Basic Bash programming

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Reminder: Sign up to Piazza

- Great knowledge and discussion platform.
- Allows students to ask and answer questions.
- Lecturers/teaching assistants will be active on the platform as well.
- We will announce course updates (for example the video recordings).
- Sign up: http://piazza.com/uio.no/fall2016/inf3331inf4331

Assignment 1

- Deadline: Extended to this Friday
- Make sure that your solutions are pushed to your github repository. All files should be in https: //github.com/UiO-INF3331/INF3331-<UiO-Username>

Assignments 2

- Assignment 2 is online now.
- Topic: Bash scripting (lecture today).



Overview of Unix shells

- The original scripting languages were (extensions of) command interpreters in operating systems
- Primary example: Unix shells
- Bourne shell (sh) was the first major shell
- C and TC shell (csh and tcsh) had improved command interpreters, but were less popular than Bourne shell for programming
- Bourne Again shell (Bash/'bash'): GNU/FSF improvement of Bourne shell
- Other Bash-like shells: Dash (dash), Korn shell (ksh), Z shell (zsh)
- Bash is the dominating Unix shell today

Why learn Bash?

- Learning Bash means learning Unix
- Learning Bash means learning the roots of scripting (Bourne shell is a subset of Bash)
- Shell scripts, especially in Bourne shell and Bash, are frequently encountered on Unix systems
- Bash is widely available (open source) and the dominating command interpreter and scripting language on today's Unix systems

Why learn Bash? (2)

- Shell scripts evolve naturally from a workflow:
 - A sequence of commands you use often are placed in a file
 - 2 Command-line options are introduced to enable different options to be passed to the commands
 - Introducing variables, if tests, loops enables more complex program flow
 - At some point pre- and postprocessing becomes too advanced for bash, at which point (parts of) the script should be ported to Python or other tools
- Shell scripts are often used to glue more advanced scripts in Perl and Python

Remark

- We use plain Bourne shell (/bin/sh) when special features of Bash (/bin/bash) are not needed
- Most of our examples can in fact be run under Bourne shell (and of course also Bash)
- In Mac OSX, the Bourne shell (/bin/sh) is just a link to Bash (/bin/bash). In Ubuntu, the Bourne shell (/bin/sh) is a link to Dash, a minimal, but much faster shell than bash.

More information

- Offline documentation: man bash
- Bash reference manual: www.gnu.org/software/bash/manual/bashref.html
- Advanced Bash-Scripting Guide: http://www.tldp.org/LDP/abs/html
- Simple and good tutorial: http: //www.linuxintro.org/wiki/Shell_scripting_tutorial

What Bash is good for

- File and directory management
- Systems management (build scripts)
- Combining other scripts and commands
- Rapid prototyping of more advanced scripts
- Very simple output processing, plotting etc.

What Bash is not good for

- Cross-platform portability
- Graphics, GUIs
- Interface with libraries or legacy code
- More advanced post processing and plotting
- Calculations, math etc.

Some common tasks in Bash

- file writing
- for-loops
- running an application
- pipes
- writing functions
- file globbing, testing file types
- copying and renaming files, creating and moving to directories, creating directory paths, removing files and directories
- directory tree traversal
- packing directory trees

Bash example 1: Hello world

```
#!/bin/bash
echo "Hello world"
```

Two options to run this script:

- Type the commands directly in the bash shell (only feasible for small scripts).
- Save the code as helloworld.sh and run with:

```
chmod a+x helloworld.sh # Make script executable
./helloworld.sh
```

Bash example 2: Hello world, v2

```
#!/bin/bash
x=world
echo "Hello ${x}!"
```

Bash example 3: Line counter

Bash variables and commands

- Assign a variable by x=3.4, retrieve the value of the variable by \$x or \${x} (also called variable substitution).
- Variables passed as command line arguments when running a script are called positional parameters.
- Bash has a number of built in commands. Type help or help
 less to get a list.
- The real power comes from all the available Unix commands, in addition to your own applications and scripts. Try:

```
curl -s http://www.yr.no/place/Norway/Oslo/Oslo/Oslo/ |
   grep -m 1 "temperature" | cut -d"\"" -f4
```

Bash variables (1)

Variables in Bash are untyped!

