECARDS



## DESCRIPTIONS OF SELECTED SUBROUTINES

The following program descriptions are in a FORTRAN-like pseudocode.

### REFORM.FOR

(This is the main driving routine.)

```
display main menu and get menu choice
if choice is "A" (make a new card file) call makecards.for
else if choice is "D" (sort a card file) call shuffle.for
else open input card file

    if choice is "B" (make a printout) open printout file
    else if choice is "C" (make a table)
        open table file
        get table entries
    else if choice is "E" (TOMGNAP file)
        display TOMGNAP menu and get choice
        open TOMGNAP file
        get plotting symbol
    else error, redisplay main menu
endif

do until end of input card file
    read next line of input file
    if this is the start of a new sample record
        (i.e.: if the first 5 characters of the
        line = "SAMPL") and it is not the first
        sample of the file, then reformat the
        last sample record and write it to the
        output file (e.g., call out1.for)
    else call appropriate subroutine to read the
        next block of data (e.g.: if first 5
        characters of the line are " CHAR" call
        readchar.for)
    endif
enddo
endif
reformat last sample record and write it to the output file
display on terminal the number of samples read, etc.
close files
```

### GETFILE

(Opens files. Parameters passed to GETFILE are unit, prompt, inout, form)

```
if inout=in
    status=old
    open: readonly, shared, sequential, formatted
else if inout=out
```

```

        status=new
        if form=formatted
            open: carriage control=list, sequential, formatted
        else if form=fortran
            open: carriage control=fortran, sequential,
                formatted
        else if form=unformatted
            open: sequential, unformatted
        endif
    else error
    endif

```

#### READ ROUTINES: READXXXX.FOR

(where "XXXX" is LOCA, FORM, ROCK, CHAR, CHEM, XRF, SHAW, NIAA, ISOT. Each of these subroutines reads a block of data from the sample entry into the appropriate variables)

```

do until error or end of block (blocks are terminated with
    the statement "COMMENT=***", or "*** for chemistry
    sections)
    read next card
    assign the data on the line to the appropriate
        variable(a)
enddo

```

#### TOMGNAP OUTPUT ROUTINES: OUT1.FOR, OUT2.FOR, OUTINAA.FOR

(These reformat chemistry data from a sample entry for TOMGNAP and write it to the output file. OUT1 reformats data from the ANALYSIS 1 chemistry section, OUT2 from ANALYSIS 2, and OUTINAA from ANALYSES 3 and part of 1)

```

get sample identifier (part of the sample number)
call fillarray (this assigns the data in the field variables
    to arrays and converts strings to real numbers)
scale numeric values for TOMGNAP (convert ppm to %, remove
    decimal points: TOMGNAP wants implicit decimal
    points)
write data, including plot symbol, in the proper format to
    the output file

```

#### GNAPSYMB

(Gets plotting symbol for TOMGNAP files. Called once by REFORM.FOR. TOMGNAP output routines write

```

plotsymb (min(# of chem samples so far, number of plot
    symbols))

```

Thus, samples beyond number of ASCII symbols in array (84) all get last symbol in array)

```
ask user if wants to assign one plotting symbol for all
  samples
if yes, read symbol and assign to each element of plot
  symbol array
else assign a unique plot symbol (ASCII characters A through
  @) to each element of plot symbol array
```

#### FILLARRY.FOR

(Called by the output routines, once for each sample)

```
Assign alphabetic-keyed variables to string array according
  to their keys.
get shawno, if present, for sample identification
convert lat/lon to decimal degrees
convert numeric-keyed variables to real numbers and assign
  them to real array according to their keys
```

#### OUTSAMP

Writes data for sample entries to a printer file, one or two samples per page, 126 columns per line. Called by REFORM.FOR, once per sample, after each sample entry's data is read from the card file.

Does no error-checking: writes fields by column position on the line so that despite invalid data or a shifted field the contents in the field's normal columns will be written.

Writes LOCATION, FORMATION, ROCK, and CHARACTERISTICS blocks even if no data in them (writes fieldnames with blanks for data). Writes ANALYSIS 1, 2, or 3, RATIOS, or ISOTOPES only if data is present.

Writes only those COMMENT, SOURCE, TEXTURE, and ISOTOPE ANALYSIS lines that are present.

#### MAKECARDS AND WRITE ROUTINES

(Prompts user for data and creates a new card file. Inserts field names and positions data in proper columns)

```
calls write subroutines for each data block
```

```
these subroutines (WRLOCA, etc.) call ASK.FOR which
prompts for the field, checks that it is not too long,
```

returns to the end of the calling write subroutine if escape character is typed. The write routines then write the data minus trailing blanks to the card file.

### SHUFFLE

(Sorts a card file according to a table. The table must contain index fields: the starting line number in the card file of the sample entry, and the number of lines in the sample entry)

```
open index (table) file, input card file, and output card
file
read input card file into string array of 80-character
elements
do until end of table file
  read indexes for next sample entry from table
  delete trailing blanks and write lines from start to
  end of sample entry per its indexes
```

### TABLE ROUTINES

The subroutines OUTTABLE.FOR, GETENTRY.FOR, INDNUM.FOR, and BREAKSN combine to produce a table of fields, with data for each sample on one line. The routines are relatively complicated because the user may specify any fields, the formats for each are different, and because most fields are truncated to enable more fields to fit on a line. For instance, some character fields are 40 characters long but commonly only the first few characters are used. Thus FORMATION NAME, for example, is 40 characters in card files but truncated to 20 characters in tables.

REFORM calls GETENTRY once, which calls INDNUM once per choice after it is specified by the user. OUTTABLE is called by REFORM once per sample, and calls BREAKSN each time if expanded sample numbers are specified.

The following table shows the data structures used by the table routines. Data flow is generally from top to bottom in the diagram. The examples show data for a table with the keys B, 102, 19, and D chosen.

# DATA STRUCTURES USED BY TABLE ROUTINES

VARIABLE	TYPE	EXAMPLE
<u>GETENTRY.FOR</u>		
expand	char*1	N (flag: break sample number into components?)
uflag	char*1	Y (flag: include index fields in table?)
tchoice	char*5	D (present choice of field key)
nchoices	int	4 (number of choices so far)
entry(nchoices)	char*3(30)	[ B 102 19 D ] (choices so far)
tablelen(nchoices)	int(150)	2 4 44 127 (integer keys as converted by [17 11 ... 11 ... 6 ... 9 ... ] INDNUM)
The tablelen block-data array holds the lengths of fields in their proper format for tables.		
ncol	int	37 (nbr of columns in table used so far)
<u>INDNUM.FOR</u>		
indnum(k)	int(150)	(converts alphabetic and numeric keys to integers 1 through 135)
tablenum(nchoices)	int(150)	[ 2 127 44 4 ] (keys A-Y => 1-25, 1-110 => 26-135)
Tablenum holds the converted keys for the fields chosen so far.		
<u>OUTTABLE.FOR</u>		
outtable(k)	int(30)	(writes data for one sample (one line) to table file)
ccateg(j)	int(30)	[127 44 ] (real categories (keys) from tablenum)
		[ 2 4 ] (character categories (keys) from tablenum)
		(j+k=nchoices)

```

id(ccateg(i), i=1, j) char*40(100) [ Old Woman Mts 30T11NR30E ]
numlist(vcateg(i), i=1, k) real(100) [ 107 1.0 ]

```

The arrays id and numlist hold data from fields with alphabetic keys and numeric keys respectively. These assignments are made by FILLARRY.FOR.

```
WRITE(FORCHR)(INDEX fields, id(ccateg(i)=1,j), numlist(rcateg(i),i=1,k))
```

This is the write statement for one line (sample) in a table. It uses the format in array FORCHR.

	2	4	44	127	converted {--keys ...]
tablefmt( )					
char*15(150)	[... A10,1X, ...	A10,1X, ...	F5.1,1X, ...	F8.4,1X, ...	

