Useful information for Unix

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General comments

Modes

One can access Unix in two modes: via telnet or via X-Windows. The first option results in a fixed-size window, and no graphics can be displayed. The X-Windows standard allows for viewing of Unix windows locally and remotely; color, graphics, resizing and cut an paste are permitted.

Important keystrokes

To exit a program, use one of the following:		
exit	exit a session and many programs	
quit	exit many programs	
ctrl-d	end of file	
ctrl-c	interrupt a program	
ctrl-z	suspend a program	

Basic commands

20010 00111100100	
pwd	list working directory
ls	list files
mkdir <i>d</i>	make directory d
cd d	change to directory d
cat <i>file</i>	list complete <i>file</i>
more file	list <i>file</i> page-by-page
exit	exit the session (also exit many other programs)
file denotes one file, many files, or a wildcard	

Help on all commands is via the "man" command.

man command provide help about command

Different Unix systems define different backspace key. One can select a key by typing: stty erase *ch* set *ch* as erase character (could be backspace or delete)

Extensions

Files can be wildcards or can be many names.

ls	a*c	list files starting with a and ending with c
ls	al a2 a3	list files a1-a3
ls	a[1-3]	as above
ls	a?	list files with names starting with a and one extra
		character
ls	~/reml/*	list all files in directory <i>reml</i> in home (~) directory
ls	[!a]*	list files that do not start with a

The same naming conventions apply to most commands. **Warning**: names in Unix are case sensitive and use / slashes

Options

Most Unix commands have a great number of options. See details in the man pages. For example, the ls command can be used as

ls -l	list files with details
ls -al	list also hidden files

Informational commands

w finger user	check who is active on the computer find name of <i>user</i> and his (her) whereabouts (if provided)	
finger user@computer As above, remotely; many systems don't allow it because of security concerns		
last -10 find	the last 10 users who logged on	
df pres	sent statistics (including free disk space) on all filesystems	
du -k sho	ws space used in current directory and its subdirectories	
whoami show	ws your userid	
hostname show	ws your computer name	
ps show	ws current processes	
ps -ef show	ws all processes	
kill -9 <i>id</i>	kill process with number <i>id</i>	

Copying and moving

ent

rm -r dir remove directory dir

Other popular file commands

head	print first 10 lines of a file
tail	print last 10 lines of a files
WC	count lines in file(s)
grep	text files finds lines in files that contain text
lpr	print

More exotic file commands

sort	sort or merge
cut	cuts specified columns from file(s)
paste	pastes two or more files line by line
join	joins lines of two files based on specified field
sed	batch editor; can insert, delete and substitute lines and
strings	
awk	fast programming language; can be used as one line
	"database" program
join sed strings	joins lines of two files based on specified field batch editor; can insert, delete and substitute lines and fast programming language; can be used as one line

Editors

vi	visual editor; works over telnet; powerful but awkward to
use	
xedit	simple X-window editor
nedit	Public domain X-Window PC-type editor
pico	easy-to-use non-graphical editor

Linking directories and files

ln

link directories; or produce alternate access to a directory. To access directory /nce/brangus/1996/data/ped as brped under the current directory, type

ln -s nce/brangus/1996/data/ped brped

Running jobs in batch

One can execute jobs system in batch by using the batch command. Below, one executes program (or script) prog. Please note that input to that program needs to come from a file, data in this example.

batch
at> prog <data</pre>

at> <ctrl-D> warning: commands will be executed using /bin/csh job 826136598.b at Wed Mar 6 12:23:18 1996

Once the program executed in the batch system completes, its output is sent by mail. Type mail to read and send mail. Type man mail to see mail options.

Progress of program in batch can be monitored by: ps -ef shows list of all processes, their memory and CPU times top shows graphically status of active jobs in the system.

The top command is a useful commands to see what is happening in the computer. It shows a screen like below:

