

## APPENDIX 1

### **THE PMAG SOFTWARE PACKAGE**

## 1.1. Getting started

After each chapter in the book, there are a number of practical examples for paleomagnetic problems using real data. The programs referred to in the examples are part of the PMAG package of programs. The distribution described here is **pmag1.7**. The most current version of the PMAG package is available by anonymous ftp at:

ftp://sorcerer.ucsd.edu/pub/pmag.

These programs were designed to operate in a UNIX type environment and we include the source code (in Fortran 77) as well as compiled versions for Linux, Solaris, and Mac OS X operating systems. There are also executables (available for pmag1.4 only) provided for running under the MSDOS prompt on a PC. There are many advantages to working in a UNIX environment so we will start with a brief discussion of how to get along in UNIX in general. There are many excellent books on UNIX and the reader is referred to them for a detailed discussion. What follows are the barest essentials for being able to use the programs in this book.

### 1.1.1. SURVIVAL UNIX

This book assumes that you have an account on a UNIX type machine and know how to log in. If you are using a Macintosh with Mac OS X, just use the terminal window. In this book, what *you* type is printed in boldface. At the end of every line, you must also type a carriage return (“Return” or “Enter” on most computer keyboards). To end input into a program, press the control key and “d” < *control* – *D* > simultaneously.

After a login message specific to your computer, you will get a command line prompt which varies widely. In this book, we use the symbol “%” as the command line prompt.

### 1.1.2. THE UNIX FILE STRUCTURE

Fundamental to the UNIX operating system is the concept of directories and files. On windows-based operating systems, directories are depicted as “folders” and moving about is accomplished by clicking on the different icons. In UNIX, the directories have names and are arranged in a hierarchical sequence with the top directory being the “root” directory, known as “/” (see Figure A.1. Within the “/” directory, there are subdirectories (e.g. **usr** and **home**). In any directory, there can also be “files” (e.g. *ex1.1*, *ex1.2* in the figure). Files can be “readable”, “writable” and/or “executable”.

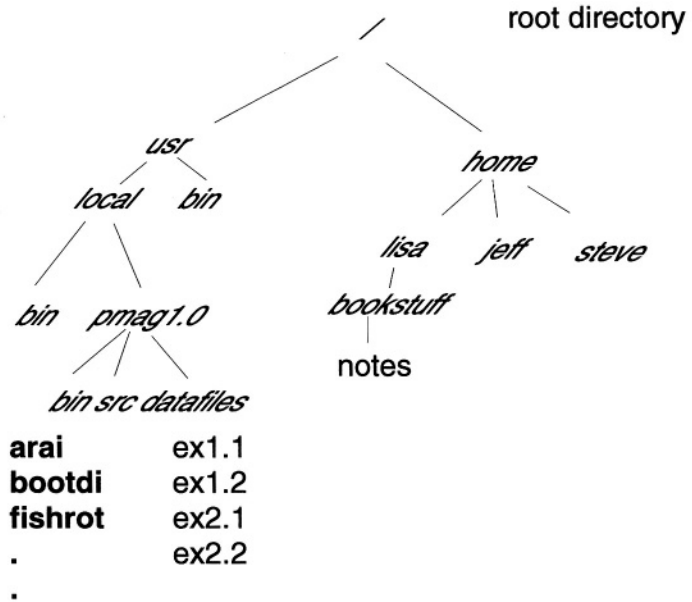


Figure A.1. Part of a UNIX filesystem tree. Directories are shown in italics, ascii files in plain text and executable programs in boldface.

When you log in you enter the computer in your “home” directory. To refer to directories, UNIX relies on what is called a “pathname”. Every object has an “absolute” pathname which is valid from anywhere on the computer. The absolute pathname always begins from the root directory */*. So the absolute pathname to the home directory *lisa* in Figure A.1 is */home/lisa*. Similarly, the absolute pathname to the directory containing *pmag1.7* executables is */usr/local/pmag1.7/bin*. There is also a “relative” pathname, which is in reference to the current directory. If user “*lisa*” is sitting in her home directory, the relative pathname for the file **notes** in the directory **bookstuff** would be **bookstuff/notes**. When using relative pathnames, it is useful to remember that *./* refers to the current directory and *../* refers to the directory “above”.

Commands typed at the command line prompt are handled by a program called the “shell”. There are many different sorts of shells (e.g. *sh*, *csh*, *jsh*, *ksh*, *bash*, *tsh*) that have a different look and feel but they all perform the task of interpreting between the user and the “kernel” which is the actual UNIX operating system. In order to execute a command, the shell needs to know where the command is. There are several “built-in” commands, but most are programs that are either part of the operating system, or something someone wrote (like the ones referred to here). There

are any number of places where programs are kept, so the shell looks in particular places determined by your “path” environment variable. To instruct the shell to look in directories other than the default directories (for example in `/usr/local/pmag1.7/bin`), ask your system administrator to add this directory to your “path”. Otherwise, you can always type the absolute pathname for any program (e.g. `/usr/local/pmag1.7/bin/fishrot`) to execute the program **fishrot**.

### 1.1.3. REDIRECTING INPUT AND OUTPUT

Most UNIX programs print output to the screen and read input from the keyboard. This is known as “standard input and output” or “standard I/O” in the following. One of the nicest things about UNIX is the ability to redirect input and output. For example, instead of typing input to a program with the keyboard, it can be read from a file using the symbol `<`. Output can either be put into a file using the symbol `>`, appended to the end of a file with `>>` or used as input to another program with the UNIX pipe facility (`|`).

### 1.1.4. WILDCARDS

UNIX has the ability to refer to a number of files and/or directories using “wildcards”. The wildcard for a single character is “?” and for any number of characters is “\*”. For example, to refer to all the files beginning with “ex” in the directory:

`/usr/local/pmag1.7/datafiles,`

we use:

`/usr/local/pmag1.7/datafiles/ex*`

To refer only to those from Chapter 1, we use:

`/usr/local/pmag1.7/datafiles/ex1.*` .

### 1.1.5. UNIX COMMANDS

Now we briefly describe essential UNIX commands.