The above block data array holds the special table formats for each field (key).

```
FORCHR x      char*9(150) [(I5, I5,2X, A16,1X,  
!-----!  
formats for  
indexes & sample #  
!-----!  
A10,1X, A10,1X,  
!-----!  
formats for  
alphanumeric keys  
!-----!  
F8.4,1X, F5.1,1X,]  
127         44  
where x is expand uflag
```

1	y	y	The format statement can have one of four forms depending
2	y	n	on the state of the expand and uflag variables.
*3	n	y	
4	n	n	

### GETENTRY

(Gets user's choices of whether to expand sample numbers and include index fields in the table, and of fields (by keys) to include in the table)

```
prompt user if wants sample numbers expanded
prompt user if wants index fields in table
display menu of choices
prompt user to enter choice
do
  if choice= "XX" stop
  else if choice= "DD" then return
  else if choice= "SS" then start over building table
  else if valid field key
    put choice in array of choices so far
    call INDNUM (convert alphabetic and numeric keys to
      sequence of integers)
    sum number of columns in table used so far
    error if more than 126 columns (start over)
  else invalid choice, prompt user again for choice
```

### INDNUM

(Converts alphabetic and numeric keys to sequence of integers. Called by GETENTRY.FOR, once for each choice.)

```
if key is numeric convert string number to integer + 25
else convert alphabetic key to integer (A=1, Y=25)
endif
```

### OUTTABLE

(Writes fields for a sample entry in their table format to table file. Called by REFORM.FOR, once for each sample entry.)

```
call fillarray (fill data arrays with current sample entry
  data)
if expand flag=yes call breaksn (expand sample nbrs into
  components)
separate table choices into array of numeric keys (rcateg)
  and array of alphabetic keys (ccateg)
assemble proper format statement into a string array,
  depending on state of expand and index flags:

  first, formats for index fields (if any) and sample
    number (expanded if specified) are assigned to the
    array
  then, formats for the alphabetic-keyed fields
  finally, formats for the numeric-keyed fields
```

write using format statement just created:

```
index fields (if any)
sample number (expanded if specified)
fields with alphabetic keys: (id(ccateg(i)), i=1,j)
fields with numeric keys: (numlist(rcateg(i)), i=1,k)
```

#### BREAKSN

(Breaks (=expands) Needles-Project format sample number into component parts to facilitate sorting)

For example, the sample number BJ87CAL-238AX becomes

```
BJ    87    CAL  238  AX
2     2     3     3    2    <- size 12 + 4 spaces=16 cols
geol year range  sn  sna    <- fields
```



## MODIFYING REFORM

The REFORM software may be modified with relative ease. The table output subroutines are somewhat obscure and are dealt with in some detail in the subroutine descriptions above.

Modifications are of four types: (1) changing database structure, (2) adding new output routines, (3) altering the software to run on another system, and (4) more complicated changes. The latter might include, for instance, boolean search and selection options, and are not considered here since they would require significant additions.

After making changes to REFORM be sure to test the new version extensively. Tests should include extreme values, improperly formatted data (to check error-trapping), data for all new fields and for all new options. Add new error-trapping if needed. Finally, update the REFORM documentation.

### (1) CHANGING DATABASE STRUCTURE

The programming complexity involved in changing the structure of the database depends on the types of modifications required. The changes to specific subroutines are outlined below for several types of modifications.

Adding more comment, source, texture, and analysis lines:

The maximum allowable number of such lines is stored as the parameters ncmt, nsrsc, ntext, and nanal, respectively, in the variable file VAR.INC. Change the appropriate parameter to the new number of lines required. Note that changing ncmt changes the number of comment lines allowed for every data block.

Adding new fields to present data blocks:

Add the new variables to VAR.INC and COMMON.INC.

If the new variables are part of an existing chemistry analysis section or are part of the CHARACTERISTICS section and are best held in array form, add the field names to the appropriate block data arrays in BLDATA.FOR, and change the appropriate parameter in VAR.INC that gives the number of fields in the section (e.g., nshaw for ANALYSIS 1 section). Note that not all CHARACTERISTICS variables are included in block data, only the ones which are most easily handled as part of an array, such as mode data.

Assign keys to the new variables, and place the desired table format for the fields in the TABLEFMT array, and their lengths in the TABLELEN array, both in BLDATA.FOR.

Add the new variables and their keys to FILLARRY.FOR.

If any new character fields (alphabetic-keyed) are longer than 40 characters, increase the size of the id array size in VAR.INC. Change AMENUMAX and NMENUMAX in VAR.INC. These specify the largest numeric and alphabetic keys. Change INIT.FOR to

initialize the new variables.

Modify the read subroutines (e.g., READLOCA) for the data blocks with new variables to read those new variables from the card file.

Modify the write subroutines (e.g., WRCHAR) to prompt for the new variables and write them to new card files.

Modify the output subroutines to write the new variables to the output files. OUTSAMP.FOR must be modified for any new variables. OUT1.FOR, OUT2.FOR, or OUTINAA.FOR must be changed only if the new variables are chemistry data.

#### Adding a new data block:

For each of the new fields in the new data block make the changes listed above.

Write a new read subroutine to read in the fields in the new block. Model it after one of the existing read subroutines.

Change REFORM.FOR to call the new read subroutine. Note the read subroutines are called twice by REFORM.FOR.

Write a new write subroutine to prompt the user for the fields in the new data block and write them to new card files.

Change MAKECARDS.FOR to call the new write subroutine.

In addition to any other changes, if the total number of variables with alphabetic keys is greater than 25:

Increase the position of the first format and length for numeric-keyed fields in TABLEFMT and TABLELEN, respectively, both in BLDATA.FOR, from 25 to the new number of alphabetic keys (or larger).

Increase the size of arrays TABLEFMT, TABLELEN, ID, NUMLIST in VAR.INC if necessary.

Change "+25" in INDNUM.FOR to the new number of alphabetic characters.

Change "25" in OUTTABLE.FOR associated with the arrays RCATEG and CCATEG to the new number of alphabetic keys.

If the number of alphabetic keys is more than 26, change FILLARRY.FOR and its IND(AKEY) function to handle double alphabetic keys (e.g., AA). Note that DD,SS and XX are used by GETENTRY.FOR as commands. They must be changed, for instance to some word. Also, INDNUM1(TKEY) in INDNUM.FOR must be changed to handle double alphabetic keys.

If the total number of numeric keys is greater than 150, increase the sizes of arrays NUMLIST, TABLEFMT, and TABLELEN.

## NEW OUTPUT ROUTINES

In general, output routines will simply write data to a file in the desired format, possibly following scaling or conversion of some kind. The present routines, OUT1, OUT2, OUTINAA, and OUTTABLE call FILLARRY which converts numeric strings to real numbers and assigns variables to string or real arrays to simplify write statements. The output routines then scale the data as required for the particular application.

New output routines may be modeled on existing ones:

OUTSAMP.FOR for text printouts

OUTTABLE.FOR for tabular data

OUT1.FOR, OUT2.FOR, and OUTINAA.FOR for numeric  
(chemistry) data.

See the descriptions of these subroutines in this manual. Binary (non-ASCII) files may be written by opening an unformatted rather than formatted file (see GETFILE).

Output routines are called each time data for one whole sample in the card file has been read. REFORM.FOR must be modified to call the new output routine in two places. Modify the REFORM menu to list the new option.

For many purposes data may be output in table form with REFORM option C and used as input by another program. In such cases a new output routine will not be needed. The fields in the tables are separated by spaces, not commas, and strings are not enclosed in quotation marks. This may pose a problem for some programs. A new output routine similar to OUTTABLE could be made which adds these delimiters.

## PORTING REFORM TO OTHER SYSTEMS

The main considerations in installing REFORM on a system other than VAX/VMS are the VAX-11 FORTRAN extensions to the FORTRAN-77 standard that are used in REFORM (see list below), the sizes of data structures, and the sizes of program files.

Data structures are not large with the exception of the array CARD used in SHUFFLE.FOR. Its size is 20,000 80-character elements. It is used to hold an entire card file in main memory during sorting. If your system cannot support an array that large, a new memory-efficient scheme for sorting a card file may be needed.

