

Basics of Unix

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What are we going to learn today?

- Move around directory structure
- Create directories
- List and explore content of directories
- View plain text files
- To copy, move and rename files
- Work with compressed files
- Use variables and lists
- To download and install software

How to connect to remote machine

```
ssh userXX@ngs-course.duckdns.org
```

Command line

To type commands (syntax):

```
name -flag(value) input > output
```

```
head -n20 file.txt > out.txt
```

What if I don't know?

```
man head
```

```
head --help
```

```
head -h
```

Take a break and check your keyboard...

`[]` squared brackets

`{}` curly brackets

`<>` angle brackets (smaller-than, bigger-than
sign)

`()` parentheses

`~` tilde

`/` slash

`\` back slash

`|` pipe

`^` caret

`$` dollar sign

`:` colon

`;` semicolon

`.` dot

`,` comma

`#` hash

`_` underscore

`-` dash

`*` asterisk

`!` exclamation mark

`?` question mark

`&` ampersand

`@` at sign

`'` quotation mark single

`"` quotation mark double

screen

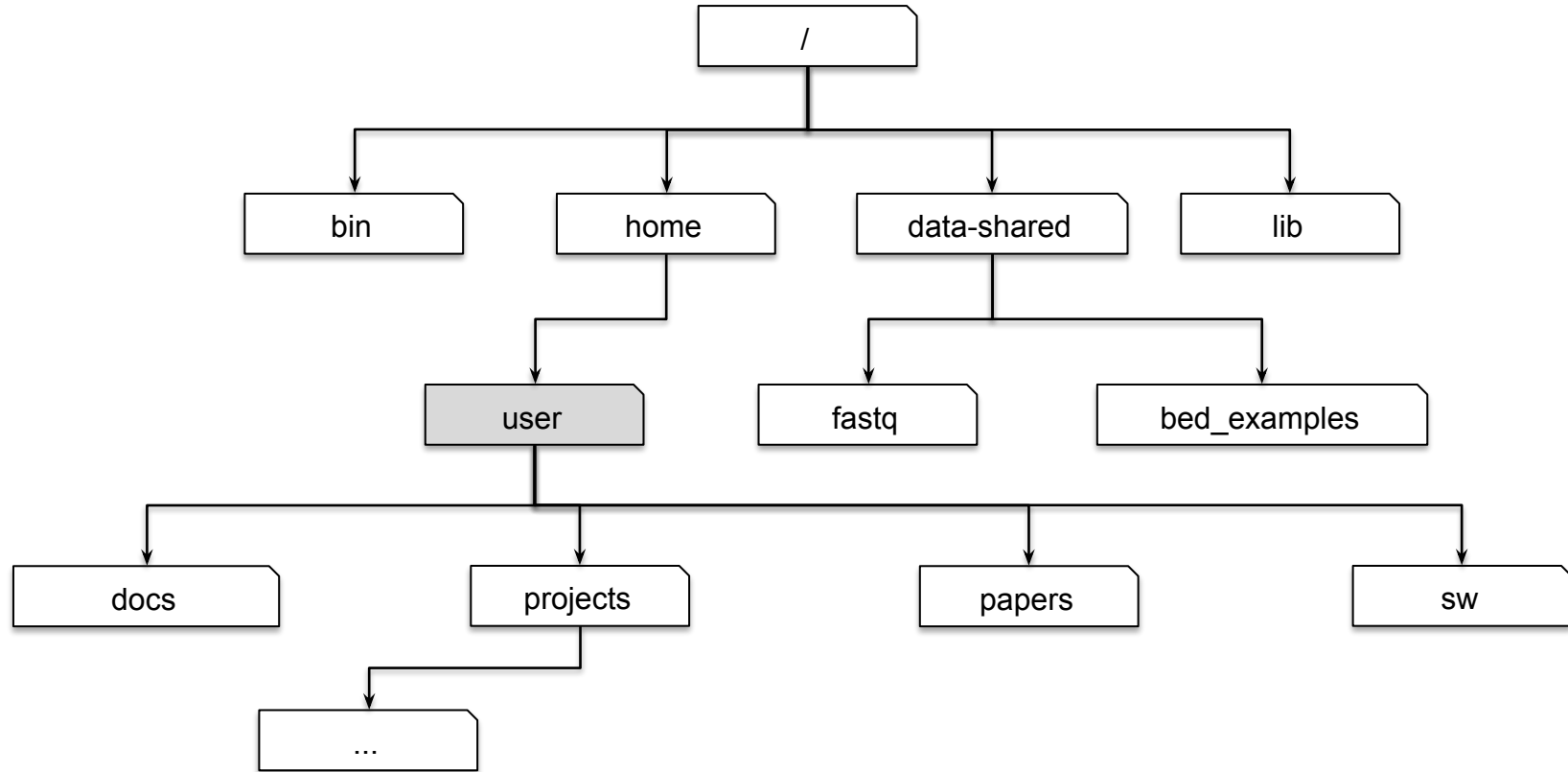
Protection from unexpected connection drop outs...

```
screen
screen -r
screen -ls
```