#### • **awk**

Usage: `awk [options] [file(s)] [Standard I/O]`

Description: There are whole books on this program. We will use **awk** in a very primitive way to select specific columns from standard input for use as input into another program. For example, if the third and fourth column of file **myfile** is desired as input to program **myprog** type:

```
% awk '{print $3, $4 }' myfile | myprog
```

#### • **cat**

Usage: `cat [options][file(s)] [Standard I/O]`

Description: **Concatinates** and displays files. It reads from standard input or from the specified file(s) and displays them to standard output.

- **cd**

Usage: cd [directory]

Description: Changes directory from current directory to the one specified.

- **cp**

Usage: cp [file1] [file2]

Description: Copies files or directories.

- **grep**

Usage: grep [options] [expression] [file(s)]

Description: **grep**, like **awk**, is a very versatile (and complicated) program about which one could read an entire book. However, we will use **grep** simply to pick out particular key strings from a file. For example, if a file **myfile** contains lines of data for many samples, and we wish to consider the data for a single sample **mysamp**, lines containing the word **mysamp** can be “gripped” out by:

```
% grep “mysamp” myfile
```

and listed to the screen, redirected to a file, or piped to a program.

- **join**

Usage: join [options] file1 file2

Description: *file1* and *file2* share a common “join” field, by default the first column. This could for example be a sample name. The output file prints the join field, followed by the rest of the line from **file1**, then the rest of the line in **file2**. Say **file1** has magnetometer output data, with lines of data containing: sample, treatment, *D*, *I*, *M* and **file2** has pertinent information such as sample orientation, with lines: sample, azimuth, plunge. We may wish to attach the sample orientation data to the magnetometer output for further processing. **join** allows us to do this by the following:

```
% join file1 file2
```

- **ls**

Usage: ls [options] [directory name]

Description: Lists the contents of the specified directory. If none specified, lists the current directory.

- **man**

Usage: man [command name]

Description: Prints the on-line documentation for the specified command.

- **mkdir**

Usage: mkdir [directory name]

Description: Makes a directory with the specified name.

- **more**

Usage: more [file name]

Description: Displays the contents of a text file on the terminal, one screenful at a time. To view line by line, type RETURN. To view page by page, use the space bar.

- **mv**

Usage: mv [file1 file2]

Description: Renames *file1* to *file2*. This also works for directory names.

- **paste**

Usage: paste [options] [file1 file2]

Description: “Pastes” lines from *file2* onto the corresponding line in *file1*.

- **pwd**

Usage: pwd

Description: Prints the absolute pathname of the working (or current) directory.

- **rm**

Usage: rm [options] [file(s)]

Description: Deletes the specified file or files.

- **rmdir**

Usage: rmdir [options] [file(s)]

Description: Deletes the specified directories. Note: the directory has to be empty.

- **tee**

Usage: tee [file name]

Description: Makes a copy of the standard input to the specified file, then passes it to standard output.

More detailed descriptions are usually available on-line with the **man** command. For example, to find out more about **cat**, type:

% **man cat**

and read all about it.

### 1.1.6. TEXT EDITORS

Text editing is a blessing and a curse in most UNIX systems. You either love it or hate it and in the beginning, you will certainly hate it. There are many ways of editing text and the subject is beyond the scope of this book. Almost all UNIX systems have some flavor of **vi** so try reading the man pages for that.

## 1.2. Getting the programs used in this book

The PMAG software distribution contains the fortran 77 source code, compiled programs for the Solaris, Linux and Mac OS X operating systems as well as executables for MSDOS (only the pmag1.4 distribution for MSDOS).

To run the compiled programs, simply “untar” them using for example the command

```
%tar xvf macosx.tar
```

Put the resulting *macosx* directory in your path or move the files in that directory to a directory that is in your path already.

We recommend that you “untar” the file *pmag1.7.tar*. This contains the source code and miscellaneous files for use with the the exercises in this book. Follow these instructions:

1) Place **pmag1.7.tar** in any directory (PARENTDIR) that you have write permission for.

Your fortran compiler must be known as “f77” or change the make.all file for the name of your compiler (eg., fort77).

2) Execute the following commands:

```
%tar xf pmag1.7.tar
```

```
%cd pmag1.7/src
```

3) Type:

```
%source make.all
```

Make sure **PARENTDIR/pmag1.7/bin** is in your path (or copy them to a directory that is in your path) and try out the programs as described in the book. The data files mentioned in the excercises can be found in **PARENTDIR/pmag1.7/datafiles** with names is given in the book.

### 1.3. Other necessary software

Included in this package of programs is a program called **plotxy**, written by Robert L. Parker and Loren Shure. This is freeware and is supplied as is. For a detailed description on how to use it, print out the file:

**pmag1.7/src/Plotxy/plotxy.doc.**

The program **plotxy** reads commands from standard input and creates a postscript file named by default *mypost*. Finally, you will need to have some sort of Postscript viewer, e.g. **pageview**, **ghostscript**, **ghostview** or commercial illustration product OR just convert the postscript files to pdf format and view them with Acrobat Reader available for free at this website:

<http://www.adobe.com>

### 1.4. Description of programs in excercises

The software described here is updated occasionally. For the description of the program you have, type:

```
% program_name -h
```

for a help message.

Here follows are hints for the use of all the programs referred to in the exercises in this book as well as some supplemental programs that are helpful.

The programs described here are part of the package **pmag1.7**. They are designed to take advantage of the power of UNIX in that they all take command line arguments. The common features are:

1) All programs respond with a help message if the **[-h]** switch is used:

```
% program_name -h
```

2) Programs that produce pictures use the freeware program **plotxy** (see section on getting and installing **plotxy**). PMAG programs output **plotxy** commands which can be piped directly to **plotxy**:

```
% program_name | plotxy
```

This causes **plotxy** to create a postscript file, by default *mypost* which can be viewed with a postscript viewer such as **pageview**, **ghostscript** or **ghostview**, or it can be printed to a postscript compatible printer. It can also be converted to *pdf* format. The **plotxy** commands generated by the PMAG programs can also be saved to a file and modified as needed:

```
% program_name > myfile.com
```

A text editor can be used to modify the commands to taste (see **plotxy** documentation **plotxy.doc** in the Plotxy directory:

```
/usr/local/pmag1.7/Plotxy),
```

or online at:

```
http://sorcerer.ucsd.edu/pmag/plotxy.html.
```

3) Programs read from standard input and write to standard output unless otherwise noted. All input files are expected to be space delimited - not tab or comma delimited. The typical syntax will be:

```
% program_name < input_file > output_file
```

The output from one program can be piped as input to another program. Here follows a brief manual for using the programs in the PMAG package. Refer to the examples at the end of each Chapter for further hints about how to use them.