Individual subroutines are not large but on a small system such as a PC not all may fit in memory at one time. It may be necessary to separate the various options into separate stand-alone programs that are called into memory as required.

### VAX FORTRAN EXTENSIONS USED IN REFORM

OPEN (...SHARED), used by GETFILE, allows programs to simultaneously access the file. If not supported then delete.

OPEN (...READONLY), used by GETFILE, write-protects files. If not supported then delete.

OPEN (CARRIAGE CONTROL=LIST), OPEN (CARRIAGE CONTROL=FORTRAN), used by GETFILE, specify carriage control, if not supported delete the parameter, and modify write statements in files opened with CARRIAGE CONTROL=LIST so that lines are single spaced (the first character of each line should be a space: ' ').

OPEN defaults to ACCESS=SEQUENTIAL. If this is not the system default, specify ACCESS=SEQUENTIAL.

"\$" format descriptor suppresses carriage return at end of line for prompts, used by REFORM, ASK, GETENTRY, GETFILE, and GNAPSYMB. If not supported then delete.

Symbolic names longer than six characters. If not supported replace by shorter names, being careful to change the names everywhere. The following routines use long names:

```
VAR.INC COMMON.INC ASK  GETENTRY GETFILE
INIT LDEBLANK OUT1 OUT2 OUTTABLE
RDEBLANK READLOCA READROCK REMCHAR SHUFFLE
```

INCLUDE/NOLIST. The NOLIST option keeps the contents of the include file from being listed in the compilation source listing, and is used in REFORM wherever INCLUDE is used (all subroutines except the short utility routines). Delete if not supported.

## COMPILING ON THE VMS/VAX

The source code for each of the REFORM subroutines is in a separate file. This allows each subroutine to be compiled individually to simplify debugging. The object files are placed in an object library which is then linked to create the executable module.

To compile new subroutines:

FOR <subroutine name> for each subroutine. This creates a compiled object file for each.

Then:

LIBR/CREATE ROCK.OLB	creates rock object library
LIBR/REPLACE ROCK *.OBJ	places the objects in the rock object library
DELETE *.OBJ.*	delete object files
FOR REFORM	compile the main loop
LINK REFORM ROCK/L/INCLUDE=BLDATA\$DATA	links modules
DELETE REFORM.OBJ.*	delete object file

```

c      reform.for
c      Reads standard card format input and outputs in new format.

      include 'var.inc/nolist'
      include 'common.inc/nolist'
      common/cardcount/ lastsamp

      character choice*1
      ncards=0
      nsamps=0
      shawtot=0
      lastsamp=1
c      xrftot=0
      do 10 i=1,80
10      blankcard(i:i)=' '

      print*, ' '
      print*, 'This is REFORM version 2.1, a database program'
      print*, ' for rock sample data.'

20      print*, ' '
      print*, 'These are your choices:'
      print*, ' '
      print*, '      A  Make a new card file'
      print*, '      B  Make a printout of a card file'
      print*, '      C  Make a table'
      print*, '      D  Sort a card file'
      print*, '      E  Reformat chem data for TOMGNAP'
      print*, ' '
      print*, '      X  (exit REFORM)'
      print*, ' '
      print '(/,1h$,a)', '*Your choice? '
      read '(a)', choice
      call lc2uc (choice)
      if (choice.eq.'A') then
         call makecards
         stop
      else if (choice.eq.'D') then
         call shuffle
         stop
      else if (choice.eq.'X') then
         stop
      else
         call getfile(21, '*Give card format infile: ', 'in', 'formatted')
         tmpfilename=filename

         if (choice.eq.'B') then
            call getfile(22, '*Give listing outfile:      ', 'out', 'fortran')

         else if (choice.eq.'C') then
            call getfile(28, '*Give table outfile:      ', 'out', 'formatted')
            call getentry

         else if (choice.eq.'E') then

```

```
c      reform.for
c      Reads standard card format input and outputs in new format.
```

```
include 'var.inc/nolist'
include 'common.inc/nolist'
common/cardcount/ lastsamp
```

```
character choice*1
ncards=0
nsamps=0
shawtot=0
lastsamp=1
```

```
c      xrftot=0
      do 10 i=1,80
10    blankcard(i:i)=' '
```

```
print*, ' '
print*, 'This is REFORM version 2.1, a database program'
print*, ' for rock sample data.'
```

```
20 print*, ' '
   print*, 'These are your choices:'
   print*, ' '
   print*, '      A  Make a new card file'
   print*, '      B  Make a printout of a card file'
   print*, '      C  Make a table'
   print*, '      D  Sort a card file'
   print*, '      E  Reformat chem data for TOMGNAP'
   print*, ' '
   print*, '      X  (exit REFORM)'
   print*, ' '
```

```
print ' (/,1h$,a)', '*Your choice? '
```

```
read ' (a)' . choice
```

call lc24c (choice)

```
if (choice.eq.'A') then
```

call makercards

stop

```

else if (choice.eq.'D') then

```

call shuffle

stop

```

else if (choice.eq.'X') then

```

stop

else

```
call getfile(21,'*Give card format infile: '.'in','formatted')
```

```
tmpfilename=filename
```

```
if (choice.eq.'B') then
```

```
call getfile(22,'*Give listing outfile:      ','out','fortran')
```

```
else if (choice.eq.'C') then
```

```
call getfile(28,'*Give table outfile:      ','out','formatted')
```

```
call getentry
```

```
else if (choice.eq.'E') then
```

```

30      print*, ' '
      print*, 'This is the Reform option E menu.'
      print*, ' '
      print*, 'The following routines reformat data
& for TOMGNAP.'
      print*, ' '
      print*, 'You may choose to reformat data from:'
      print*, ' '
      print*, '      1  Analysis 1'
      print*, '      2  Analysis 2'
      print*, '      3  Analyses 3 and 1 combined'
      print*, ' '
      print*, '      X  (exit REFORM)'
      print*, ' '
      print ' (/,1h$,a)', '*Your choice? (1,2,3,or X): '
      read '(a)', choice
      if (choice.eq.'x'.or.choice.eq.'X') then
         close(21)
         stop
      else if (choice.eq.'1'.or.choice.eq.'2'.or.choice.eq.'3') then
         call getfile(30,'*Give tomgnap outfile:      ','out','formatted')
         call gnapsymb
      else
         print*, 'Try again'
         print*, ' '
         goto 30
      endif
endif

      else
         print*, 'Try again'
         print*, ' '
         goto 20
      endif
endif

100  read(21,'(a80)',end=900) card
      ncards=ncards+1
      datatype=card(1:5)

      if(card.eq.blankcard) then
         continue
      else if(datatype.eq.'SAMPL') then
c      New sample...output previous sample data...
         if(nsamps.gt.0) then
            if (choice.eq.'B')call outsamp
            if (choice.eq.'C')call outtable
            if (choice.eq.'1')call out1 (symdex)
            if (choice.eq.'2') call out2 (symdex)
            if (choice.eq.'3') call outinaa (symdex)
         endif
         call init
         sample=card(11:30)
         nsamps=nsamps+1
      else if(datatype.eq.' LOCA') then
         call readloca
      else if(datatype.eq.' FORM') then

```

```

        call readform
    else if(datatype.eq.' ROCK') then
        call readrock
    else if(datatype.eq.' CHAR') then
        call readchar
    else if(datatype.eq.' CHEM') then
        call readchem
    else if(datatype.eq.' ISOT') then
        call readisot
    else
        print *, ' Unknown datatype at line',ncards
        print *, '   after sample number... ',sample
        print *, ' line reads:'
        print *,   card
        print *, ' Trying to continue...'
    endif
endif

goto 100

c      Output last sample if not already done...
900    if(sample.ne.' ') then
        if (choice.eq.'B')call outsamp
        if (choice.eq.'C')call outtable
        if (choice.eq.'1')call out1   (symdex)
        if (choice.eq.'2')call out2 (symdex)
        if (choice.eq.'3')call outinaa (symdex)
    endif
        if (choice.eq.'1'.or.choice.eq.'2'.or.choice.eq.'3')
&      print*, 'Number of analysis samples read = ',shawtot

print*, 'Number of cards read = ',ncards
print*, 'Number of samples read = ',nsamps

close(21)
close(22)
close(28)
close(30)
stop
end