<pre>last pid: 7477; load averages: 0.86, 0.83 14:33:55 33 processes: 31 sleeping, 1 running, 1 on c Cpu states: 0.0% idle, 1.3% user, 57.9% ker Memory: 158M real, 680K free, 184M swap, 282M</pre>	pu mel, 23.2% iowait, 17.5% swap
PID USERNAME PRI NICE SIZE RES STATE T	IME WCPU CPU COMMAND
1775 ignacy -25 19 340M 254M run 73	.9H 12.20% 74.75% jaadomnlarge
7477 ignacy 33 0 2576K 1424K cpu 0	0:00 0.03% 0.20% top
104 root 33 0 1376K 320K sleep 0	:39 0.02% 0.10% in.routed
308 root 33 0 2864K 464K sleep 0	:02 0.00% 0.00% rpc.ttdbserver
213 root 33 0 2264K 552K sleep 0):01 0.00% 0.00% vold
169 root 33 0 1456K 432K sleep 0	0:01 0.00% 0.00% cron
1 root 33 0 1632K 248K sleep 0):01 0.00% 0.00% init
175 root -3 0 1976K 648K sleep 0):00 0.00% 0.00% nscd
120 root -3 0 1776K 216K sleep 0	0:00 0.00% 0.00% kerbd
1770 ignacy 10 2 880K 176K sleep 0	0:00 0.00% 0.00% ax
7256 root 13 0 1464K 632K sleep 0	0:00 0.00% 0.00% in.telnetd
):00 0.00% 0.00% sh
1769 ignacy 13 2 232K OK sleep 0):00 0.00% 0.00% sh
	0:00 0.00% 0.00% keyserv
5887 root 23 0 1816K 616K sleep 0	0:00 0.00% 0.00% inetd

It shows processes' ids, priorities, amount of memory requested and used, CPU time and name. Once top is running, type h for help or q to quit. Top can be used to reduce priority of long-running programs by typing:

r -20 PID

where PID is a process number.

Top lines of the top command show decomposition of the CPU time. Too much time spent in iowait and swap is an indication that programs are too many/large for the memory available.

Unix tools as filters

General syntax

command {options] [filename(s)]

if filename missing, input from standard input output can be redirected (>) or piped (|) to other commands.

Formats of records

field1 field 2 field 3 ... field n $\$

n.p = field n, starting from character p

standard separator - space other separators can be set by options

Commands

cut	cuts characters or fields from records
	cut -c -5, 10-20,25- cuts characters 1-5, 10-20, 25 to end from each record
	cut -f 1,5,7 cuts fields 1, 5 and 7
paste	past f1,f2,fn Pastes records f1,,fn on the same line; space is a
separator	Tastes records 11,,in on the same line, space is a
grep	
	grep -n -i string file lists files containing string; option -i makes matching case-insensitive, -n precedes lines by their line numbers
sort	
	<pre>sort f1 >f2 sort -o f1 f1 sort alphanumerically on all fields in ascending order; /tmp used for intermediate storage</pre>

-C	to each line add count of repeated lines
+1 -2	take only the specified fields in determining the
	uniqueness (format as in sort)

join

joins two files (sorted) based on provided fields. See man pages for description

tr

tr ` ' `0' fl >f. translate all spaces into zeroes

3)

Examples

Find all lines of fortran programs that contain subroutine "adjust"

grep -n adjust *f

Prepare list of sires and number of daughters from file f1; sires in columns 11-18

```
cut -c11-18 f1|sort|uniq -c >f.
```

Other utilities

sed - batch editor: replacement, deletion, and insertion of strings

sed `s/1990/1992/g' f1 >f. replace all 1990 into 1992

tee

list input and at the same time copy it to a file

```
mtdfreml|tee f1
```

script f1

copies all terminal traffic to file f1

script f1
.....
exit (or ctr-d)

AWK

a programming language, similar in ease but more powerful (and cryptic) than BASIC, with capabilities for line (and much more) processing

'program' fl -f program fl				
\$0 \$1 \$2 subs	entire line first field second field str(\$0,5,10)	10 characters of string \$0 beginning with character 5		

NR current record numberNF number of fields in current line

General structure of awk file

BEGIN {action} condition {action} END {action}

> BEGIN & END lines not obligatory if no condition, only {action} performed if no {action}, list entire line

<u>examples</u>

- awk '\$1 > 100' f1 list lines in f1 where field 1 is > 100; conditions specified as in C language
- awk `{print NR,\$0}' precede each record with record number
- awk `{printf("%10s%20s%15s\n",\$1,\$2,\$3}'
 lists fields 1-3 in 10, 20 and 15 character space, respectively;
 formats are described as in C language

Problem 1

Select only first-parity records from file with the following format:

Problem 2

A file contains a sire in column 1-10 and a performance of its daughters in 11-15. List each sire with a consecutive number and a mean performance of its daughters. The file is sorted by sire. List the total number of sires at the end.

awk -f progfile

progfile:

BEGIN	{nsire=0; oldid=""; sum=0} {sire=substr(\$0,1,10); perf=substr(\$0,11,5);
	sum=sum+perf
	if sire != oldid
	{nsire=nsire+1
	<pre>print sire,nsire,sum/nsire}</pre>
END	} {print sire,nsire,sum/nsire; print nsire}