Generally treated as character arrays, but permit simple arithmetic and other operations

Variables can be explicitly declared to integer or array;

Bash variables (2)

The echo command is used for writing:

```
s=42
echo "The answer is $s"
```

and variables can be inserted in the text string (variable interpolation)

Frequently seen variables:

Command line arguments:

```
$1 $2 $3 $4 and so on
```

- All command line arguments as array: \$0
- Number of command line arguments: \$#
- The exit status of the last executed command: \$? (is 0 if command was succesfull)

Bash variables (3)

Comparison of two integers use a syntax different from comparison of two strings:

Unless you have declared a variable to be an integer, assume that all variables are strings and use double quotes (strings) when comparing variables in an if test

```
if [ "$?" != "0" ]; then # this is safe if [ $? != 0 ]; then # might be unsafe
```

Convenient debugging tool: -x

Each source code line is printed prior to its execution if you add -x as option to /bin/bash

Either in the header

```
#!/bin/bash -x
```

or on the command line:

```
unix> /bin/bash -x hw.sh
unix> sh -x hw.sh # only works if sh links to bash
unix> bash -x hw.sh
```

Very convenient during debugging

Combining bash commands (1)

- The power of Unix lies in combining simple commands into powerful operations
- Standard bash commands and Unix applications normally do one small task
- Text is used for input and output easy to send output from one command as input to another

Combining bash commands (2)

Two standard ways to combine commands:

• The pipe, sends the output of one command as input to the next:

```
ls -1 | grep 3331
```

Will list all files having 3331 as part of the name

Executing a command, storing the result as a variable:

```
time=$(date)
time='date'
```

Combining bash commands (3)

More useful applications of pipes:

```
# send files with size to sort -rn
# (reverse numerical sort) to get a list
# of files sorted after their sizes:
/bin/ls -s | sort -rn

cat $case.i | oscillator
# is the same as
oscillator < $case.i</pre>
```

Make a new application: sort all files in a directory tree root, with the largest files appearing first, and equip the output with paging functionality:

```
du -a root | sort -rn | less
```

Bash redirects

Redirects are used to pass output to either a file or stream.

```
echo "Hei verden" > myfile.txt  # Save output to file
wc -w < myfile.txt  # Use file content as command input
```

Note: Pipes can be (in a clumpsy way) reimplemented with redirects:

```
prog1 > myfile && prog2 < myfile
is the same as
prog1 | prog2
```

Redirects and stdin, stdout, stderr

You can print to stderr with:

```
echo "Wrong arguments" >&2
```

Redirects and pipes can be combined:

```
./compile 2>&1 | less # View both stdout and stderr in less
```

Example: the classical Unix script

A combination of commands, or a single long command, that you use often;

```
./pulse_app -cmt WinslowRice -casename ellipsoid < ellipsoid.i | tee main_output
```

(should be a single line) In this case, flexibility is often not a high priority. However, there is room for improvement:

- Not possible to change command line options, input and output files
- Output file main_output is overwritten for each run
- Can we edit the input file for each run?

Problem 1; changing application input

In many cases only one parameter is changed frequently;

Still not very flexible, but in many cases sufficient. More flexibility requires more advanced parsing of command line options, which will be introduced later.

Problem 2; overwriting output file

A simple solution is to add the output file as a command line option, but what if we forget to change this from one run to the next?