```





```

dimension alysis(nanal), alyst(nanal), mineral(nanal),
&    age(nanal), isotratio(nanal)
character alysis*5, alyst*10, mineral*5, age*20, isotratio*25
character*50 isotcmt(ncmt)

```

c table variables:

```

character expand*1, entry(30)*3, tablefmt(150)*15, tmpfilename*13
character uflag*1
integer tablelen(150), tablenum(150), nchoices

```







```

&          'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,',
&          'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,',
&          'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,',
&          'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,',
&          'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,', 'F9.4,1X,' /
data (tablefmt(i), i=126,150) /
&          'F8.4,1X,',
&          'F8.4,1X,', 'F8.4,1X,', 'F8.4,1X,', 'F9.3,1X,', 'F9.3,1X,',
&          'F7.2,1X,', 'F8.4,1X,', 'F7.2,1X,', 'F8.4,1X,', ' ',
&          ' ', ' ', ' ', ' ', ' ', ' ', ' ',
&          ' ', ' ', ' ', ' ', ' ', ' ', ' ',
&          ' ', ' ', ' ', ' ', ' ', ' ', ' /

data tablelen/17,11,16,11,16,21,21,21,09,21,06,11,06,06,11,05,
&          21,21,21,04,12,0,0,0,0,
&          8,9,7,5,8,6,6,6,6,6,6,6,6,6,6,6,6,
&          6,6,6,6,6,5,6,6,6,4,6,6,6,6,5,6,6,6,7,6,
&          6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,6,
&          6,6,6,6,6,6,6,7,6,6,6,7,7,7,10,10,10,10,10,10,
&          10,10,10,10,10,10,10,10,10,10,10,10,10,10,10,10,
&          10,10,10,10,10,9,9,9,9,10,10,8,9,8,9,0,0,0,0,0,0,
&          0,0,0,0,0,0,0,0,0,0 /
end

```

```

c cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      subroutine breaksn(samp,geol,ye ar,rang,sn,sna)
c   Breaks a sample number into...
c       geologist
c       year
c       range
c       number (numeric part)
c       number (alphabetic part, if any)

      character*(*) samp,geol,ye ar,rang,sn,sna

      l=len(samp)
      i1=numfirst(samp)
      geol=samp(1:i1-1)
      ye ar=samp(i1:i1+1)

      i2=numfirst(samp(i1+2:l))+i1+1
      i3=chrfirst(samp(i2:l))+i2-1
      if(samp(i2-1:i2-1).eq.'-') samp(i2-1:i2-1)= ' '
      rang=samp(i1+2:i2-1)
      sn=samp(i2:i3-1)
      sna=' '
      if(chrfirst(samp(i2:l)).ne.0) sna=samp(i3:l)

c   Right-justify the fields

      call rjust (geol)
      call rjust (ye ar)
      call rjust (rang)
      call rjust (sn)
      call rjust (sna)

      return
end

c cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      subroutine carderror(nkards)
c   Sends error message if error in card format.

      integer nkards

      print *, ' Error trying to read in line', nkards

      return
end
```

```

function chrfirst(string)
c    Finds the position of the first alphabetic character in a string.
    character string*(*)

```

```
l=len(string)
chrfirst=0
```

```
do 20 i=1,1
if(string(i:i).ge.'A'.and.string(i:i).le.'Z') then
  chrfirst=i
  return
```

```
endif
20 continue
```

```
return
end
```

```

c
c      subroutine ckblank(array,n,notblank)
c      Counts how many entries in a character array are not blank.
c      character blank*80, array(n)*(*)

```

```
blank( : )=' '
notblank=0
```

```

      do 10 i=1,n
10      if(array(i).ne.blank) notblank=notblank+1

```

```
return
end
```





```

xlat=decdeg(xlatd,xlatm,xlats)

numlist(1)=xlat
numlist(2)=xlon
numlist(3)=rlnu(sg)
numlist(4)=rlnu(ci)
numlist(5)=rlnu(mag)
do 110 i=6,24
110 numlist(i)=rlnu(charval(i-5))
do 120 i=25,54
120 numlist(i)=rlnu(shawval(i-24))
do 130 i=55,71
130 numlist(i)=rlnu(xrfval(i-54))
do 140 i=72,100
140 numlist(i)=rlnu(niaaval(i-71))
do 150 i=101,106
150 numlist(i)=rlnu(ratio(i-100))
numlist(107)=rlnu(age(1))
numlist(108)=rlnu(isotratic(1))
numlist(109)=rlnu(age(2))
numlist(110)=rlnu(isotratic(2))

return

end

```

```

c      subroutine getentry
c          Prompts user for which fields are to make up the table,
c          any combination up to 126 columns wide (except that the
c          sample number is automatically included as a choice)

        include 'var.inc/nolist'
        include 'common.inc/nolist'
        character tchoice*5

10     print*, ' '
        print'(/,1h$,a)', '*Expand (Needles format) sample numbers
& (Y/N)? '
        read '(a)', expand
        call lc2uc (expand)

        print*, ' '
        print'(/,1h$,a)', '*Index fields (Y/N)? '
        read '(a)', uflag
        call lc2uc (uflag)

        print*, ' '
        print*, 'Enter a key or one of the following command keys:'
        print*, ' '

        nchoices=0

100    print*, ' '
        print*, '           SS   start over building table '
        print*, '           DD   done building table, start processing'
        print*, '           XX   (exit to VAX system) '
        print*, ' '

        print'(/,1h$,a)', 'Choice? '
        read '(a)', tchoice
        call lc2uc (tchoice)

        call ldeblank(tchoice,tchoice,lnt)
        call rdeblank(tchoice,tchoice,lngth)
        if (lngth.gt.3) then
            print*, 'Invalid choice (too long). Try again please.'
            goto 100
        endif
        if (tchoice.eq.'XX') then
            stop
        else if (tchoice.eq.'DD') then
            goto 200
        else if (tchoice.eq.'SS') then
            print*, 'Start over now.'
            goto 10
        else if ((lngth.eq.1).and.(tchoice.ge.'A'.and.tchoice.le.
& amenumax)) then
            goto 120
        else if (tchoice(1:1).lt.':') then
            if (rlnu(tchoice).ge. 1.0 .and.rlnu(tchoice)
& .le. nmerumax) then

```

```

        goto 120
    else
        goto 170
    endif
120    nchoices=nchoices+1
        entry(nchoices)=tchoice
        call rjust(entry(nchoices))
        print*, 'Your choices so far are: ', (entry(i), i=1, nchoices)
        call indnum(nchoices)
        ncols=17
        do 150 i=1, nchoices
            ncols=ncols+tablelen(tablenum(i))
150        continue
            if (uflag.eq.'Y') ncols=ncols+12
            if (ncolms.gt.126) then
                print*, 'You''ve used ', ncols, ' columns which is
&                more than 126; please start over'
                goto 10
            else
                print*, 'Columns used up so far = ', ncols
            endif
        else
170        print*, 'Invalid choice, try again please.'
        endif
        goto 100

200    return
    end