#### • **arai**

Usage: **arai [-sfmd] [min max field] [Standard I/O]**

Description: makes an Arai plot from input data.

Example 3.5

Options:

- s sets fit from [min] to [max]
- f sets lab field to [field] (in Tesla)
- m uses .mag file as input

Defaults:

finds “optimum” interval - beware! You may have to “tweak” the [min] and [max] values using the [-s] switch



default laboratory field is  $40 \mu\text{T}$ .

Input options:

1em Default input:

Sample treatment intensity  $D I$

.mag file [-m] option:

Sample treatment CSD intensity  $D I$

treatment steps are coded as follows:

XXX.YY where XXX is the temperature and

YY is as follows:

NRMdata: .00

pTRM: .11

pTRM check: .12

Output: plotxy commands.

### • bootams

Usage: bootams [-pP] [Standard I/O]

Description: calculates bootstrap eigenparameters from input file.

Example: 5.9

Options:

-p option specifies parametric (by sample) bootstrap

-P option specifies parametric (by site) bootstrap

Input:  $s_1 s_2 s_3 s_4 s_5 s_6 \sigma$

Output: bootstrap error statistics:

$\tau_1 \sigma D I \eta D_\eta I_\eta \zeta D_\zeta I_\zeta$

$\tau_2 \sigma D I \eta D_\eta I_\eta \zeta D_\zeta I_\zeta$

$\tau_3 \sigma D I \eta D_\eta I_\eta \zeta D_\zeta I_\zeta$

### • bootdi

Usage: bootdi [-pPv] [Standard I/O]

Description: calculates bootstrap statistics for input file.

Example: 4.9

Options:

-p option selects parametric bootstrap

-P works on principal eigenvectors

-v spits out bootstrapped means

Defaults:

simple bootstrap

works on Fisher means

Input:  $D I [\kappa N]$

Output:

Fisher statistics if Fisherian, otherwise bootstrap ellipses for one or two modes:

Mode  $\eta, D_\eta I_\eta \zeta D_\zeta I_\zeta$ .

or if -v selected, bootstrapped eigenvectors  $DI$

• **bootstrap**

Usage: bootstrap [-pb] [Nb] [Standard I/O]

Description: Calculates bootstrap statistics from input data.

Example: 4.8

Options:

-p plot a histogram

-b sets number of bootstraps [Nb](< 10000)

Input: single column of numbers

Output:

if no plot, then:

NNbmeanbounds\_containing\_95%\_of\_means

if plot, then output is series of plotxy commands

Defaults:

no plot and nb=1000

• **cart\_dir**

Usage: cart\_dir [Standard I/O]

Description: Converts cartesian data to geomagnetic elements

Example: 1.2

Input:  $x_1$   $x_2$   $x_3$

Output:  $D$   $I$  magnitude

• **cart\_hist**

Usage: cart\_hist [-dcbpr][dec inc][file1 file2] [Standard I/O]

Description: Makes histograms of cartesian coordinates of input.

Example: 4.11

Options:

-d compares with direction [dec inc]

-c compares two files [file1 file2]

-b plots confidence bounds

-p specifies parametric bootstrap

-r flips second mode for reversals test

Input:  $D$   $I$  [ $\kappa$   $N$ ]

Output is plotxy command file

Defaults:

standard input of single file

no confidence bounds

simple bootstrap

no reversals test

• **curie**

Usage: curie -[lspt] [smooth] [low hi step] [Tmin Tmax] [Standard I/O]

Description: Analyzes Curie temperature data.

Example: 2.2

Options:

-l smooth over [smooth] data points  
 NB: [smooth] must be an odd number  $\geq 3$   
 -s scan range of smoothing intervals  
 [low] to [hi] using a spacing of [step]  
 [low],[hi] and [step] must be odd  
 -p plot option on to generate Plotxy command file  
 can be piped directly to plotxy and viewed:  
 curie -p < filename | plotxy; ghostview mypost  
 printed:  
 curie -p < filename | plotxy; lpr mypost  
 or saved to a file for modification:  
 curie -p < filename > eqarea.com  
 -t truncates to interval between [Tmin] and [Tmax]  
 input:  
 temperature,magnetization

Defaults:

no smoothing  
 plot option off  
 uses entire record

● **di\_geo**

Usage: di\_geo [Standard I/O]

Description: Rotates directions from specimen to geographic coordinates.

Example: 3.2

Input: *D I* azimuth plunge

Output: *D I* (in geographic coordinates)

Notes: the azimuth and plunge are the declination and inclination of the arrow used for reference during the measurements.

● **dir\_cart**

Usage: dir\_cart [-m] [Standard I/O]

Description: Converts geomagnetic elements to cartesian coordinates.

Example: 1.1

Option: -m read magnitude field

Input: *D I* [magnitude]

Output:  $x_1 x_2 x_3$

Notes: VGP longitude, latitude can be substituted for *D, I*.

● **di\_tilt**

Usage: di\_tilt [Standard I/O]

Description: rotates directions from geographic to tilt adjusted coordinates.

Example: 3.2

Input: *D I* strike dip

Output: *D I* (in adjusted coordinates)

Notes: convention is that dip is to the “right” of strike.

- **di\_vgp**

Usage: di\_vgp [Standard I/O]

Description: Transforms declination/inclination to VGP.

Example: 1.5

Input:  $D I$  lat.(° N) long.(° E)

Output: Pole\_Longitude Pole\_Latitude

Notes: convention is positive: North, negative: South and positive: East, negative: West

- **eigs\_s**

Usage: eigs\_s [Standard I/O]

Description: Converts eigenparameters to .s format.