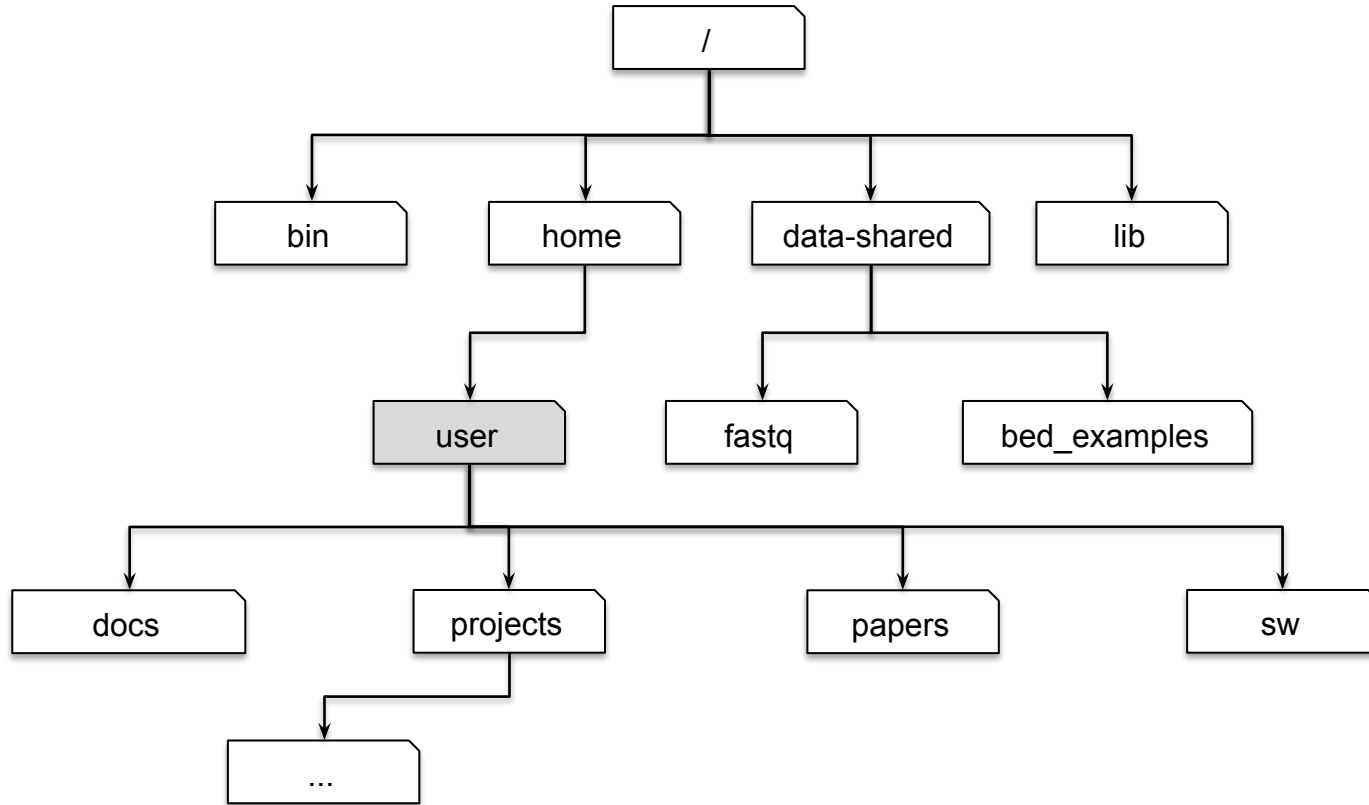
```
# inside screen
ctrl+a c      # new window
ctrl+a space  # switch between multiple windows
ctrl+a d      # detach from the screen
```

