

Introduction to The Command Line II

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A good start

- These commands, and a few more we'll learn today, are enough to get started with using the command line.
- As you have seen, Unix commands are simple, and are designed to do one specific thing.
- By combining these commands together we will be able to do more interesting things.

Our commands

```
echo arg          echo the argument
pwd              present working directory
ls [dir]         list the directory contents
cd [dir]         change directory
history [num]    print the shell history
man cmd          command's man page
cp file1 file2   copy a file
mv file1 file2   move/rename a file
rm file         delete a file
mkdir dir        create a directory
rmdir dir        delete a directory
more file        scroll through file
less file        scroll through file
cat file         print the file contents
cmd > file       redirect output to file
cmd >> file      append output to file
cmd < file       use file as input to cmd
head file        print first 10 lines of file
tail file        print last 10 lines of file
wc file          word count data of file
```

Example problem: Cochlear implant study

To participate, subjects visited the laboratory and one of the lab techs played an audio sample, and recorded their data. Data consists of:

- date of the test.
- information relating to the sounds played and heard during the test.
- subject ID.
- birth month/year.

Each set of test results were written out to a text file, one set per file.



source: wikipedia.org

Copying the data

```
[brelier.IBM-c5724425579] > pwd
```

```
/home/mobaxterm
```

```
[brelier.IBM-c5724425579] > df -h
```

Filesystem	Size	Used	Available	Use%	Mounted on
C:/Users/IBM_AD~1/AppData/Local/Temp/MOBAXT~1.1/bin					
	29.9G	16.8G	13.1G	56%	/usr/bin
C:/Users/IBM_AD~1/AppData/Local/Temp/MOBAXT~1.1/lib					
	29.9G	16.8G	13.1G	56%	/usr/lib
C:/Users/IBM_AD~1/AppData/Local/Temp/MobaXterm7.1					
	29.9G	16.8G	13.1G	56%	/
C:/Users/IBM_AD~1/DOCUME~1/FakeHome					
	29.9G	16.8G	13.1G	56%	/home/mobaxterm
C:	29.9G	16.8G	13.1G	56%	/drives/c
E:	7.2G	736.2M	6.5G	10%	/drives/e

```
[brelier.IBM-c5724425579] > cp -r /drives/e/SCMP101_shell .
```

```
[brelier.IBM-c5724425579] > cd SCMP101_shell
```

```
[brelier.IBM-c5724425579] >
```

Copying the data, continued

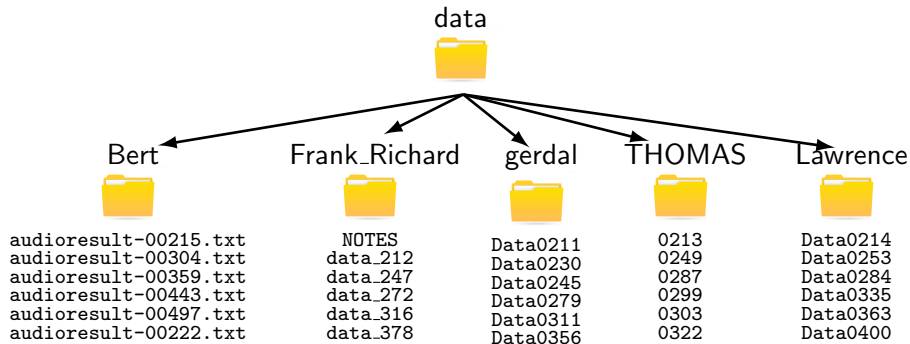
```
[brelier.IBM-c5724425579] > ls
BashSessionData.tar.gz
-----
[brelier.IBM-c5724425579] > tar -z -x -f BashSessionData.tar.gz
-----
[brelier.IBM-c5724425579] >
-----
[brelier.IBM-c5724425579] > ls -F
data/ BashSessionData.tar.gz
```

What happened?