Example: 5.2

Input:

$\tau_3 D\mathbf{V}_3 I\mathbf{V}_3 \tau_2 D\mathbf{V}_2 I\mathbf{V}_2 \tau_1 D\mathbf{V}_1 I\mathbf{V}_1$

Output:  $s_1 s_2 s_3 s_4 s_5 s_6$

Notes:  $\tau_1$  is the largest eigenvalue and  $\mathbf{V}_i$  are the associated eigenvectors.

- **eqarea**

Usage: eqarea [Standard I/O]

Description: Makes an equal area projection of input data.

Example: 1.3

Input:  $D I$

Output: plotxy commands

- **fishdmag**

Usage: fishdmag [-fdm] [beg end ta] [Standard I/O]

Description: Calculates Fisher mean from specified portion of demagnetization data.

Example: 4.3

Options:

-f Fisher mean from [beg] to [end] steps

-d uses .dat file as input

if [ta] = 0 (default), uses geographic ( $D_g, I_g$ )

if [ta] = 1 uses tilt adjusted ( $D_{ta}, I_{ta}$ )

-m uses .mag file as input

where [beg] and [end] are the number of the treatment (i.e. 1st, 2nd, 3rd).

Input options:

Default input:

Sample treatment intensity  $D I$

.mag file [-m] option:

Sample treatment CSD intensity  $D I$

.dat file [-d] option

Sample position treatment CSD intensity  $D_g I_g D_{ta} I_{ta}$

Output: Sample f n beg end  $\alpha_{95} \bar{D} \bar{I}$

• **fisher**

Usage: fisher -kns [kappa] [N] [seed] [Standard I/O]

Description: generates set of Fisher distributed data from specified distribution.

Example: 4.1

Options:

-k specifies  $\kappa$  as [kappa]

-n specifies number as [N]

-s specifies random seed (non-zero integer) as [seed]

Defaults:

$\kappa = 30$

$N = 100$

seed = 1200

• **fishqq**

Usage: fishqq [Standard I/O]

Description: plots Q-Q diagram for input against Fisher distribution.

Example: 4.5

Input: *DI*

Output: plotxy commands

• **fishrot**

Usage: fishrot -kndis [kappa] [N][dec][inc] [seed] [Standard I/O]

Description: draws a Fisher distribution with mean of [dec] [inc] and [kappa], [N], using random seed [seed].

Example: 4.4

Options:

-k specifies  $\kappa$  as [kappa]

-n specifies number as [N]

-d specifies *D* as [dec]

-i specifies *I* as [inc]

-s specifies [seed] for random number generator (non-zero)

Defaults:

$\kappa = 30$

$N = 100$

$D = 0$

$I = 90$

• **foldtest**

Usage: foldtest [-p] [Standard I/O]

Description: Performs bootstrap fold test.

Example: 4.13

Options:

-p option selects parametric bootstrap

Input: *D I* strike dip [ $\kappa$  N]

Output: plotxy commands

Notes: unfolding curve of data is solid line, pseudo samples are dashed histogram is fraction of  $\tau_1$  maxima. Also, dip is to the “right” of strike.

● **gauss**

Usage: gauss -msni [mean] [sigma] [N] [seed] [Standard I/O]

Description: draws a set of Gaussian distributed data from specified distribution.

Example: 4.6

Options:

- m sets the mean to [mean]
- s sets the standard deviation to [sigma]
- n sets the number of points to [N]
- i sets integer random seed to [seed]

Defaults:

- [mean] is 1
- [sigma] is .5
- [N] is 100

● **gofish**

Usage: gofish [Standard I/O]

Description: calculates Fisher statistics from input file.

Example: 4.2

Input:  $D I$

Output:  $\bar{D} \bar{I} N R k \alpha_{95}$

● **goprinc**

Usage: goprinc [Standard I/O]

Description: calculates principal component from input data.

Example: 4.2

Input:  $D I$

Output:  $D I N \tau_1$

● **gtcirc**

Usage: gtcirc [-gdm] [beg end ta] [Standard I/O]

Description: calculates best-fit plane through specified input data.

Example: 3.3

Options:

- g best-fit great circle (plane) from [beg] to [end] steps  
[beg] and [end] are the numbers of the treatment step.  
For example the kNRM step is [1], the second step is [2], etc.
- d uses .dat file as input  
if [ta] = 0 (default), uses geographic ( $D_g, I_g$ )  
if [ta] = 1 uses tilt adjusted ( $D_{ta}, I_{ta}$ )
- m uses .mag file as input

Input options:

Default input:

Sample treatment intensity  $D I$

.mag file [-m] option:

Sample treatment CSD intensity  $D I$

.dat file [-d] option

Sample position treatment CSD intensity  $D_g I_g D_{ta} I_{ta}$  Output:

Sample  $g$   $N$  beg end MAD  $D I$

where  $D$  and  $I$  are for the pole to the best-fit plane

and [beg] and [end] are the numbers of the treatment step.

### • **histplot**

Usage: histplot [-lb] [bin] [Standard I/O]

Description: creates a histogram of input data.

Example: 4.6

Options:

-l plots the distributions of logs

-b sets bin size to [bin]

Input: single column of data

Output: Plotxy commands

Defaults:

not logs

auto binning

### • **hystcrunch**

Usage: hystcrunch [-mpatl] [trunc\_deg] [label] [Standard I/O]

Description: plots hysteresis loop data and massages them.

Example: 2.3

Options:

-m Micromag data file

-p do not plot

-a do not adjust slope

-t truncate to trunc\_deg harmonics

-l label plot with label

Defaults:

-xy data file

-retain 99 terms of FFT

-adjust for high field slope

-no plot label

-generate plotxy commands

### • **igrf**

Usage: igrf [Standard I/O]

Description: calculates reference field vector at specified location and time uses appropriate IGRF or DGRF for date > 1945.

Example: 1.4

Input: year altitude latitude longitude

where year is decimal year, altitude is in kilometers, latitude is in ° N and longitude is in ° E.

Output:  $D I B$  (nT)

• **incfish**

Usage: incfish [Standard I/O]

Description: calculates an estimated inclination, assuming a Fisher distribution, for data with only inclinations. Uses the method of McFadden and Reid [1982].

