



# **URPP** tutorial

Basic bash scripting 2 & Introduction to cluster submission system

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# What is a bash script?

- Mainly a set of commands that can be executed in the terminal
  - Text file with commands
  - Should begin with the shebang line (#!/bin/bash)
  - Executed line by line 

    new line = new command
- Example: script.sh

```
#!/bin/bash
#this is a comment
echo "Hello World"

chmod +x script.sh
./script.sh
Hello World

shebang line
A comment (ignored)
Print something in terminal

Give execution permission
Run script
```

# Variables and arrays

#### Variables:

- Store data and configuration options
- Create: name followed by "=" and the value
- Call: "\$" in front of the name, enclose the name in {} if directly followed by something else

```
a=/home/user
echo "Data folder: ${a}/data"
```

## **Arrays:**

- Arrays are variables containing multiple values
- Create: name followed by "=()", values are space separated
- Values can be accessed by their index (number) starting from 0

```
col=(red blue yellow)
echo "The sun is ${col[2]}"
```

# For loop

```
for VARIABLE in 1 2 3
do
command1
done
```

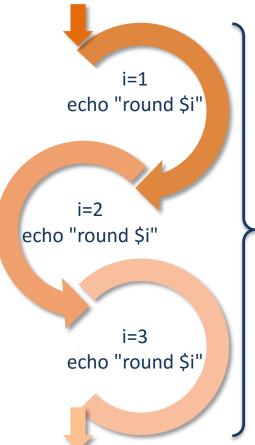
Can be used to repeat certain tasks

## • Examples:

```
#!/bin/bash
for i in 1 2 3
do
    echo "round $i"
done
```

```
#!/bin/bash
for i in {1..3}
do
   echo "round $i"
done
```

```
round 1
round 2
round 3
```



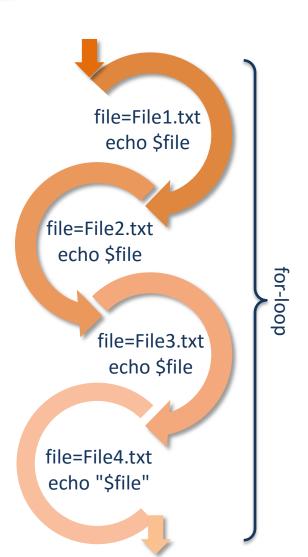
# For loop

Examples:

## Go through all files in a folder:

```
#!/bin/bash
for file in /home/user/*
do
   echo $file
done
```

```
File1.txt
File2.txt
File3.txt
File4.txt
```



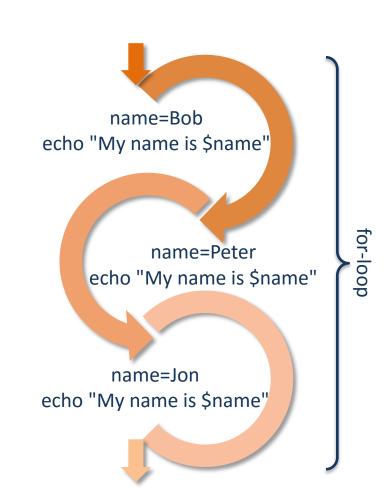
# For loop

Examples:

## Go through all items of an array:

```
#!/bin/bash
names=(Bob Peter Jon)
for name in ${names[*]}
do
   echo "My name is $name"
done
```

```
My name is Bob
My name is Peter
My name is Jon
```



# While loop

Can also be used to repeat certain tasks:

```
while [[ something ]]
do
   command
done
```

• Example:

```
#!/bin/bash
count=0
while [[ $count -lt 4 ]]
do
    echo $count
    let count+=1
done
echo "done"
```

```
0
1
2
3
done
```

 If statements can be used to check for something and do something else depending on the outcome of the check

```
if [[ someting ]]
then
   command1
elif [[ something ]]
then
   command2
else
   command3
fi
```

## Boolean operators:

• **-e FILE**: True if file exists

```
#!/bin/bash
if [[ -e log.txt ]]
then
   echo "log file exist"
else
   echo "log file doesn't exist"
fi
```

#### Boolean operators:

• **-e FILE**: True if file exists

STRING = STRING: True if first string is identical to the second

STRING != STRING: True if first string is not identical to the second

STRING < STRING: True if first string sorts before the second</li>

STRING > STRING: True if first string sorts after the second

```
#!/bin/bash
name=Bob
if [[ $name = "Rod" ]]
then
   echo "Your name is Rod"
else
   echo "Your name is not Rod"
fi
```

# #!/bin/bash a=10 if [ \$a -eq 7 ] then echo "Not equal to 7" else echo "You guessed \$a" fi

#### Boolean operators:

• **-e FILE**: True if file exists

STRING = STRING: True if first string is ider nical to the second

STRING != STRING: True if first string is not identical to the second

STRING < STRING: True if first string sorts before the second</li>

STRING > STRING: True if first string sorts after the second

• **INT -eq INT**: True if both integers are identical

• **INT -ne INT**: True if integers are not identical

INT -It INT: True if first integer is less than the second

• INT -gt INT: True if first integer is greater than the second

INT -le INT: True if first integer is less than or equal to the second

INT -ge INT: True if first integer is greater than or equal to the second

```
#!/bin/bash
a=Bob
if [[ $a = "Rob" || $a = "Tod" ]]
then
    echo "Your name is Rod or Tod"
else
    echo "Your name is not Rod or Tod"
fi
```