Unix - Part 1

Ways to deal with long names

- file manager (in X)
- cut and paste (in \dot{X} or OS/2)
- wildcards
- command editing

Wildcards

home directory
anything
one character
[ab] one of a or b
[!a] not a
[a-g] one of a to g

Example

ls ?[a-d]*

/home/ignacy ls ?[a-d]* aa abc backcmd* mbox tar.log watorold.c wc.c

Redirection and pipes

mtdfreml <input >output

Terminal will be locked

mtdfreml <input >output&

Terminal won't be locked; better submit long programs by batch

File ff: animal_id sire_id WW

1,000,000 records

Find animal 1234567; don't tie the terminal

grep 1234567 ff&

or

awk '\$2 == 1234567' ff&

Find progeny of 1234567 with highest WW awk '2 = 1234567' ff | sort +2 | head -1

/home/ignacy du | more 52 ./sas 2 ./.wastebasket 2 ./ignacy1 2 ./.cetables 212 ./Http/gifs ./Http 568 ./ab 548 14 ./nsmail ./back 2 1198 ./idled/idled-1.05 1524 ./idled ./inip94/600 2 7212 ./inip94 ./.tt 2 30178 ./temp ./wpfiles/krysia 110 ./wpfiles 3100 ./.mosaic-personal-annotations 4 2 ./tmp 610 ./sasuser ./.spa 6 1114 ./meishan/dfreml/DF93/PRE ./meishan/dfreml/DF93/UNI 2872 ./meishan/dfreml/DF93/MUV 4036 ./meishan/dfreml/DF93/MUW 1712 ./meishan/dfreml/DF93/MUX 122 ./meishan/dfreml/DF93/LIB 400 302 ./meishan/dfreml/DF93/INC ./meishan/dfreml/DF93/SPA 170 ./meishan/dfreml/DF93/bin 4544/ioffice/auxiliary 14 ./ioffice 16 ./mtdfreml 9166 26 ./system ./.helplus 2 215570 . /home/ignacy

Simplifying work

need to repeat the same command or string, e.g.,

/opt1/emacs/bin/emacs largedata

1. Create an alias

alias em /opt1/emacs/bin/emacs

em largedata

(works for programs only)

2. Setting the variable

set a =/opt1/emacs/bin/emacs

\$a largedata

(works for names and programs)

3, Creating a symbolic link on your directory

ln -s /opt1/emacs/bin/emacs . emacs largedata

4. Create a script

Configuration of nce.ads.uga.edu

448 Mbytes of memory3 x 9 Gbyte applcation disks4 x 4 Gbyte user disks2 Gbyte for system

Names of files can be readable

To describe 1996 angus raw data:

angus.96r or angus.1996.raw

angus.1996.results.provenbulls.complete or angusrp.com Long names too long to type. To avoid typing:

use file manager use cut-and-paste specify names with wildcards:

an* an*1996* a*raw [ab][nb]*1996* [!a]*

list files, and then remove the listed files

>ls very-long-name*

very-long-name1 very_long_name2 >^ls^rm

!p !25 history

Vi editor

Full screen editor that works on every Unix machine in a non graphical environment; small subset of options shown below. To start:

vi filefor one filevi *.fall files *.f (:n moves to next one)vi +25 fileset the cursor on line 25

<u>Cursor mode</u>		<u>Type mode</u>
	1	ESC
CTRL-F (go forward) CTRL-B (go backward) 6789 (or) hjkl		
 a (insert after cursor) A (insert at end of line i (insert before cursor) I (insert at beginning of line 	±	
x (delete 1 character) 5x (delete 5 characters)		
dd (delete 1 line) 5dd (delete 1 line)		
U (undelete)		
5yy (copy 5 lines)		
p (insert copied or deleted text after the cursor)		
/text (search forward) ? (search backward) 7G (go to line 7) G (go to last line)		
ZZ (save and exit)		
:q! (quit without saving)		

4. Find who logged more time last week: matt or daniel (last, wc, grep)

/home/ignacy finger matt Login name: matt In real life: Matt Culbertson Directory: /home/matt Shell: /bin/csh On since May 9 11:15:30 on pts/10 from 128.192.43.68:0.0 5 hours 54 minutes Idle Time No unread mail No Plan.

/home/ignacy finger daniel Login name: daniel In real life: Daniel deMattos Directory: /home/daniel Shell: /bin/csh On since May 9 14:55:36 on pts/17 from nce 2 hours 13 minutes Idle Time No unread mail No Plan.