Example: 4.4

Input: inclinations

Output:  $\langle I \rangle$ , upper and lower bounds,  $N$ , estimated  $\kappa$  and  $\alpha_{95}$ .

• **jackstrat**

Usage: jackstrat Standard I/O

Description: calculates magnetostratigraphic jackknife parameter  $J$ .

Example: 6.1

Input: VGP latitudes or inclinations in stratigraphic order

Output: plotxy commands

• **k15\_hext**

Usage: k15\_hext [-tga] [Standard I/O]

Description: calculates Hext statistics from 15 measurements uses Jelinek's 15 measurement scheme.

Examples: 5.5 & 5.6 .

Options:

- a average whole file
- g geographic coordinates
- t geographic tilt coordinates

Default: average by sample

Input: 1 line with sample name, [azimuth, plunge, strike, dip] followed by 3 rows of 5 measurements for each specimen in the following order (see Chapter 5):

$K_1 K_2 K_3 K_4 K_5$   
 $K_6 K_7 K_8 K_9 K_{10}$   
 $K_{11} K_{12} K_{13} K_{14} K_{15}$

Output: Hext statistics

[if individual samples, id and bulk chi]

$F F_{12} F_{23}$   
 $\tau_1 D I \epsilon_{12} D I \epsilon_{13} D I$   
 $\tau_2 D I \epsilon_{12} D I \epsilon_{13} D I$   
 $\tau_3 D I \epsilon_{12} D I \epsilon_{13} D I$

• **k15\_s**

Usage: k15\_s [-gt] [Standard I/O]



Description: calculates  $\bar{s}$  from 15 measurements scheme ( $K_i$ ) (see Chapter 5).

Example: 5.4

Options:

- g geographic rotation
- t geographic AND tectonic rotation

Input: (see k15\_hext)

Output:  $\bar{s}_1 \bar{s}_2 \bar{s}_3 \bar{s}_4 \bar{s}_5 \bar{s}_6 \sigma$

#### • **lnp**

Usage: lnp [-f] [infile] [Standard I/O]

Description: calculates Fisher mean from combined directed lines and great circles using the method of McFadden and McElhinny [1988].

NB: this program has changed somewhat since the first distribution!

Example: 4.3

Options:

- f calculates mean from data in [infile], one site at a time

Input: output file from **pca**, **gtcirc**, **fishdmag** programs, i.e.

Sample [fpg] n beg end  $\alpha_{95}$  MAD *D I*

sample name convention: ABC123D[1]

where ABC is a study designator of letters (any length)

123 is the site number (any length)

D is a (single) letter for each separately oriented sample

[1] is an optional (single digit) specimen number

Default: assumes whole file is one site.

Output:

site *NL NG*  $\kappa$   $\alpha_{95}$  *D I*

where *NL* is the number of directed lines

where *NG* is the number of great circles

#### • **mag\_dat**

Usage: mag\_dat -n nfofile [Standard I/O]

Description: Converts magnetometer data file to format with geographic and tilt adjusted coordinates. Also pastes in stratigraphic position data.

Uses **.nfo** files made by **mk\_nfo**.

Example: 6.4

Input:

magnetometer data (.mag file format):

sample[optional specimen number] treatment CSD intensity *D I*  
(.nfofile format):

Sample position [cpsdu] NBaz NBpl LABaz LABpl NBstr strike dip

Output:

(.dat file format)

Sample position treatment CSD intensity  $D_g I_g D_{ta} I_{ta}$

where  $D_g$  and  $I_g$  are  $D, I$  in geographic coordinates and  $D_{ta}$  and  $I_{ta}$  are  $D, I$  in tilt adjusted coordinates.

• **mk\_nfo**

Usage: `mk_nfo [-HbBDstfk][strike dip][magdec][lat. long. delta T][cpsdu][az_add az_mult pl_add pl_mult][basename][keyboard input]`

Description: Makes an information file(s) for conversion of data into geographic, tectonic and stratigraphic references.

Example: 6.3

Options:

-H stratigraphic position data

-b structural strike/dip for each sample

-B structural for entire suite read as [strike dip]

-D add [magdec] to all declination/strike info

-s sun compass data using [lat. long. dT]

lat/long of study area

and dT is time difference from GMT

-t type of conversion from

notebook azimuth (= NBaz)/ notebook plunge (= NBpl) to

lab azimuth (= LABaz)/ lab plunge (= LABpl)

[c]ube:

NBaz/NBpl are strike and dip on face perpendicular  
to face with lab arrow

LABaz=NBaz-90, LABpl=NBpl-90

[p]omeroy orientation device:

NBaz/NBpl are direction of drill and angle from vertical

LABaz=NBaz, LABpl=-NBpl

[s]trike/dip:

NBaz/NBpl are strike and dip on face with lab arrow

LABaz=NBaz+90, LABpl=NBpl

[d]rill direction/dip:

NBaz/NBpl are direction of drill (az/pl) in plane perpendicular  
to face with lab arrow

LABaz=NBaz, LABpl=NBpl-90

[u]ser defined conversion

input [az\_add az\_mult pl\_add pl\_mult] to

specify affine for NBaz/NBpl to LABaz/LABpl

e.g. for [p] above, az\_add=0, az\_mult=1

pl\_add=-90, az\_mult=-1

-f output file specified as [basename]

output file will be appended to basename.nfo and if

sun compass data calculated, basename.snfo

-k input from keyboard with prompts

<cntl-D>. to quit.

Input:

Sample NBaz NBpl [position] [strike dip] [yyyymmdd hhmm shadow]

Output:

Sample position type NBaz NBpl LABaz LABpl NBstr strike dip

Defaults:

read/write from Standard I/O

input only: sample, NBaz, NBpl

no declination adjustment

LABaz=NBaz; LABpl=NBpl

### • **pca**

Usage: pca [-pmd] [beg end] [ta] [Standard I/O]

Description: Calculates best-fit line through specified input data.

Example: 3.3

Options:

-p PCA from [beg] to [end] steps

[beg] and [end] are the numbers of the treatment step.

For example the NRM step is [1], the second step is [2], etc.