- A 'tar' file (also called a 'tarball') is a file in which has been bundled a number of other files, for easy of moving around.
- `tar` handles a tar file,
 - ▶ `-z` means gunzip it.
 - ▶ `-x` means extract the contents of the file.
 - ▶ `-f` specifies which file you are applying this command to.

The data is now in the 'data' directory.

Data is a complete mess



Multiple directories, inconsistent file names, some directories have a NOTES file...

Goal: create a single file format, using CLI.

Pipelines of commands

- So far we've used the following technique to combine commands:

```
[brelrier.IBM-c5724425579] > pwd
/home/mobaxterm/Desktop
[brelrier.IBM-c5724425579] > cd data/Lawrence
[brelrier.IBM-c5724425579] > wc Data* > all-wcs
[brelrier.IBM-c5724425579] > more all-wcs
```

which creates a temporary file we don't care about; we just want to scroll through the wc results.

- This combination of actions, output of one command goes straight to another, is so common and useful that the shell has a special feature to do this:

```
[brelrier.IBM-c5724425579] > wc Data* | more
```

- The 'pipe' allows you to chain together small commands.

The sort command

The sort command can take a number of important flags:

- `-n`: sort by number (not lexicographic; `101 < 30` without `-n`).
- `-k [num]`: sort by the k'th column.
- `-r`: reverses order.

```
[brelier.IBM-c5724425579] > sort -n -k 3
all-wcs
...
9 24 150 Data0515
9 24 153 Data0214
468 1246 7644 total
-----
[brelier.IBM-c5724425579] > wc Data* |
sort -n -k 3
...
9 24 150 Data0515
9 24 153 Data0214
```

More commands

<code>wget url</code>	downloads the url
<code>tar file</code>	handles tar files
<code>cmd1 cmd2</code>	pipe cmd1 output to cmd2
<code>sort file</code>	sorts the lines of file
<code>arg</code>	mandatory argument
<code>[arg]</code>	optional argument

Pop quiz!

Modify this code to print only the smallest, and then only the largest, data file.

```
[brelhier.IBM-c5724425579] > wc Data* |  
sort -n -k 3  
...  
9 24 150 Data0515  
9 24 153 Data0214  
468 1246 7644 total
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
arg	mandatory argument
[arg]	optional argument

Pop quiz!

Modify this code to print only the smallest, and then only the largest, data file.

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3  
...  
9 24 150 Data0515  
9 24 153 Data0214  
468 1246 7644 total
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 | head -n 1  
9 24 144 Data0234
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
arg	mandatory argument
[arg]	optional argument

Pop quiz!

Modify this code to print only the smallest, and then only the largest, data file.

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3
```

...

```
9 24 150 Data0515
```

```
9 24 153 Data0214
```

```
468 1246 7644 total
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 | head -n 1
```

```
9 24 144 Data0234
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 -r | head -n 2
```

```
468 1246 7644 total
```

```
9 24 153 Data0214
```

More commands

wget **url** downloads the url

tar **file** handles tar files

cmd1 | **cmd2** pipe cmd1 output to cmd2

sort **file** sorts the lines of file

arg mandatory argument

[**arg**] optional argument

Pop quiz!

Modify this code to print only the smallest, and then only the largest, data file.

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3  
...  
9 24 150 Data0515  
9 24 153 Data0214  
468 1246 7644 total
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 | head -n 1  
9 24 144 Data0234
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 -r | head -n 2  
468 1246 7644 total  
9 24 153 Data0214
```

```
[brelier.IBM-c5724425579] > wc Data* |  
sort -n -k 3 -r | head -n 2 | tail -n 1  
9 24 153 Data0214
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
arg	mandatory argument
[arg]	optional argument

Our first shell script

It's time to write our first shell script. To do this we need to use a text editor. Linux has many text editors, and they're all difficult to use.