* ***tmux*** - a bit more user friendly and powerful alternative to *screen*

Basic directory structure of unix

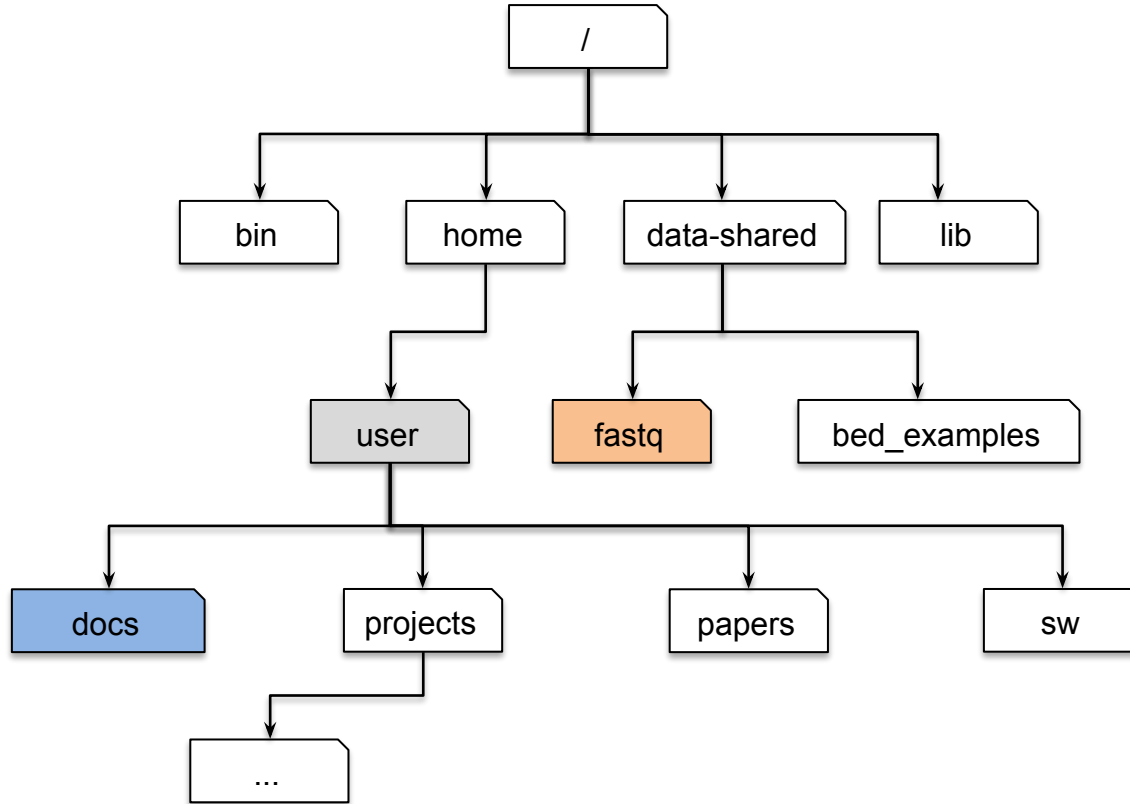


Moving around



```
pwd
ls
ls ~
ls /
ls ..
ls ../..
cd
cd ~
cd /
cd ..
cd -
```