-d uses .dat file as input

if [ta ] = 0 (default), uses geographic ( $D_g, I_g$ )

if [ta] = 1 uses tilt adjusted ( $D_{ta}, I_{ta}$ )

-m uses .mag file as input

Input options:

Default input:

Sample treatment intensity  $D I$

.mag file [-m] option:

Sample treatment CSD intensity  $D I$

.dat file [-d] option

Sample position treatment CSD intensity  $D_g I_g D_{ta} I_{ta}$

Output:

Sample p N beg end MAD  $D I$

where  $D$  and  $I$  are for the principal component

### • **plotams**

Usage: plotams [-BpPvxjn] [name] [Standard I/O]

Description: Plots AMS data from  $s$  data

Example: 5.10

Options:

-B do not plot simple bootstrap ellipses

-p plot parametric (sample) bootstrap ellipses

-P plot parametric (site) bootstrap ellipses

-v plot bootstrap eigenvectors - not ellipses

-x plot Hext [1963] ellipses

-j plot Jelinek [1978] ellipses

-n use [name] as plot label

Default: plot only the simple bootstrap

Input:  $s_1 s_2 s_3 s_4 s_5 s_6 [\sigma]$

Output: plotxy commands

### ● **plotdi**

Usage: plotdi [-pPv] [Standard I/O]

Description: Makes equal area plot of input data, with uncertainties.

Example: 4.10

Options:

-p parametric bootstrap

-P works on principal eigenvector

-v plots bootstrapped eigenvectors

Defaults:

simple bootstrap

works on Fisher means

plots estimated 95% conf. ellipses

Input:  $D I [k N]$

Output: plotxy commands

### ● **plotdike**

Usage: plotdike [-BpPvn] [name] [Standard I/O]

Description: Makes a plot of ams data for dike margins. Designed for estimating flow directions.

Example 6.5

-B DONT plot simple bootstrap ellipses

-p plot parametric (sample) ellipses

-P plot parametric (site) ellipses

-v plot bootstrap eigenvectors

-n use [name] as plot label

Input:

-one or both files called:

e.s and w.s containing:

$s_1 s_2 s_3 s_4 s_5 s_6 [\sigma]$  for the nominal east and west margins respectively

- a file called dike.dd containing one or more measurements of the dip direction and dip of the dike

-optional files:

struct.dat: contains first and second tectonic corrections as strike and dips

lin.di: contains dec,inc of lineation data

Output: plotxy commands and a file:

fort.20 is summary file

Default: plot only the simple bootstrap confidence ellipses

• **plotdmag**

Usage: plotdmag [-pgfrmd] [beg end] [D] [ta] [Standard I/O]

Description: Makes orthogonal and equal area projections of input demagnetization data.

Example: 3.3

Options:

- p PCA from [beg] to [end] steps
- g gtcirc from [beg] to [end] steps
- f fisher mean from [beg] to [end] steps
- r plot horizontal axis = [D] degrees
- m uses .mag file format
- d uses .dat file format

Defaults:

- North on horizontal ([D] = 0)
- no PCA, gtcircle, or fisher calculations

Input options:

Default input:

Sample treatment intensity  $D I$

.mag file [-m] option:

Sample treatment CSD intensity  $D I$

.dat file [-d] option

Sample position, treatment CSD intensity  $D_g I_g D_{ta} I_{ta}$

Output: plotxy commands

Notes: select either p, f OR g

• **pseudot**

Usage: pseudot [-sm] [min] [ta] [Standard I/O]

Description: Analyses pseudo-Thellier data

Example 3.6

Input options:

- s sets the minimum field to [min]
- m sets input file to .mag format

Input options:

Default input:

Sample tr int dec inc

.mag file option

Sample tr csd int dec inc

treatment steps are coded as follows:

XXX.XY where XXX.X is the AF level and

Y is as follows:

NRM data: 0

ARM: 1

Output: plotxy command file

• **qqplot**

Usage: qqplot [Standard I/O]

Description: Plots data against Normal Quantile

Example: 4.7

Input: single column of data

Output: plotxy commands

Notes: On the plot, there are these parameters:

$N$ : the number of data points

mean: the Gaussian mean

$\sigma$ : the standard deviation

$D$ : the Kolmogorov-Smirnov D-statistic

$D_c$ : the critical value given  $N$  at 95% confidence

(if  $D > D_c$ , distribution is not Gaussian at 95% confidence)

• **s\_eigs**

Usage: s\_eigs [Standard I/O]

Description: Converts  $s$  format data to eigenparameters.

Example: 5.1

Input:  $s_1 s_2 s_3 s_4 s_5 s_6$

Output:

$\tau_1 D\mathbf{V}_1 I\mathbf{V}_1 \tau_2 D\mathbf{V}_2 I\mathbf{V}_2 \tau_3 D\mathbf{V}_3 I\mathbf{V}_3$ .

Notes:  $D\mathbf{V}$ ,  $I\mathbf{V}$  are the directions of the eigenvectors corresponding to the eigenvalues,  $\tau$  and  $\tau_1$  is the largest eigenvalue.

• **s\_flinn**

Usage: s\_flinn [-pl] [Standard I/O]

Description: Plots Flinn diagram of input  $s$  data.

Example: 5.13

Options:

-p parametric bootstrap

-1 plots  $L'$  versus  $F'$

Input:  $s_1 s_2 s_3 s_4 s_5 s_6 [\sigma]$

Output: plotxy commands

• **s\_geo**

Usage: s\_geo [Standard I/O]

Description: Rotates  $s$  data to geographic coordinates

Example: 5.3

Input:  $s_1 s_2 s_3 s_4 s_5 s_6$  azimuth plunge

Output:  $s_1 s_2 s_3 s_4 s_5 s_6$  rotated to geographic coordinates

• **s\_hext**

Usage: s\_hext [Standard I/O]

Description: Calculates Hext statistics from input  $s$  data.

Example: 5.7

Input:  $s_1 s_2 s_3 s_4 s_5 s_6$

Output: Hext statistics

$$F \ F_{12} \ F_{23} \ \sigma$$

$$\tau_1 \ D \ I \ \epsilon_{12} \ D \ I \ \epsilon_{13} \ D \ I$$

$$\tau_2 \ D \ I \ \epsilon_{21} \ D \ I \ \epsilon_{23} \ D \ I$$

$$\tau_3 \ D \ I \ \epsilon_{32} \ D \ I \ \epsilon_{31} \ D \ I$$

● **s\_hist**

Usage: `s_hist (-cbpP123t)[file1 file2] [standard I/O]`

Description: Plots histograms of bootstrapped eigenparameters of  $s$  data.