```

```

c      subroutine getfile(unit,question,inout,form)
c      Opens files...filenames not recognized by VAX will generate
c      an error message and request a second try.

c      Variables:
c      unit = io unit
c      question = query requesting file name
c      inout = 'in' or 'out'
c      form = 'unformatted' (binary) or
c            'formatted' (ascii with no carriage control) or
c            'fortran' (ascii with fortran carriage control).

c      Sample call:
c      call getfile(21,'*Give input file:', 'in','formatted')

include 'var.inc/nolist'
include 'common.inc/nolist'
integer unit
character status*8
character*(*) question, inout, form

10 write(*,101) question
101 format(/,'$',a,' ')
read '(a50)', filename

if(inout.eq.'in') then
    status='old'
    open(unit,file=filename,form=form,status=status,err=900,
&        readonly,shared)
c    Note that readonly and shared are VAX specific...

else if(inout.eq.'out') then
    status='new'

    if(form.eq.'formatted') then
        open(unit,file=filename,form=form,status=status,
&        carriagecontrol='list',err=900)
    else if(form.eq.'fortran') then
        open(unit,file=filename,form='formatted',status=status,
&        carriagecontrol='fortran',err=900)
    else if(form.eq.'unformatted') then
        open(unit,file=filename,form=form,status=status,err=900)
    endif
else
    print *, 'ERROR...inout variable in sub getfile not recognized.'
    print *, '   inout =', inout
    stop
endif
return

900 print *,' ERROR IN OPENING ',filename
print *,'       try again.....'
close(unit)
go to 10
end

```





```

c cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      subroutine init
c      Reinitializes variables.

      include 'var.inc/nolist'
      include 'common.inc/nolist'
      blankcard=' '

c      sample & category:
          sample=blankcard
          do 10 i=1,10
10       category(i)=blankcard
          shawno=blankcard

c      location:
          range=blankcard
          quad=blankcard
          trs=blankcard
          do 20 i=1,ncmt
20       locacmt(i)=blankcard
          do 25 i=1,nsrc
25       source(i)=blankcard
          lat=blankcard
          lon=blankcard

c      formation:
          formsymb=blankcard
          formname=blankcard
          do 30 i=1,ncmt
30       formcmt(i)=blankcard

c      rock types:
          rocktype=blankcard
          grainsize=blankcard
          do 40 i=1,ntext
40       texture(i)=blankcard
          do 50 i=1,ncmt
50       rockcmt(i)=blankcard

c      characteristics:
          cps=blankcard
          sg=blankcard
          ci=blankcard
          mag=blankcard
          do 60 i=1,nch
60       charval(i)=blankcard
          othermin=blankcard
          do 70 i=1,ncmt
70       charcmt(i)=blankcard

c      chemistry:
          do 80 i=1,nshaw
80       shawval(i)=blankcard
          do 82 i=1,nxrf
82       xrfval(i)=blankcard
          do 84 i=1,nniaa

```



```

84   niaaval(i)=blankcard
      do 90 i=1,nrat
90   ratio(i)=blankcard
      do 100 i=1,ncmt
100  chemcmt(i)=blankcard
      analyst1=blankcard
      analyst2=blankcard
      analyst3=blankcard

c   isotopes:
      do 110 i=1,nanal
      alysis(i)=blankcard
      alyst(i)=blankcard
      mineral(i)=blankcard
      age(i)=blankcard
110  isotratio(i)=blankcard
      do 120 i=1,ncmt
120  isotcmt(i)=blankcard

      return
      end

```



```

subroutine makecards

```

```
c      Asks for data interactively and makes standard card format output.
c      (Also deletes trailing blanks from long strings (e.g., COMMENTS))
```

```
include 'var.inc/nolist'
include 'common.inc/nolist'
```

```
call getfile(22,'*Give card outfile: ','out','formatted')
```

```
100 print '(/,a,/)', ' Answer with "'\' to skip a block or to quit.'
    print '(a./)', ' Answer with "return" for a blank.'
```

```
call init
```

```
call ask('sample no (20)= ',sample,*900)
call rdblank(sample,sample,1)
write(22,1100) sample(1:1)
```

```
1100 format('SAMPLE NO=', a)
```

```
call wrloca
call wrform
call wrrock
call wrchar
call wrchem
call wrisot
```

nota 100

```
900  close(22)
      stop
      end
```

```

subroutine wrloca

```

c Writes cards in a location block.

```
include 'var.inc/nolist'
include 'common.inc/nolist'
```

```
print *, 'LOCATION BLOCK:'
```

```
call ask('mountain range (20)=' ,range,*999)
call rdeblank (range,range,1)
write(22,1100)
```

```
1100 format(' LOCATION:')
```

```
write(22,1110) range(1:1)
```

```
1110 format(10x,'MOUNTAIN RANGE=',.a)
```

```
call ask('latitude (6)=' ,lat,*900)
call ask('longitude (7)=' ,lon,*900)
write(22,1200) lat,lon
```

```
1200 format(10x,'LATITUDE=',a6,4x,'LONGITUDE=',a7)
```







```

        call ask('chemistry??? [\ or return]: ',chemans,*999)

        write(22,1000)
1000 format(' CHEMISTRY:')

```

#### c...Shaw block

```

        print *, ' ANALYSIS 1 BLOCK: (eg:SHAW)'
        call ask('analysis 1 by(50):',analyst1,*199)
        call rdeblank(analyst1,analyst1,1)
        do 110 i=1,nshaw
110    call rjust(shawname(i))
        call ask(shawname(i)//' =' ,shawval(1),*199)
        do 150 i=2,nshaw
150    call ask(shawname(i)//' =' ,shawval(i),*190)
190    write(22,1100)analyst1(1:1)
1100 format(10x,' ANALYSIS 1 BY: ',a)
        write(22,1110) (shawname(i),shawval(i),i=1,nshaw)
1110 format((10x,5(a6,'=' ,a7)))
        write(22,1120)
1120 format(10x,8x,'###')
199    continue

```

#### c...xrf block

```

        print *, ' ANALYSIS 2 BLOCK: (eg:USGS(XRF))'
        call ask('analysis 2 by(50):',analyst2,*299)
        call rdeblank(analyst2,analyst2,1)
        do 210 i=1,nxrf
210    call rjust(xrfname(i))
        call ask(xrfname(i)//' =' ,xrfval(1),*299)
        do 250 i=2,nxrf
250    call ask(xrfname(i)//' =' ,xrfval(i),*290)
290    write(22,2000)analyst2(1:1)
2000 format(10x,' ANALYSIS 2 BY: ',a)
        write(22,1110) (xrfname(i),xrfval(i),i=1,nxrf)
        write(22,1120)
299    continue

```

#### c...niaa block

```

        print *, ' ANALYSIS 3 BLOCK: (eg:USGS(INAA))'
        call ask('analysis 3 by(50):',analyst3,*399)
        call rdeblank(analyst3,analyst3,1)
        do 310 i=1,nniaa
310    call rjust(niaaname(i))
        call ask(niaaname(i)//' =' ,niaaval(1),*399)
        do 350 i=2,nniaa
350    call ask(niaaname(i)//' =' ,niaaval(i),*390)
390    write(22,3000)analyst3(1:1)
3000 format(10x,' ANALYSIS 3 BY: ',a)
        write(22,1115) (niaaname(i),niaaval(i),i=1,nniaa)
1115 format((14x,4(a2,'=' ,a10,1x)))
        write(22,1120)
399    continue

```





```

299  continue

      do 500 i=1,ncmt
      call ask('comment (50)=' , isotcmt(i),*900)
      call rdeblank(isotcmt(i),isotcmt(i),1)
      write(22,1500) isotcmt(i)(1:1)
1500 format(10x,'COMMENT=',a)
500  continue

900  write(22,1400)
1400 format(10x,'COMMENT=###')

999  return
      end

```







```

        write(30,1100)plotsymb(min(shawtot+1,nsyms)),idata(25),
&      idata(27),idata(28),idata(29),
&      idata(31),idata(32),idata(33),idata(34),idata(36),idata(26),
&      idata(35),idata(30),idata(38),ident

        write(30,1200)plotsymb(min(shawtot+1,nsyms)),idata(40),
&      idata(41),idata(42),idata(43),idata(44),idata(45),idata(46),
&      idata(47),idata(48),idata(49),idata(50),idata(51),idata(52),
&      idata(53),idata(54),ident

1100      format( 'NRM',a1,12I4,3x,I4,2x,3x,3x,3x,2x,a8)
1200      format( 'NRM',a1,15I4,8x,a8)
        shawtot=shawtot+1

endif

return
end