Simple solution to ensure data is never over-written:

Problem 2; overwriting output file (2)

Alternative solutions;

- Use process ID of the script (\$\$, not really unique)
- mktemp can create a temporary file with a unique name, for use by the script
- Check if subdirectory exists, exit script if it does;

```
dir=$case
# check if $dir is a directory:
if [ -d $dir ]
    #exit script to avoid overwriting data
    then
        echo "Output directory exists, provide a different name"
        exit
fi
mkdir $dir # create new directory $dir
cd $dir # move to $dir
```

Alternative if-tests

As with everything else in Bash, there are multiple ways to do if-tests:

```
# the 'then' statement can also appear on the next line:
if [ -d $dir ]; then
    exit
fi

# another form of if-tests:
if test -d $dir; then
    exit
fi

# and a shortcut:
[ -d $dir ] && exit
test -d $dir && exit
```

Problem 3; can we edit the input file at run time?

- Some applications do not take command line options, all input must read from standard input or an input file
- A Bash script can be used to equip such programs with basic handling of command line options
- We want to grab input from the command line, create the correct input file, and run the application

File reading and writing

File writing is efficiently done by 'here documents':

```
cat > myfile <<EOF
multi-line text
can now be inserted here,
and variable substition such as
$myvariable is
supported.
EOF</pre>
```

The final EOF must start in column 1 of the script file.

EOF is an arbitrary keyword here.

Parsing command-line options

Alternative to case: if

case "\$option" in

case is standard when parsing command-line arguments in Bash, but if-tests can also be used. Consider

```
-m)
            m=$1; shift; ;; # load next command-line arg
        -b)
            b=$1; shift; ;;
        *)
            echo "$0: invalid option \"$option\""; exit ;;
     esac
versus
     if [ "$option" == "-m" ]; then
         m=$1; shift; # load next command-line arg
     elif [ "$option" == "-b" ]; then
         b=$1: shift:
     else
         echo "$0: invalid option \"$option\""; exit
     fi
 echo "Command line arguments:"
 [ -n "$m" ] && echo "m=$m"
 [ -n "$b" ] && echo "b=$b"
```

After assigning variables, we can write the input file

Then execute the program as usual

Redirecting input to read from the new input file

```
../pulse_app < $infile
```

We can add a check for successful execution. The shell variable \$? is 0 if last command was successful, otherwise \$? != 0.

```
if [ "$?" != "0" ]; then
  echo "running pulse_app failed"; exit 1
fi
# exit n sets $? to n
```

Other uses of cat

For-loops

What if we want to run the application for multiple input files?

```
./run.sh test1.i test2.i test3.i test4.i
or
./run.sh *.i
```

A for-loop over command line arguments

```
for arg in $0; do
    ../../build/app/pulse_app < $arg
done</pre>
```

Can be combined with more advanced command line options, output directories, etc...

For-loops (2)

For loops for file management:

```
files='ls *.tmp'
for file in $files
do
  echo removing $file
rm -f $file
done
```

Declare an integer counter:

```
declare -i counter
counter=0
# arithmetic expressions must appear inside (( ))
((counter++))
echo $counter # yields 1

For-loop with counter:

declare -i n; n=1
for arg in $@; do
    echo "command-line argument no. $n is <$arg>"
        ((n++))
done
```

C-style for-loops

```
declare -i i
for ((i=0; i<$n; i++)); do
  echo $c
done</pre>
```

Example: bundle files

Idea:

- Scripts packs a series of files into one file
- Executing this single file as a Bash script packs out all the individual files again

Usage:

```
bundle file1 file2 file3 > onefile # pack
bash onefile # unpack
```

Writing bundle is easy:

```
#/bin/sh
for i in $0; do
    echo "echo unpacking file $i"
    echo "cat > $i <<EOF"
    cat $i
    echo "EOF"
done</pre>
```

The bundle output file

```
Consider 2 fake files; file1
 Hello, World!
 No sine computations today
and file2
 1.0 2.0 4.0
 0.1 0.2 0.4
Running bundle file1 file2 yields the output
 echo unpacking file file1
 cat > file1 <<EOF
 Hello, World!
 No sine computations today
 EOF
 echo unpacking file file2
 cat > file2 <<EOF
 1.0 2.0 4.0
 0.1 0.2 0.4
 EOF
```