Example: 5.11

Options:

- c compares [file1] and [file2]
- b plot 95% confidence bounds
- p sample parametric bootstrap
- P Site parametric bootstrap
- 1 plots principal eigenvector
- 2 plots major eigenvector
- 3 plots minor eigenvector

-t plots eigenvalues

Defaults:

- simple bootstrap
- all eigenparameters
- no confidence limits

Input:  $s_1 s_2 s_3 s_4 s_5 s_6 [\sigma]$

Output: plotxy command for histograms of eigenvalues and eigenvectors of bootstrap samples

● **s\_jel78**

Usage: `s_jel78 [Standard I/O]`

Description: Calculates Jelinek (1978) statistics from  $s$  data.

Example: 5.8

Input:  $s_1 s_2 s_3 s_4 s_5 s_6$

Output: Jelinek statistics

$$\tau_1 \ D \ I \ \epsilon_{12} \ D \ I \ \epsilon_{13} \ D \ I$$

$$\tau_2 \ D \ I \ \epsilon_{21} \ D \ I \ \epsilon_{23} \ D \ I$$

$$\tau_3 \ D \ I \ \epsilon_{32} \ D \ I \ \epsilon_{31} \ D \ I$$

● **s\_pt**

Usage: `s_pt [-p] [Standard I/O]`

Description: Makes a Jelinek plot of input  $s$  data.

Example: 5.13

Options: -p parametric bootstrap

Input:  $s_1 s_2 s_3 s_4 s_5 s_6 [\sigma]$

Output: plotxy commands for P' versus T diagram

- **s\_tern**

Usage: s\_tern [-p] [Standard I/O]

Description: Makes a Ternary projection of input *s* data.

Example: See Chapter 5

Options: -p parametric bootstrap

Input: *s*<sub>1</sub> *s*<sub>2</sub> *s*<sub>3</sub> *s*<sub>4</sub> *s*<sub>5</sub> *s*<sub>6</sub> [ $\sigma$ ]

Output: plotxy command file of ternary diagram

Notes: triangles are data, dots are bootstrapped means

- **s\_tilt**

Usage: s\_tilt [Standard I/O]

Description: Rotates *s* data into tilt adjusted coordinates

Example: 5.3

Input: *s*<sub>1</sub> *s*<sub>2</sub> *s*<sub>3</sub> *s*<sub>4</sub> *s*<sub>5</sub> *s*<sub>6</sub> strike dip

Output: *s*<sub>1</sub> *s*<sub>2</sub> *s*<sub>3</sub> *s*<sub>4</sub> *s*<sub>5</sub> *s*<sub>6</sub> [in tilt adjusted coordinates]

- **splint**

Usage: splint [-i] [interval] [Standard I/O]

Description: Calculates spline interpolation of input.

Option: -i uses interpolation interval [interval]

Default: [interval] = 1

Input: x y data with monotonic increasing x

Output: interpolated x,y

- **stats**

Usage: stats [standard I/O]

Description: Calculates Gauss statistics for input data

Example: 4.6

Input: single column of numbers

Output: N mean sum  $\sigma$  (%) stderr 95%conf.

Notes:  $\sigma$  is the standard deviation % is  $\sigma$  as percentage of the mean stderr is the standard error and 95% conf.=  $1.96\sigma/\sqrt{N}$

- **sundec**

Usage: sundec [-u] [delta T] [Standard I/O]

Description: Calculates declination from sun compass measurements.

Example: 3.1

Options: -u sets the time difference [delta T] in hours from universal time (e.g. -5 for EST)

Input: latitude longitude year month day hours minutes shadow\_angle

Output: *D*

Notes: positive: North, East; negative: South, West

- **vgp\_di**

Usage: vgp\_di [Standard I/O]

Description: Transforms VGP to equivalent *D*, *I*

Example: 1.6



Input: pole\_latitude longitude site\_latitude longitude.

Output:  $D I$

Notes: convention is positive: North, negative: South and positive: East, negative: West

• **vspec**

Usage: vspec [Standard I/O]

Description: Calculates a vector average for multiple measurements of a single specimen at a single treatment step.

Example: 6.2

Input:

(.mag file format - see mag\_dat)

Sample treatment CSD intensity  $D I$

Output:

Sample treatment CSD/ $R^*$  intensity  $\bar{D} \bar{I}$

Unique specimen/treatment data are simply copied.

$R^*$  is the vector resultant (including intensity) over the sum of all intensities and ranges from 0 to 1.

## Appendix 2

### **TERMS, ACRONYMS AND CONSTANTS**

Symbol	Term and Definitions
<b>Paleomagnetic acronyms and terms</b>	
$\tau$	Relaxation time: Section 2.6.1; equation 2.18
$\theta_m$	Magnetic co-latitude: Section 1.3.3; equation 1.14
$[a_m]$	Magnetic activity: Section 3.10.2
APWP	Apparent polar wander path: Section 6.3
AF	Alternating field demagnetization: Section 3.5
ARM	Anhysteretic remanent magnetization: Section 2.6.4
CRM	Chemical remanent magnetization: Section 2.6.5
$D$	Declination: Section 1.3.1; equation 1.6
DGRF	Definitive geomagnetic reference field: Section 1.3.3
DRM	Detrital remanent magnetization: Section 2.6.6
GAD	Geocentric axial dipole: Section 1.3
GPTS	Geomagnetic polarity time scale: Section 1.4.2; Section 6.1
IGRF	International geomagnetic reference field: Section 1.3.3
IRM	Isothermal remanent magnetization: Section 2.4.1
$M_{eq}$	Equilibrium magnetization: Section 2.6.2
MD	Multi-domain: Section 2.5
NRM	Natural remanent magnetization: Section 2.6.7
pARM	Partial anhysteretic remanence: Section 2.6.4
pDRM	Post-depositional detrital remanent magnetization: Section 2.6.6
PSD	Pseudo-single domain: Section 2.5
PSV	Paleosecular variation of the geomagnetic field: Section 1.4.1
pTRM	Partial thermal remanence: Section 2.6.3
sIRM	Saturation IRM: See $M_r$
SD	Single domain: Section 2.5
SP	Superparamagnetic: Section 2.6.1
SV	Secular variation: Section 1.4.1
TRM	Thermal remanent magnetization: Section 2.6.3
VADM	Virtual axial dipole moment: Section 1.3.5
VDM	Virtual dipole moment: Section 1.3.5; equation 1.19
VDS	Vector difference sum: Section 3.7
VGP	Virtual geomagnetic pole: Section 1.3.4
VRM	Viscous remanent magnetization: Section 2.6.2
<b>Miscellaneous terms</b>	
GHA	Greenwich hour angle: Section 3.1
SQUID	Superconducting quantum interference device: Section 3.4
UT	Universal time (Greenwich mean time): Section 3.1