```



```

c      subroutine outinaa (symdex)
c      Outputs 3 analysis cards for tomgnap
c      of analysis 1 and analysis 3 chemistry

      include 'var.inc/nolist'
      include 'common.inc/nolist'
c      common/plotf/ ckey, nkey
      common/index/ i

      character ident*8
      integer idata (100)


c      Get last 8 characters of sample number for ident
      call rdeblank(sample,sample1,ns)
      if(ns.gt.8) then
        call remchar ('-',sample,sample1,12)
        call rdeblank(sample1,sample1,ns)
      else
      endif
      ident=sample1(ns-7:ns)


      call fillarry


      idata(1)=0
      do 210 i=25,38
210    idata(i)=nint(100.0*numlist(i))


      idata(40)=nint(numlist(40))
      idata(41)=nint(numlist(41))
      idata(42)=nint(numlist(42))
      idata(44)=nint(numlist(44))
      idata(53)=nint(numlist(53))
      idata(54)=nint(numlist(54))
      idata(75)=nint(numlist(75))
      idata(83)=nint(numlist(83))


      idata(80)=nint(10.0*numlist(80))
      idata(81)=nint(10.0*numlist(81))
      idata(87)=nint(10.0*numlist(87))
      idata(91)=nint(10.0*numlist(91))


      idata(73)=nint(100.0*numlist(73))
      idata(74)=nint(100.0*numlist(74))
      idata(79)=nint(100.0*numlist(79))
      idata(89)=nint(100.0*numlist(89))
      idata(90)=nint(100.0*numlist(90))
      idata(92)=nint(100.0*numlist(92))
      idata(94)=nint(100.0*numlist(94))
      idata(96)=nint(100.0*numlist(96))


      idata(72)=nint(1000.0*numlist(72))

```

```

idata(76)=nint(1000.0*numlist(76))
idata(77)=nint(1000.0*numlist(77))
idata(78)=nint(1000.0*numlist(78))
idata(84)=nint(1000.0*numlist(84))
idata(85)=nint(1000.0*numlist(85))
idata(86)=nint(1000.0*numlist(86))
idata(88)=nint(1000.0*numlist(88))
idata(93)=nint(1000.0*numlist(93))
idata(97)=nint(1000.0*numlist(97))
idata(98)=nint(1000.0*numlist(98))
idata(99)=nint(1000.0*numlist(99))

idata(82)=nint(10000.0*numlist(82))
idata(95)=nint(10000.0*numlist(95))

write(29,1100)plotsymb(min(shawtot+1,nsyms)),idata(25),
& idata(26),idata(27),idata(28),idata(29),idata(30),idata(31),
& idata(32),idata(33),idata(34),idata(35),idata(36),idata(37),
& idata(38),idata(40),idata(41),idata(42),idata(44),ident

write(29,1200)plotsymb(min(shawtot+1,nsyms)),idata(53),
& idata(54),idata(72),idata(73),idata(74),idata(75),idata(76),
& idata(77),idata(78),idata(79),idata(80),idata(81),idata(82),
& idata(83),idata(84),idata(85),ident

write(29,1300)plotsymb(min(shawtot+1,nsyms)),idata(86),
& idata(87),idata(88),idata(89),idata(90),idata(91),idata(92),
& idata(93),idata(94),idata(95),idata(96),idata(97),idata(98),
& idata(99),ident

1100 format( 'NRM',a1,I4,I3,3I4,I2,4I4,I3,3I4,I3,2I4,I3,2X,a8)
1200 format( 'NRM',a1,i4,i2,i5,3i4,i5,i6,2i4,i5,4i4,i5,a8)
1300 format( 'NRM',a1,i4,4i5,4i4,i5,2i4,i5,i4,6x,a8)
shawtot=shawtot+1

return
end

```



```

c-----
      subroutine outsamp
c      Outputs a sample.

```

```
c      Outputs a sample.
```

```
include 'var.inc/nolist'
```

include 'common-inc/polist'

```
character spacer*126, space1*126, space2*126
```

```
do 10 i=1,126
```

```
spacer(i:i)=' $'
```

```
space1(i:i)=' -'
```

```
10 space2(i:i)=' '
```

```
write(22,1000) spacer
```

```
1000 format(/, '1', a126)
```

do 1008 i=1,3

```
write(22,1005)blankcard
```

1008 continue

```
1005 format(' ',a80)
```

C... LOCATION

```
write(22, 1010) 'SAMPLE:', sample, 'RANGE:', range, 'QUAD:', quad,
```

```
& 'TRS:', trs
```

```
1010 format('/', ' ', a7, 1x, a20, 5x, a6, 1x, a20, 5x, a5, 1x, a30, 5x, a4, 1x, a10)
```

```
write(22, 1012) space1(1:20)
```

```
1012 format(' ', 8x, a20)
```

```
write(22,1016) lat(1:2),lat(3:4),lat(5:6).
```

```
& lon(1:3),lon(4:5),lon(6:7)
```

```
1016 format(' ',18x,'lat=',a2,':',a2,':',a2,5x,'lon=',a3,':',a2,':',a2)
```

```
do 25 i=1,nsrc
```

```
25  if(source(i).ne.blankcard)      write(22,1025) source(i)
```

```
1025  format(' ',18x,' source: ',a50)
```

```
do 20 i=1,ncmt
```

```
20  if(locacmt(i).ne.blankcard)      write(22,1020) locacmt(i)
```

```
1020      format(' ',18x,'      comment: ',a50)
```

### C...FORMATION

```
write(22,1040) 'FORMATION:',formsymb,formname
```

```
1040 format('/', ' ', a15, 3x, 'symbol=', a20, ' name=', a40)
```

```
do 40 i=1,ncont
```

```
40      if(formcnt(i).ne.blankcard)      write(22,1020) formcnt(i)
```

C... ROCK TYPE

```
write(22,1060) 'ROCK TYPE:',rocktype,grainsize
```

```
1060 format('/', ' ', a15, 3x, a40, 5x, 'grain size: ', a40)
```

```
do 60 i=1,ntext
```

```
60    if(texture(i).ne.blankcard)      write(22,1062) texture(i)
```

```
1062 format(' ',18x,' texture: ',a50)
```

```
do 65 i=1, ncont
```

```
65   if(rockcmt(i).ne.blankcard)      write(22,1020) rockcmt(i)
```

### c...CHARACTERISTICS

```
      write(22,1080) 'CHARACTER:',cps,sg,ci,mag
1080 format(/,' ',a15,3x,'cps=',a12,' sg=',a5,'      ci=',a5,' mag=',a7)
      write(22,1081) (charname(i),charval(i), i=1,5)
1081 format(' ',18x,'slab:      ',7(1x,a4,' : ',a5))
      write(22,1082) (charname(i),charval(i),i=6,12)
1082 format(' ',18x,'thin section:',7(1x,a4,' : ',a5))
      write(22,1084) (charname(i),charval(i),i=13,19)
1084 format(' ',18x,'modes:      ',7(1x,a4,' : ',a5,'%'))
      write(22,1086) othermin
1086 format(' ',18x,'other minerals = ',a25)
      do 80 i=1,ncmt
80   if(charcmt(i).ne.blankcard)      write(22,1020) charcmt(i)
```