In MobaXterm try Tools > MobaTextEditor, or you can use Notepad. Create a file called 'biggest':

```
#!/bin/bash
wc * | sort -n -k 3 | tail -n 2 | head -n 1
```

And run it by typing

```
[brelrier.IBM-c5724425579] > rm all-wcs
[brelrier.IBM-c5724425579] > source biggest
9 24 153 Data0214
```

MobaXterm makes the file executable by default. As such, we can just run it like a regular program:

```
[brelrier.IBM-c5724425579] > ./biggest
9 24 153 Data0214
```

Largest range - grep

Number of characters is probably unimportant for our analysis. What about the data with the largest range?

```
[brelrier.IBM-c5724425579] > grep Range  
Data0352  
Range: 2
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

grep prints the lines from the input that contain the search argument.

Largest range - grep

Number of characters is probably unimportant for our analysis. What about the data with the largest range?

```
[brelhier.IBM-c5724425579] > grep Range  
Data0352  
Range: 2  
-----  
[brelhier.IBM-c5724425579] > grep Range *
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

grep prints the lines from the input that contain the search argument.

Largest range - grep

Number of characters is probably unimportant for our analysis. What about the data with the largest range?

```
[brelier.IBM-c5724425579] > grep Range
Data0352
Range: 2
-----
[brelier.IBM-c5724425579] > grep Range *
...
Data0544:Range: 6
Data0554:Range: 2
-----
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

grep prints the lines from the input that contain the search argument.

Largest range - grep

Number of characters is probably unimportant for our analysis. What about the data with the largest range?

```
[brelier.IBM-c5724425579] > grep Range
Data0352
Range: 2
-----
[brelier.IBM-c5724425579] > grep Range *
...
Data0544:Range: 6
Data0554:Range: 2
-----
[brelier.IBM-c5724425579] > grep -v Range
Data0352
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

grep prints the lines from the input that contain the search argument.

Largest range - grep

Number of characters is probably unimportant for our analysis. What about the data with the largest range?

```
[brelier.IBM-c5724425579] > grep Range
Data0352
Range: 2
-----
[brelier.IBM-c5724425579] > grep Range *
...
Data0544:Range: 6
Data0554:Range: 2
-----
[brelier.IBM-c5724425579] > grep -v Range
Data0352
#
Reported: Mon Jul 25 14:01:36 2011
Subject: babyMcCartney281
Year/month of birth: 1991/07
Sex: M
CI type: 14
Volume: 1
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

grep prints the lines from the input that contain the search argument.

grep -v prints the lines from the input that *don't* contain the search argument.

Pop quiz!

Create a new script, called 'biggestRange', which prints out the data file which has the biggest Range.

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

Pop quiz!

Create a new script, called 'biggestRange', which prints out the data file which has the biggest Range.

```
[brelhier.IBM-c5724425579] > cat biggestRange
#!/bin/bash

# a comment! Use the # sign.

grep Ra * | sort -n -k 2 | tail -n 1
[brelhier.IBM-c5724425579] > ./biggestRange
Data0531:Range: 9
[brelhier.IBM-c5724425579] >
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
arg	mandatory argument
[arg]	optional argument

Cutting up the output

How do keep just part of the output?

Suppose we want just the file name?

```
[brelier.IBM-c5724425579] > grep Ra * |  
sort -n -k 2 | tail -1  
Data0531:Range: 9  
-----  
[brelier.IBM-c5724425579] > grep Ra * |  
sort -n -k 2 | tail -1 | cut -c -8  
Data0531
```

Suppose we just want the range?

```
[brelier.IBM-c5724425579] > grep Ra * |  
sort -n -k 2 | tail -1 | cut -c 10-  
Range: 9
```

Suppose we just want something else?

```
[brelier.IBM-c5724425579] > grep Ra * |  
sort -n -k 2 | tail -1 | cut -c 2-5  
ata0
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
arg	mandatory argument
[arg]	optional argument

- "-c" tells cut to cut characters.
- "-8" means keep up-to-and-including character eight.
- "10-" means keep 10 and higher.

Suppose I need to save information

If I need to save something, I use variables.