Running an application

Running in the foreground:

```
cmd="myprog -c file.1 -p -f -q";
$cmd < my_input_file

# output is directed to the file res
$cmd < my_input_file > res

# process res file by Sed, Awk, Perl or Python
```

Running in the background:

```
\label{eq:myprog} \mbox{ -c file.1 -p -f -q < my_input_file \&}
```

or stop a foreground job with Ctrl-Z and then type bg

Functions

```
function system {
# Run operating system command and if failure, report and abort
  "$@"
  if [ $? -ne 0 ]; then
    echo "make.sh: unsuccessful command $0"
    echo "abort!"
   exit 1
 fi
# function arguments: $1 $2 $3 and so on
# return value: last statement
# call:
name = mydoc
system pdflatex $name
system bibtex $name
```

How to return a value from a function? Define a new variable within the function - all functions are global!

File globbing, for loop on the command line

List all .ps and .gif files using wildcard notation:

```
files='ls *.ps *.gif'
# or safer, if you have aliased ls:
files='/bin/ls *.ps *.gif'
# compress and move the files:
gzip $files
for file in $files; do
    mv ${file}.gz $HOME/images
```

Testing file types

```
if [ -f $myfile ]; then
    echo "$myfile is a plain file"
fi
# or equivalently:
if test -f $myfile; then
    echo "$myfile is a plain file"
fi
if [ ! -d $myfile ]; then
    echo "$myfile is NOT a directory"
fi
if [ -x $myfile ]; then
    echo "$myfile is executable"
fi
[ -z $myfile ] && echo "empty file $myfile"
```

Rename, copy and remove files

```
# rename $myfile to tmp.1:
mv $mvfile tmp.1
# force renaming:
mv -f $myfile tmp.1
# move a directory tree my tree to $root:
mv mytree $root
# copy myfile to $tmpfile:
cp myfile $tmpfile
# copy a directory tree mytree recursively to $root:
cp -r mytree $root
# remove myfile and all files with suffix .ps:
rm myfile *.ps
# remove a non-empty directory tmp/mydir:
rm -r tmp/mydir
```

Directory management

The find command

Very useful command!

find visits all files in a directory tree and can execute one or more commands for every file

Basic example: find the oscillator codes

```
find $scripting/src -name 'oscillator*' -print
```

Or find all PostScript files

```
find $HOME \( -name '*.ps' -o -name '*.eps' \) -print
```

We can also run a command for each file:

```
find rootdir -name filenamespec -exec command {} \; -print
# {} is the current filename
```

Applications of find (1)

Find all files larger than 2000 blocks a 512 bytes (=1Mb):

```
find $HOME -name '*' -type f -size +2000 -exec ls -s \{\}\ \;
```

Remove all these files:

```
find $HOME -name '*' -type f -size +2000 \
-exec ls -s {} \; -exec rm -f {} \;
```

or ask the user for permission to remove:

```
find $HOME -name '*' -type f -size +2000 \
    -exec ls -s {} \; -ok rm -f {} \;
```

Applications of find (2)

Find all files not being accessed for the last 90 days:

Tar and gzip

The tar command can pack single files or all files in a directory tree into one file, which can be unpacked later

```
tar -cvf myfiles.tar mytree file1 file2
# options:
# c: pack, v: list name of files, f: pack into file
# unpack the mytree tree and the files file1 and file2:
tar -xvf myfiles.tar
# options:
# x: extract (unpack)
```

The tarfile can be compressed:

```
gzip mytar.tar
# result: mytar.tar.gz
```

Two find/tar/gzip examples

Pack all PostScript figures:

```
tar -cvf ps.tar 'find $HOME -name '*.ps' -print'
gzip ps.tar
```

Pack a directory but remove CVS directories and redundant files