/home/ignacy last matt|grep May|wc
 26 260 1924
/home/ignacy last daniel|grep May|wc
 36 360 2665

Unix tools as filters

General syntax

command {options] [filename(s)]

if filename missing, input from standard input output can be redirected (>) or piped (|) to other commands.

Formats of records

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n.p = field n, starting from character p

standard separator - space other separators can be set by options

Commands

cut	cuts characters or fields from records	
	cut -c -5, 10-20,25- cuts characters 1-5, 10-20, 25 to end from each record	
	cut -f 1,5,7 cuts fields 1, 5 and 7	
paste	paste f1,f2,fn Pastes records f1,,fn on the same line; space is a separator	
grep	grep -n -i string file lists files containing string; option -i makes matching case-insensitive, -n precedes lines by their line numbers (see also egrep and fgrep)	
egrep	egrep '(string1 string2 string3)' file matches any of the strings	
sort		

sort f1 >f2
sort -o f1 f1
sort alphanumerically on all fields in ascending order;
/tmp used for intermediate storage
sort +2 -3 +0 -1
sorting fields:
 skip first 2, until 3 (resulting primary sorting field
 3)

Skip first 0 until 1 (resulting secondary sorting field 1)

sort -T./. +.015 -0.18 +0.5 -0.7 sort on characters 16-18 and 6-7; intermediate directory is the current directory

other options

-r reverse order

-n numeric sort

sort f1 f2 f3 sort&merge uniq

options

-c	to each line add count of repeated lines
+1 -2	take only the specified fields in determining

+1 -2	take only the specified fields in determining the		
	uniqueness (format as in sort)		

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joins two files (sorted) based on provided fields. See man pages for description

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tr ' ' '0' f1 >f. translate all spaces into zeroes

Examples

Find all lines of fortran programs that contain subroutine "adjust"

grep -n adjust *f

Prepare list of sires and number of daughters from file f1; sires in columns 11-18

```
cut -c11-18 f1 | sort | uniq -c \geq f2
```

Other utilities

sed - batch editor: replacement, deletion, and insertion of strings

sed 's/1990/1992/g' f1 >f2 replace all 1990 into 1992

tee - list input and at the same time copy it to a file

mtdfreml|tee f1

script - copies all terminal traffic to file

script f1 exit (or ctr-d) AWK

a programming language, similar in ease but more powerful (and cryptic) than BASIC, with capabilities for line (and much more) processing

awk 'program' f1 awk -f program f1

\$0 entire line
\$1 first field
\$2 second field
\$ubstr(\$0,5,10) 10 characters of string \$0 beginning with character 5

NR current record numberNF number of fields in current line

General structure of awk file

BEGIN {action} condition {action} END {action}

> BEGIN & END lines not obligatory if no condition, only {action} performed if no {action}, list entire line

examples

- awk '\$1 > 100 && \$2=="May" f1 list lines in f1 where field 1 is > 100 and \$2 is May; conditions specified as in the C language
 - == equal != not equal && and || or
- awk '{print NR,\$0}' precede each record with record number
- awk '{print log(\$1), \$2+\$3}' write functions of fields in the record
- awk '{printf("%10s%20s%15s\n",\$1,\$2,\$3}'
 lists fields 1-3 in 10, 20 and 15 character space, respectively;
 formats are described as in C language

Problem 1

Select only first-parity records from file with the following format: 1-38 fixed fields

- 39 starting parity numbers
- 40 number of records in line
- 41-47 first record
- 48-.. later records

awk 'substr(\$0,39,1)==1 {printf(``%39s%1s%7s'',substr(\$0,39,39),1,substr(\$0,41,7))}'

Problem 2

data

1-10 sire no 11-15 WW

sorted by sire

1. List each sire with a consecutive number and a mean performance of its daughters.

2. List the total number of sires at the end.

awk -f progfile datafile

progfile:

```
BEGIN {nsire=0; oldid=""; sum=0
{sire=substr($0,1,10);
perf=substr($0,11,5);
sum=sum+perf
if sire != oldid
{nsire=nsire+1
print sire,nsire,sum/nsire}
}
END {print nsire}
```

Homework

Select a data file with at least 10,000 but no more than 1,000,000 records. Time each command

For the homework, please use the data set /scr/ignacy1/example_file. This file contains 100,000 records, and the sire field is in columns 21-28.