```
[brelier.IBM-c5724425579] > grep Ra * |
sort -n -k 2 | tail -1 | cut -c -8
Data0531
-----
[brelier.IBM-c5724425579] > i='grep Ra * |
sort -n -k 2 | tail -1 | cut -c -8'
-----
[brelier.IBM-c5724425579] >
-----
[brelier.IBM-c5724425579] > echo i
i
-----
[brelier.IBM-c5724425579] > echo $i
Data0531
```

NOTE: The ' symbol used here is the single quote in the upper-left corner of your keyboard!! Using this single quote will execute the command.

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
arg	mandatory argument
[arg]	optional argument

What happens if you use the other single quote?

Arguments for bash scripts

- We'd like to use our script on each directory, but not have a copy in each one.
- Move biggestRange down one directory, and change it so that it works on any directory's files:

```
[brelier.IBM-c5724425579] > pwd
/home/mobaxterm/Desktop/data/Lawrence
-----
[brelier.IBM-c5724425579] > mv biggestRange ..
-----
[brelier.IBM-c5724425579] > cd ..
-----
[brelier.IBM-c5724425579] > cat biggestRange
#!/bin/bash
grep Range ${1}/* | sort -n -k 2 | tail -1
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
arg	mandatory argument
[arg]	optional argument

- When a command is run in the shell, its name is put in $\${0}$.
- All other arguments are put in $\${1}$, $\${2}$...

Arguments for bash scripts

```
[brelier.IBM-c5724425579] > pwd
/home/mobaxterm/Desktop/data
-----
[brelier.IBM-c5724425579] > ls
Bert Frank Richard Lawrence THOMAS
alexander biggestRange gerdal jamesm
-----
[brelier.IBM-c5724425579] > ./biggestRange
Bert
Bert/audioresult-00384.txt:Range: 10
-----
[brelier.IBM-c5724425579] > ./biggestRange
THOMAS
THOMAS/0336:Range: 10
-----
[brelier.IBM-c5724425579] > ./biggestRange
james
grep: james/*: No such file or directory
-----
[brelier.IBM-c5724425579] > ./biggestRange
jamesm
jamesm/data_517.txt:Range: 10
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
arg	mandatory argument
[arg]	optional argument

Loops for bash scripts

Bash has loops, just like any language:

```
[brelier.IBM-c5724425579] >  
[brelier.IBM-c5724425579] > cat  
loop_biggestRange #!/bin/bash  
for dir in alexander Bert Frank.Richard  
do  
echo "The biggest range in " ${dir} " is:"  
./biggestRange ${dir}  
done
```

```
[brelier.IBM-c5724425579] >  
[brelier.IBM-c5724425579] >  
./loop_biggestRange  
The biggest range in alexander is:  
alexander/data_462.DATA:Range: 10  
The biggest range in Bert is:  
Bert/audioreresult-00384.txt:Range 10  
The biggest range in Frank.Richard is:  
Frank.Richard/data_538:Range: 10
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
for ..do..done	for loop in bash

arg	mandatory argument
[arg]	optional argument

If statements for bash scripts

Bash has if statements, just like any other language:

```
[brelier.IBM-c5724425579] > pwd /home/-
mobaxterm/Desktop/data
-----
[brelier.IBM-c5724425579] > cat if_test.sh
#!/bin/bash
cd ${1}
for filename in *
do

if [ $filename == "NOTES" ]
then
echo "I'm a NOTES file."
fi

done
-----
[brelier.IBM-c5724425579] > ./if_test.sh
jamesm
I'm a NOTES file.
-----
[brelier.IBM-c5724425579] > ./if_test.sh
alexander
```

More commands

wget url	downloads the url
tar file	handles tar files
cmd1 cmd2	pipe cmd1 output to cmd2
sort file	sorts the lines of file
source file	run the cmds in file
grep arg file	search for arg in file
cut flags output	cut part of output
for.. do .. done	for loop in bash
if.. then .. fi	if statement in bash
arg	mandatory argument
[arg]	optional argument

An easier way to avoid the NOTES file would be

```
cd ${1}
ls * | grep -v NOTES
```

Assignment:

Write a script, called `cleanedData.sh`, which does the following, when run from the data directory:

- Copies all of the data files from `data/alexander`, `data/Bert`, etc. to a new directory, `'data/cleaneddata'`.
- Renames the data files in `cleaneddata`; the names must have the format `"Data.NNN.txt"`, where `NNN` is the three-digit number that goes with the original file.