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**Physical Parameters and Constants**


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$\alpha$	Co-inclination: Section 4.5
$\chi$	Magnetic susceptibility: The slope relating induced magnetization to an applied field: Section 1.1
$\chi_b$	Bulk magnetic susceptibility: Section 5.3; equation 5.23
$\chi_d$	Diamagnetic susceptibility: Section 2.1.1
$\chi_f$	Ferromagnetic susceptibility: Section 2.2; equation 2.11
$\chi_p$	Paramagnetic susceptibility: Section 2.1.2; equation 2.9
$\Delta M$ curve	Curve defined by subtracting the ascending from the descending curves in a hysteresis loop: Section 2.4.1
$\lambda, \phi$	Latitude, Longitude
$\mu_0$	Permeability of free space: ( $4\pi \times 10^{-7} \text{ Hm}^{-1}$ ): Section 1.1.3
$\theta$	Co-latitude: Section 1.3
$\Theta$	Curie temperature. Section 2.2
$a_{ij}$	Direction cosines: Section 3.2
$a$	The radius of the Earth ( $6.371 \times 10^6 \text{ m}$ ): Section 1.3.3
AMS	Anisotropy of magnetic susceptibility: Section 5.1
<b>B</b>	Magnetic induction: Section 1.1.2
$C$	Frequency factor ( $10^{-10} \text{ s}^{-1}$ ): Section 2.6.1
$D$	Declination: Section 1.3.1
$g_m^l, h_m^l$	Gauss coefficients: Section 1.3.3
<b>H</b>	Magnetic field: Section 1.1.1
$H_{cr}$	Coercivity of remanence; field required to reduce saturation IRM to zero: Section 2.4.1
$H_c$	Coercivity; the magnetic field required to change the magnetic moment of a particle from one easy axis to another: Section 2.4
$H_s$	Saturating field; field required to impart $M_s$ : Section 2.4.1
$I$	Inclination: Section 1.3.1; equation 1.6
$k$	Boltzmann's constant ( $1.381 \times 10^{-23} \text{ JK}^{-1}$ ): Section 2.1.1
$K_i$	AMS measurement: Section 5.1
$K_u$	Constant of uniaxial anisotropy energy: Section 2.4
<b>m</b>	Magnetic moment: Section 1.1.3
$m_b$	Bohr magneton ( $9.27 \times 10^{-24} \text{ Am}^2$ ): Section 2.0
<b>M</b>	Magnetization: Section 1.1.3
$M_r$	Saturation remanence (also sIRM): Section 2.4.1
$M_s$	Saturation magnetization; the magnetization measured in the presence of a saturating field: Section 2.4.1
$P_l^m$	Schmidt polynomials: Section 1.3.3
<b>s</b>	Six elements of $\chi_{ij}$ ; $s_1 = \chi_{11}$ , $s_2 = \chi_{22}$ , $s_3 = \chi_{33}$ , $s_4 = \chi_{12}$ , $s_5 = \chi_{23}$ , $s_6 = \chi_{13}$ : Section 5.1; equation 5.4
$T$	Absolute temperature (in kelvin)
$T_b$	Blocking temperature: Section 2.6.3
$v$	Volume
$v_b$	Blocking volume: Section 2.6.5

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**Statistical parameters**


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$\alpha_{95}$	Circle of 95% confidence (Fisher): Section 4.1, equation 4.4
$\delta$	Residual errors for AMS measurements: Section 5.2, equation 5.20
$\epsilon_{ij}$	Semi-angles of Hext uncertainty ellipses: Section 5.3; equation 5.21
$\kappa$	Fisher precision parameter: Section 4.1
$\eta, \zeta$	Semi-angles of bootstrap uncertainty ellipses: Section 4.10; equation 4.16
$\tau, \mathbf{V}$	Eigenvalues and eigenvectors of tensors
$\hat{\kappa}$	Estimate of $\kappa$ : Section 4.1
CSD	Circular standard deviation (Fisher): Section 4.1
$dm$	Uncertainty in the meridian (longitude) of a paleomagnetic pole: Section 4.1; equation 4.5
$dp$	Uncertainty in the parallel (latitude) of a paleomagnetic pole: Section 4.1; equation 4.5
$F, F_{12}, F_{23}$	Significance tests for anisotropy (Hext): Section 5.3; equation 5.22
$J$	Magnetostratigraphic jackknife parameter: Section 6.1.1
LPA	Linear Perturbation Analysis: Section 5.2
MAD	Maximum angular deviation of principal eigenvector (Kirschvink): Section 3.8.2; equation 3.9
$MAD_{plane}$	MAD of the pole to a best-fit plane (Kirschvink): Section 3.8.2; Equation 3.10
$M_u, M_e$	Significance tests for uniform and exponential distributions: Section 4.6
$N$	Number of samples, specimens or sites
$n_f$	Number of degrees of freedom
$R$	Resultant vector length (Fisher): Section 4.2
$R_o$	Critical value of $R$ for non-random distribution (Watson): Section 4.2; equation 4.6
$S_o$	Residual sum of squares of errors (Hext): Section 5.2; equation 5.14
$T$	Orientation tensor: Section 3.8.1; equation 3.6

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