Then:

- 1. Identify one sire; Find how many records contain that sire id (head, grep, wc)
- 2. Create a file that only contains sire ids (cut or awk)
- 3. Sort that file (sort)

4. Find number of progenies per each sire; find the sire with the largest number of progeny (uniq, sort, head)

Put the name of the file in variable \$a so that it does not have to be retyped

```
/home/ignacy set a = /scr/ignacyl/example_file
/home/ignacy echo $a
/scr/ignacyl/example file
```

Get id of the first sire

/home/ignacy head -1 \$a|cut -c21-28 41572878

Command set b = `xxx` puts results of command xxx in variable \$b

```
/home/ignacy set b = ` head -1 $a|cut -c21-28`
/home/ignacy echo $b
41572878
```

Search the sire.

```
/home/ignacy time grep $b $a
8007841458800715900441572878....
1.0u 0.0s 0:01 57% 0+0k 0+0io 0pf+0w
Searching took 1.0 second of users time and 0.01 wall clock time, and took 57% of processor's
time.
```

Extract sire ids

```
/home/ignacy time cut -c21-28 $a >a
22.0u 0.0s 0:28 78% 0+0k 0+0io 0pf+0w
/home/ignacy head -5 a
41572878
41428104
41527802
41515207
41457932
```

Sort and check

/home/ignacy time sort -o a a
4.0u 0.0s 0:07 52% 0+0k 0+0io 0pf+0w
/home/ignacy head -5 a

20255163 20255163 20255163 20259668 20260599

Summarize

```
/home/ignacy
/home/ignacy uniq -c a|sort -n -r|head -10
2551 41626813
2026 41667366
1462 41773417
1444 41650414
1253 41723741
1156 41491007
1038 41724657
1013 41810969
881 41672151
867 41635843
```

Sire 41626813 has 2551 progeny in the file.

Unix scripts and shells

Shells are programs that accept user's input. There are 3 popular shells in Unix: sh, csh and ksh. We dealt with csh only, and commands below apply to csh only. There are similar instructions for other shells too.

Variables

Unix variable contains text string.

/home/ignacy set a = /usr/local/bin

```
/home/ignacy echo $a
/usr/local/bin
```

Look for all files usr/local/bin/d*

ls \$a/d*

To check on a variable:

echo \$a

Backquotes around command(s) return text that would otherwise be displayed by the execution of the command(s).

Set \$a to the first file starting in beef

```
set a = `ls beef*|head -1`
```

Commands useful for scripts

Cat The cat command can be used to create text files.

```
/home/ignacy cat <<AA >b
>beef.5.96
>1992
>1994
AA
/home/ignacy cat b
beef.5.96
1992
1994
```

Foreach

Commands are repeated with \$a assigned to consecutive elements of the list

```
foreach a (list)
.....
commands
.....
end
```

foreach h2 (.3 .4 .5)
blub \$h2
end

To change all files ending with .f into .for

```
foreach a (*.f)
mv $a $aor
end
```

If statement

```
if (condition) command
if ($1 >= 235) mtdfreml
if (`ls mtdfreml.output` != `' ) exit
```

```
operators as in awk (==, !=, >=, <=).
```

Script

A unix script is a text file that contains Unix instructions and that has execute permissions. In a simple case, it contains commands that otherwise would be typed on the command line

```
# This scripts runs several evaluation programs
# sequentially
#
blupedit <ed.par
bluprun <run.par
blupaccur <acc.par
blupfinish <fin.par</pre>
```

Instead of having to set up parameter files in advance, one can put them into the script

```
# Parameter file created in the script so that one does
# not have to edit parameter files separately
#
cat <<AA >ed.par
angus.sires
1970
1990
1000
AA
blupedit <ed.par
...
```

If a program needs just one line of input, it can be generated with the echo command:

echo 1000 .001 | bluprun

When one program fails, there is no need to continue

```
# This script exits before running bluprun if file
```

```
# edit.out does not exist
#
blupedit <ed.par
if (`ls edit.out` == `')exit
bluprun <run.par
...</pre>
```

If script is called with parameters, e.g., beefeval par1 par2 par3 then inside the script, \$1 will have value par1, \$2 will have value par2 etc.

One script for many breeds

breedeval angus

Assume that:

1. Breeds are in directory /nce/breed

2. Program flows are identical

Then one can create a more universal script

```
# change to breed directory
cd /nce/$1
# create parameters in ed.par
cat <<AA >ed.par
$1.sires
1970
1990
1000
AA
# Run
blupedit <ed.par
.....</pre>
```