The final result should be 351 files, ranging from `Data.211.txt` to `Data.561.txt`, in the `cleaneddata` directory. No `NOTES` files should be present.

Be sure to ask questions. Use `man` pages if you need to. If there is some functionality that you would like, and you hope that a command exists that does that, ask me. You should be able to do this with everything you have been taught thus far.

Assignment:

```
#!/bin/bash
mkdir cleaneddata
for i in alexander Bert Frank_Richard gerdal jamesm Lawrence THOMAS
do
for filename in $i/*
do
if [ $filename != "$i/NOTES" ]
then
if [ $i == "alexander" ]
then
tmp='echo $filename| cut -c 16-18'
echo "cp $filename cleaneddata/Data.$tmp.txt"
cp $filename cleaneddata/Data.$tmp.txt
fi
if [ $i == "Bert" ]
then
tmp='echo $filename| cut -c 20-22'
echo "cp $filename cleaneddata/Data.$tmp.txt"
cp $filename cleaneddata/Data.$tmp.txt
fi
if [ $i == "Frank_Richard" ]
then
tmp='echo $filename| cut -c 20-22'
echo "cp $filename cleaneddata/Data.$tmp.txt"
cp $filename cleaneddata/Data.$tmp.txt
fi
if [ $i == "gerdal" ]
then
tmp='echo $filename| cut -c 13-15'
echo "cp $filename cleaneddata/Data.$tmp.txt"
cp $filename cleaneddata/Data.$tmp.txt
fi
fi
```

Assignment:

```
if [ $i == "jamesm" ]
then
    tmp='echo $filename| cut -c 13-15'
    echo "cp $filename cleaneddata/Data.$tmp.txt"
    cp $filename cleaneddata/Data.$tmp.txt
fi
if [ $i == "Lawrence" ]
then
    tmp='echo $filename| cut -c 15-17'
    echo "cp $filename cleaneddata/Data.$tmp.txt"
    cp $filename cleaneddata/Data.$tmp.txt
fi
if [ $i == "THOMAS" ]
then
    tmp='echo $filename| cut -c 9-11'
    echo "cp $filename cleaneddata/Data.$tmp.txt"
    cp $filename cleaneddata/Data.$tmp.txt
fi
fi
done
done
```

Assignment: other solution using function

```
#!/bin/bash
function MyCopy {
    num=$((($2 + 2))
    tmp='echo $1| cut -c $2-$num'
    echo "cp $1 cleaneddata/Data.$tmp.txt"
    cp $1 cleaneddata/Data.$tmp.txt
}
mkdir cleaneddata
for i in alexander Bert Frank_Richard gerdal jamesm Lawrence THOMAS
do
for filename in $i/*
do
if [ $filename != "$i/NOTES" ]
then
if [ $i == "alexander" ]
then
    MyCopy $filename 16
fi
if [ $i == "Bert" ]
then
    MyCopy $filename 20
fi
if [ $i == "Frank_Richard" ]
then
    MyCopy $filename 20
fi
if [ $i == "gerdal" ]
then
    MyCopy $filename 13
fi
fi
fi
fi
```

Assignment: other solution using function

```
if [ $i == "jamesm" ]
then
    MyCopy $filename 13
fi
if [ $i == "Lawrence" ]
then
    MyCopy $filename 15
fi
if [ $i == "THOMAS" ]
then
    MyCopy $filename 9
fi
fi
done
done
```