### c...CHEMISTRY

```
      call ckblank(shawval,nshaw,numshaw)
      call ckblank(xrfval,nxrf,numxrf)
      call ckblank(niaaval,nniaa,numniaa)
      call ckblank(ratio,nrat,numrat)

      if(numshaw.ne.0.or.numxrf.ne.0.or.numniaa.ne.0.or.
& numrat.ne.0) then
      write(22,2100) 'CHEMISTRY:'
2100 format(/,' ',a15,3x,a108)

      if(numshaw.ne.0) then
      write(22,2110) 'analyzed by: ',analyst1
2110 format(' ',18x,a12,a50)
      write(22,2250) space2(1:108)
      write(22,2200) (shawname(i), i=1,15)
2200 format(' ',15x,3x,4x,15(a5,2x))
      write(22,2210) (shawval(i), i=1,15)
2210 format(' ',18x,' % ',15a7)
      write(22,2250) space1(1:108)
      write(22,2220) (shawname(i), i=16,30)
2220 format(' ',18x,5x,15(a5,2x))
      write(22,2230) (shawval(i), i=16,30)
2230 format(' ',18x,' ppm ',15a7)
      write(22,2250) space2(1:108)
2250 format(' ',18x,a108)
      endif

      if(numxrf.ne.0) then
      write(22,2110) 'analyzed by: ',analyst2
      write(22,2250) space2(1:108)
      write(22,2200) (xrfname(i), i=1,10)
      write(22,2210) (xrfval(i), i=1,10)
      write(22,2250) space1(1:108)
      write(22,2220) (xrfname(i), i=11,16)
      write(22,2230) (xrfval(i), i=11,16)
      write(22,2250) space2(1:108)
      endif
```

```

        if(numniaa.ne.0) then
            write(22,2110) 'analyzed by: ',analyst3
            write(22,2250) space2(1:108)

            write(22,2270) (niaaname(i), i=1,3)
2270 format(' ',15x,4x,9(5x,a2,4x))
            write(22,2275) (niaaval(i), i=1,3)
2275 format(' ',18x,' % ',3(1x,a10))
            write(22,2250) space1(1:108)

            write(22,2270) (niaaname(i), i=4,12)
            write(22,2280) (niaaval(i), i=4,12)
2280 format(' ',18x,' ppm ',9(1x,a10))
            write(22,2250) space1(1:108)

            write(22,2270) (niaaname(i), i=13,21)
            write(22,2280) (niaaval(i), i=13,21)
            write(22,2250) space1(1:108)

            write(22,2270) (niaaname(i), i=22,28)
            write(22,2280) (niaaval(i), i=22,28)
            write(22,2250) space2(1:108)
        endif

        if(numrat.ne.0) then
            write(22,2110) 'ratios:'
            write(22,2250) space2(1:108)
            write(22,2300) (rationame(i), i=1,6)
2300 format(' ',15x,3x,4x,6(7x,a5,2x))
            write(22,2310) (ratio(i), i=1,6)
2310 format(' ',18x,4x,6(7x,a7))
            write(22,2250) space2(1:108)
        endif

        do 280 i=1,ncmt
280  if(chemcmt(i).ne.blankcard)      write(22,1020) chemcmt(i)

        endif

c... ISOTOPES

        if(alysis(1).ne.blankcard) then
            write(22,2100) 'ISOTOPES:'

            do 300 i=1,nanal
300  if(alysis(i).ne.blankcard)
            & write(22,3100) alysis(i),alyst(i),mineral(i),age(i),isotratio(i)
3100 format(' ',18x,' analysis=',a5,2x,' analyst=',a10,2x,' mineral=',
            & a5,2x,' age=',a20,2x,' ratio=',a25)
c      write(22,2250) space2(1:108)
            do 380 i=1,ncmt
380  if(isotcmt(i).ne.blankcard)      write(22,1020) isotcmt(i)
            endif
            return
        end

```

```

c      subroutine outtable
c      Outputs a 126-column table of user specified data.

      include 'var.inc/nolist'
      include 'common.inc/nolist'
c      common/plotf/ ckey, nkey
      common/index/ i
      common/cardcount/ lastsamp

      character space2*126, tstfmt*30
      character*9 forchr1(40), forchr2(40), forchr3(40), forchr4(40)
      character geol*2, year*2, rang*3, sn*3, sna*2
      integer ccateg(30),rcateg(30)

      data (forchr1(i), i=1,7) /' (i5,', 'i5,2x,', 'a2,1x,', 'a2,1x,',
&      'a3,1x,', 'a3,1x,', 'a2,1x,' /
      data (forchr2(i), i=1,5) /' (a2,1x,', 'a2,1x,', 'a3,1x,', 'a3,1x,',
&      'a2,1x,' /
      data (forchr3(i), i=1,3) /' (i5,', 'i5,2x,', 'a16,1x,' /
      data forchr4(1)      /' (a16,1x,' /

      do 10 i=1,126
10      space2(i:i)=' '

      newsamp=ncards
      nfirst=lastsamp
      numfirst=newsamp-lastsamp

      call fillarry

      if (expand.eq.'Y') call breaksn(sample,geol,year,rang,
&      sn,sna)

      j=0
      k=0
      do 200 i=1,nchoices
          if (tablenum(i).gt.25) then
              k=k+1
              rcateg(k)=tablenum(i)
          else
              j=j+1
              ccateg(j)=tablenum(i)
          endif
200      continue

      if (expand.eq.'Y') then
          if (uflag.eq.'Y') then

              do 300 i=8,j+7
                  forchr1(i)=tablefmt(ccateg(i-7))
300              continue
              do 400 i=8+j,j+k+7
                  forchr1(i)=tablefmt(rcateg(i-7-j))
400              continue

```

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        forchr1(j+k+7)(8:8)=')'
        write(28, forchr1) nfirst, numfirst, geol, year, rang, sn, sna,
&      (id(ccateg(i)), i=1,j), (numlist(rcateg(i)-25), i=1,k)
    else
        do 500 i=6,j+5
            forchr2(i)=tablefmt(ccateg(i-5))
500        continue
            do 600 i=6+j,j+k+5
                forchr2(i)=tablefmt(rcateg(i-5-j))
600        continue
            forchr2(j+k+5)(8:8)=')'
            write(28, forchr2) geol, year, rang, sn, sna,
&      (id(ccateg(i)), i=1,j), (numlist(rcateg(i)-25), i=1,k)
        endif
    else
        if (uflag.eq.'Y') then
            do 700 i=4,j+3
                forchr3(i)=tablefmt(ccateg(i-3))
700        continue
                do 800 i=4+j,j+k+3
                    forchr3(i)=tablefmt(rcateg(i-3-j))
800        continue
                forchr3(j+k+3)(8:8)=')'
                write(28, forchr3) sample, nfirst, numfirst,
&      (id(ccateg(i)), i=1,j), (numlist(rcateg(i)-25), i=1,k)
            else
                do 900 i=2,j+1
                    forchr4(i)=tablefmt(ccateg(i-1))
900        continue
                    do 1000 i=j+2,k+j+1
                        forchr4(i)=tablefmt(rcateg(i-j-1))
1000       continue
                    forchr4(j+k+1)(9:9)=')'
                    write(28, forchr4) sample, (id(ccateg(i)), i=1,j),
&      (numlist(rcateg(i)-25), i=1,k)
                endif
            endif

        lastsamp=newsamp

        return
    end

```



```

c
subroutine readchar
  Reads cards in characteristics block.

  include 'var.inc/nolist'
  include 'common.inc/nolist'

  icmt=0
100  read(21,'(a80)',end=900) card
     ncards=ncards+1
     if(card.eq.blankcard) then
       continue

     else if(card(1:10).ne.'          ') then
       print *, 'Charact. block not properly ended before line',ncards
       call badcard (card,sample)
       print*, 'Trying to continue ...'
       return

     else if(card(11:21).eq.'COMMENT=$$$') then
       return
     else if(card(11:14).eq.'CPS=') then
       read(card,1010,err=990) cps, sg, ci, mag
1010  format(14x,a12,4x,a5,4x,a5,5x,a7)

     else if(card(11:13).eq.'SN=') then
       read(card,1020,err=990) (charval(i), i=1,5)
1020  format(13x,a5,4(4x,a5))

     else if(card(11:13).eq.'TN=') then
       read(card,1030,err=990) (charval(i), i=6,12)
1030  format(13x,a5,6x,a5,3(4x,a5),2(5x,a5))

     else if(card(11:14).eq.'%Q1=') then
       read(card,1040,err=990) (charval(i), i=13,19)
1040  format(14x,a5,6(5x,a5))
     else if(card(11:25).eq.'OTHER MINERALS=') then
       othermin=card(26:50)

     else if(card(11:18).eq.'COMMENT=') then
       icmt=icmt+1
       call newcmt(charcmt,icmt,card,ncards)

     else
       print *, 'Unrecognized line in a charact. block =', ncards
       call badcard (card,sample)
       print *, 'Trying to continue...'
     endif
     goto 100