### Boolean operators:

• -e FILE: True if fil fi

• **STRING** = **STRING**: True if first string is identical to the second

STRING != STRING: True if first string is not identical to the second

STRING < STRING: True if first string sorts before the second</li>

• **STRING > STRING**: True if first string sorts after the second

INT -eq INT: True if both integers are identical

INT -ne INT: True if integers are not identical

INT -It INT: True if first integer is less than the second

• INT -gt INT: True if first integer is greater than the second

INT -le INT: True if first integer is less than or equal to the second

• INT -ge INT: True if first integer is greater than or equal to the second

! EXPR: Inverts the result of the expression (logical NOT)

EXPR && EXPR: True if both expressions are true (logical AND)

• **EXPR** | **EXPR**: True if either expression is true (logical OR)

## **Functions**

- Functions: blocks of commands
- Code that you may call multiple times within your script
- Arguments can be passed to functions and be accessed by \$1 (first argument), \$2 (second argument),...
- Example:

```
#!/bin/bash
sum() {
  echo "$1 + $2 = $(($1 + $2))"
}

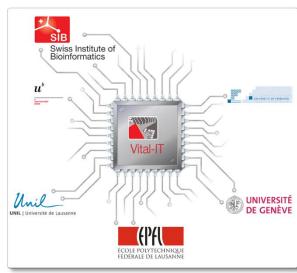
sum 1 4
sum 8 7

Call function

1 + 4 = 5
8 + 7 = 15
```

# **Cluster submission systems**

- Usually a cluster consist of
  - Front-end machine (where user log in)
    - → where you submit jobs
  - Compute nodes
    - → where computations take place
- Specialized software for submitting jobs to compute nodes
  - LSF (Load Sharing Facility):Vital-IT (SIB)
  - SGE (Sun Grid Engine):Wagner-cluster (UZH)
  - Slurm workload manager:
     Hydra/ Piz Dora (S3IT UZH)



## When should I use a cluster?

- Application
  - runs too long on a single computer
  - takes too much CPU
- The problem (input data) can be split into pieces which can be executed in parallel Examples:
  - BLAST 1 million sequences
  - simulate model 1 million times
- Every program (pipeline) to be executed on a cluster needs to be defined as a "job" with
  - Executable , input data, output data
  - Job characteristics (CPU, memory, run time,...)

## **Basics**

Get info about job status:

```
- LSF: bjobs
```

- SGE: qstat

- Slurm: squeue

```
squeue
JOBID PARTITION
                NAME
                      USER ST
                               TIME NODES NODELIST (REASON)
65646
         batch
                chem mike R 24:19
                                        2 adev[7-8]
               bio joan R 0:09
65647
         batch
                                        1 adev14
65648
                      phil PD
         batch
               math
                               0:00
                                         6 (Resources)
```

Kill jobs:

```
- LSF: bkill jobID
```

- SGE: qdel jobID

- Slurm: squeue jobID

# Submit job

## Example command we want to execute:

bsub < ./blast.sh

```
blastall -p blastp -d "swiss" -i p123456.seq
```

#### – LSF:

Write shell script with job description: blast.sh

```
#!/bin/bash

#BSUB -o blast-output.txt → Standard output is redirected to this file

#BSUB -e blast-error.txt → Standard error is redirected to this file

#BSUB -J blastp → Job name

#BSUB -R "mem > 4000" → Minimal memory in MB

#BSUB -n 4

#BSUB -R "span[ptile=4]" → CPUs per process (multithreading)

blastall -p blastp -d "swiss" -i p123456.seq
```

# Submit job

#### — SGE:

Write shell script with job description: sge-blast.sh

```
qsub ./sge-blast.sh
```

# Submit job

#### – Slurm:

Write shell script with job description: slurm-blast.sh

```
sbatch ./slurm-blast.sh
```

# **Array job**

- Array job: submit the same job multiple times
  - Simulation has to run 20 times
    - → submit a single job rather than 20 individual ones
- LSF:
  - mysim.sh

```
#!/bin/bash

#BSUB -J array[1-20] → Will start 20 subjobs

#BSUB -o output-%I.txt → %I: get index of job array in header

#BSUB -e error-%I.txt

run-mysim --seed=$LSB_JOBINDEX → get index of job array
```

```
bsub < ./mysim.sh</pre>
```

# **Array job**

#### • SGE:

- mysim.sh

qsub ./mysim.sh

# **Array job**

#### • Slurm:

- mysim.sh

```
#!/bin/bash
#SBATCH --array=1-20 → Will start 20 subjobs
#SBATCH --time=1:0:0
#SBATCH --mem=16g
#SBATCH --output=output-%a.txt → get index of job array in header
#SBATCH --error=error-%a.txt
#SBATCH --job-name=array-job

srun run-mysim --seed=$SLURM_ARRAY_TASK_ID
```

sbatch ./mysim.sh

get index of job array

# **Acknowledgment**

## Bash scripting:

- http://www.allaboutlinux.eu/bash-script-for-beginners/
- http://www.howtogeek.com/67469/the-beginners-guide-to-shell-scripting-the-basics/
- http://bash.cyberciti.biz/guide/Main\_Page
- http://www.arachnoid.com/linux/shell\_programming.html
- http://www.tldp.org/LDP/Bash-Beginners-Guide/html/
- http://mywiki.wooledge.org/BashGuide
- http://ryanstutorials.net/bash-scripting-tutorial/bash-functions.php
- There are a lot of other online tutorials available and forums discussing diverse kinds of topics!

#### Cluster:

- SIB: HPC in Life Science workshop
- http://www.ceci-hpc.be/slurm\_tutorial.html
- http://www.s3it.uzh.ch/infrastructure/hydra/usage/