990  call badcard (card,sample)
     call carderror(ncards)
     print*, 'Trying to continue ...'
     goto 100
900  return
end

```





```

        icmt=icmt+1
        call newcmt(chemcmt,icmt,card,ncards)

    else
        print *, 'Unrecognized line in a chemistry block =',ncards
        call badcard (card,sample)
        print *, 'Trying to continue...'
    endif

    goto 100

990  call badcard (card,sample)
    call carderror(ncards)
    print*, 'Trying to continue ...'

    goto 100

900  return

end

```

```

c
subroutine readform
  Reads cards in a formation block.

  include 'var.inc/nolist'
  include 'common.inc/nolist'

  icmt=0

100  read(21,'(a80)',end=900) card
     ncards=ncards+1

     if(card.eq.blankcard) then
       continue

     else if(card(1:10).ne.'          ') then
       print *, 'Formation block not properly ended before line',ncards
       call badcard (card,sample)
       print*, 'Trying to continue ...'
       return

     else if(card(11:21).eq.'COMMENT=$$$') then
       return

     else if(card(11:27).eq.'FORMATION SYMBOL=') then
       formsymb=card(28:47)

     else if(card(11:25).eq.'FORMATION NAME=') then
       formname=card(26:65)

     else if(card(11:18).eq.'COMMENT=') then
       icmt=icmt+1
       call newcmt(formcmt,icmt,card,ncards)

     else
       print *, 'Unrecognized line in a formation block =', ncards
       call badcard (card,sample)
       print *, 'Trying to continue...'
     endif

     goto 100

900  return
end

```

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```

c      subroutine readisot
c      Reads cards in an isotope block.

      include 'var.inc/nolist'
      include 'common.inc/nolist'

      icmt=0
      ianal=0

100    read(21,'(a80)',end=900) card
      ncards=ncards+1

      if(card.eq.blankcard) then
        continue

      else if(card(1:10).ne.'          ') then
        print *, ' Isotope block not properly ended before line',ncards
        call badcard (card,sample)
        print*, 'Trying to continue ...'
        return

      else if(card(11:21).eq.'COMMENT=$$$') then
        return

      else if(card(11:19).eq.'ANALYSIS=') then
        ianal=ianal+1
        if(ianal.gt.nanal) then
          print *, 'Too many analysis lines at line',ncards
          print *, 'Trying to continue...'
        else
          read(card,1010,err=990) alysis(ianal),alyst(ianal),
            &      mineral(ianal)
1010    format(19x,a5,9x,a10,9x,a5)
        endif

      else if(card(11:14).eq.'AGE=') then
1020    read(card,1020,err=990) age(ianal), isotratio(ianal)
        format(14x,a20,7x,a25)

      else if(card(11:18).eq.'COMMENT=') then
        icmt=icmt+1
        call newcmt(isotcmt,icmt,card,ncards)

      else
        print *, 'Unrecognized line in an analysis block =', ncards
        call badcard (card,sample)
        print *, 'Trying to continue...'
      endif
      goto 100
990    call badcard (card,sample)
      call carderror(ncards)
      print*, 'Trying to continue ...'
      goto 100
900    return
      end

```







```

c      subroutine readshaw
c      Reads cards in chemistry block.

      include 'var.inc/nolist'
      include 'common.inc/nolist'

100    read(21,'(a80)',end=900) card
      ncards=ncards+1

      if(card.eq.blankcard) then
        continue

      else if(card(1:12).ne.'          ') then
        print *, ' Analysis 1 block not properly ended before
& line',ncards
        call badcard (card,sample)
        print*, 'Trying to continue ...'
        return

      else if(card(19:21).eq.'$$$') then
        return

      else if(card(13:17).eq.'SiO2=') then
        read(card,1010,err=990) (shawval(i), i=1,5)
1010    format(10x,5(7x,a7))

      else if(card(13:17).eq.' MNO=') then
        read(card,1010,err=990) (shawval(i), i=6,10)

      else if(card(13:17).eq.' P2O5=') then
        read(card,1010,err=990) (shawval(i), i=11,15)

      else if(card(13:17).eq.'   Y=') then
        read(card,1010,err=990) (shawval(i), i=16,20)

      else if(card(13:17).eq.'   TH=') then
        read(card,1010,err=990) (shawval(i), i=21,25)

      else if(card(13:17).eq.'   NI=') then
        read(card,1010,err=990) (shawval(i), i=26,30)

      else
        print *, 'Unrecognized line in Analysis 1 block =' ,ncards
        call badcard (card,sample)
        print *, 'Trying to continue...'
      endif

      goto 100

990    call badcard (card,sample)
      call carderror(ncards)
      print*, 'Trying to continue ...'
      goto 100
900    return
      end

```

```

c
subroutine readxrf
  Reads cards in chemistry block.

  include 'var.inc/nolist'
  include 'common.inc/nolist'

100  read(21,'(a80)',end=900) card
     ncards=ncards+1

     if(card.eq.blankcard) then
       continue

     else if(card(1:12).ne.'          ') then
       print *, ' Analysis 2 block not properly ended before
&   line',ncards
       call badcard (card,sample)
       print*, 'Trying to continue ...'
       return

     else if(card(19:21).eq.'$$$') then
       return

     else if(card(13:17).eq.'SiO2=') then
       read(card,1010,err=990) (xrfval(i), i=1,5)
1010  format(10x,5(7x,a7))

     else if(card(13:17).eq.' MGO=') then
       read(card,1010,err=990) (xrfval(i), i=6,10)

     else if(card(13:17).eq.'  Y=') then
       read(card,1010,err=990) (xrfval(i), i=11,15)

     else if(card(13:17).eq.'  NB=') then
       read(card,1010,err=990) (xrfval(i), i=16,17)

     else
       print *, 'Unrecognized line in Analysis 2 block =',ncards
       call badcard (card,sample)
       print *, 'Trying to continue...'
     endif

     goto 100

990  call badcard (card,sample)
     call carderror(ncards)
     print*, 'Trying to continue ...'

     goto 100

900  return

end

```



```

cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
  subroutine remchar (ch, string1, string2, l2)
c    Removes character from string and returns length of new string

  character*1 ch
  character*(*) string1, string2

  l1=len(string1)
  index=1

  do 10 i=1,l1
  if (string1(i:i).ne.ch) then
    string2(index:index)=string1(i:i)
    index=index+1
  else
    endif
10  continue

  l2=len(string2)
  return
end

```

```

cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
  subroutine rjust(string)
c    Right justifies a string in its allotted length.

  character*(*) string
  character*120 temp
  temp=' '

  l=len(string)

c    Find last non-blank character on right
  do 10 i=l,1,-1
10  if(string(i:i).ne.' ') go to 20

20  ilast=i

  temp(l-ilast+1:l)=string(1:ilast)
  string(1:l)=temp(1:l)

  return
end

```



```

cccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
      subroutine shuffle
c       Sorts a card format input file according to an index.
c       20000 cards max

      parameter (maxdim=20000)
      character*80 card(maxdim), blankcard

      do 5 i=1,80
5        blankcard(i:i)=' '

      call getfile(21,'*Give index file: ','in','formatted')
      call getfile(22,'*Give infile:      ','in','formatted')
      call getfile(23,'*Give outfile:     ','out','formatted')

      n=1
10      read(22,'(a80)',end=50) card(n)
         n=n+1
         if(n.gt.maxdim) then
            print *, 'ERROR...array not dimensioned big enough'
            stop
         endif
         go to 10

50      read(21,'(2i5)',end=900) nfirst,num
         do 60 j=nfirst,nfirst+num-1
            call rdeblank(card(j), card(j), 1)
60      write(23,'(a)') card(j)(1:1)
            go to 50

900    print*, 'Number of cards read=',n
         stop
      end

```