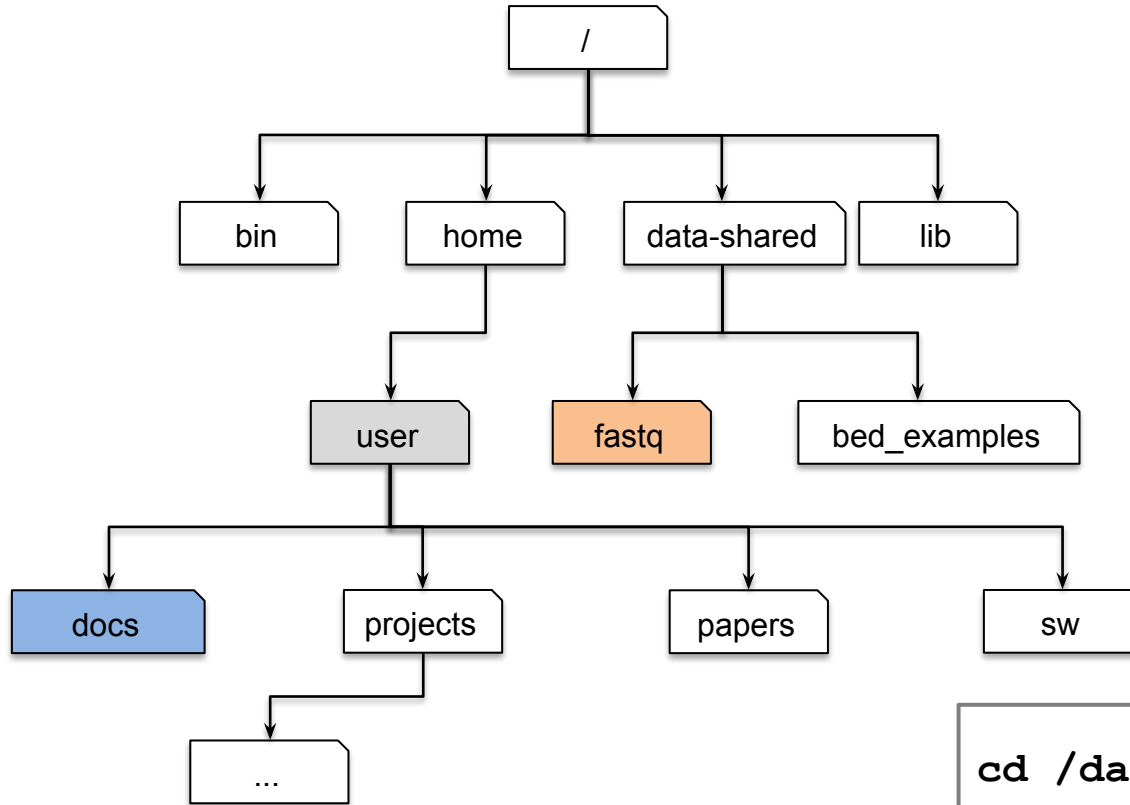

Absolute vs. relative path



Exercise:

Use absolute and relative path to move from 'docs' (blue) to 'fastq' (red)

Absolute vs. relative path

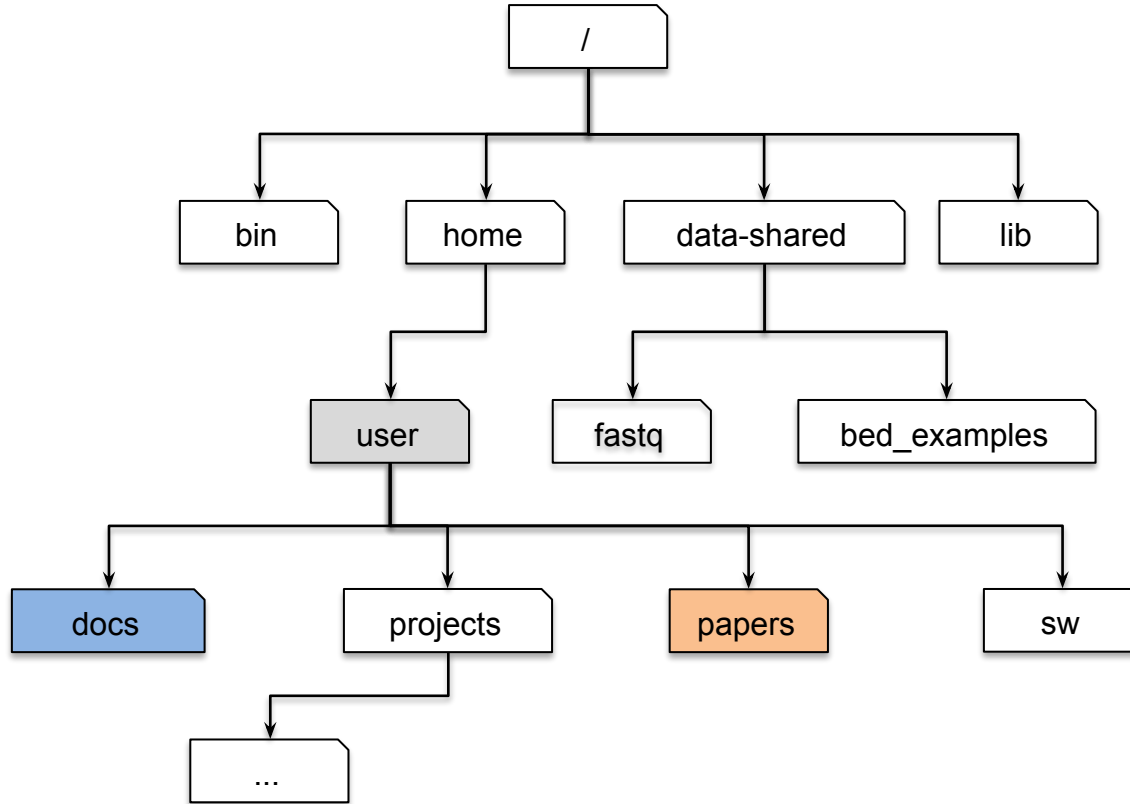


Exercise:

Use absolute and relative path to move from 'docs' (blue) to 'fastq' (red)

```
cd /data/fastq  
cd ../../../../data/fastq
```

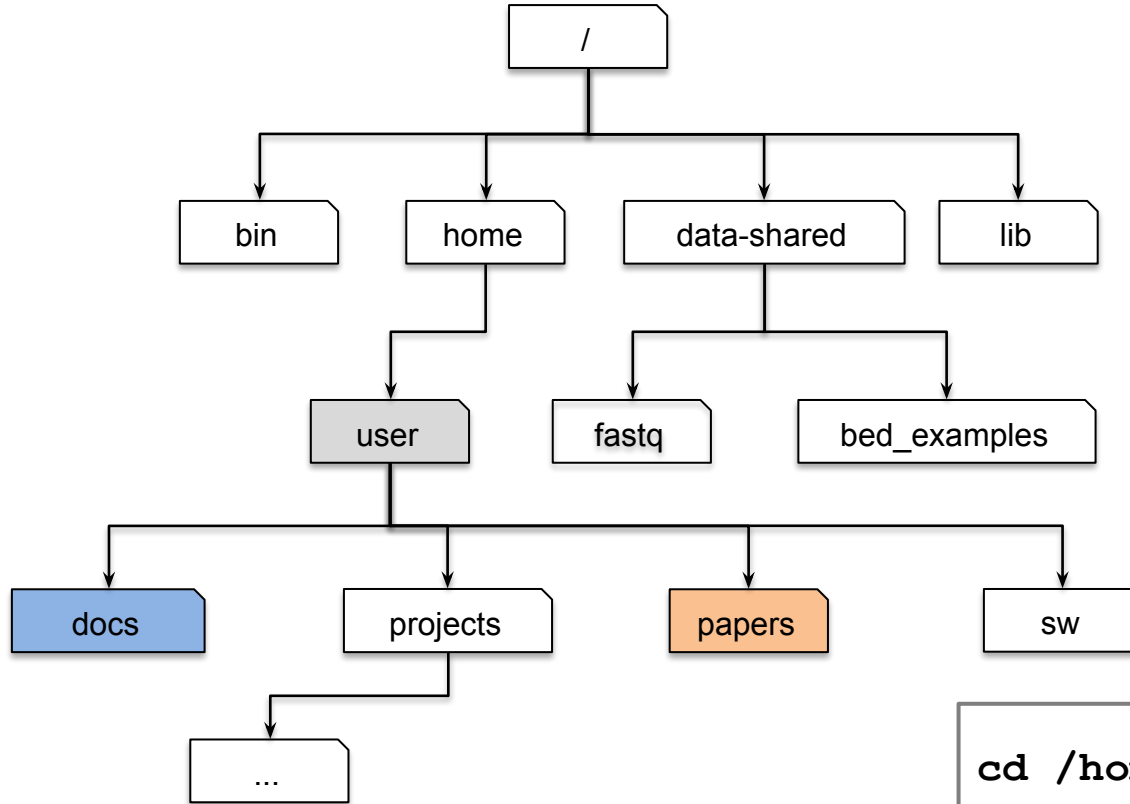
Absolute vs. relative path



Exercise:

Use absolute and relative path to move from 'docs' (blue) to 'projects' (red)

Absolute vs. relative path



Exercise:

Use absolute and relative path in to move from 'docs' (blue) to 'projects' (red)

```
cd /home/user/projects  
cd ../projects
```

Moving and copying files or directories

Try these tools to:

- make new files/(sub)directories
- move and rename them
- remove them

```
touch          # make empty file(s)
mv             # move/rename files
cp (-r)        # copy files (-r directories)
mkdir (-p)     # make directory (-p subdirectory)
rm (-r)        # remove file (-r non-empty directory)
```

Viewing plain text file content

```
less -SN
```

```
tail -n8
```

```
head -n8
```

```
cat
```

```
nano
```

Work with compressed data

```
# only gzipped (only one file)
```

```
gunzip file.gz
```

```
# view content of a compressed file
```

```
zcat fastq.gz | less
```

```
# gzipped tarball archive
```

```
tar -xzvf fastq.tar.gz
```

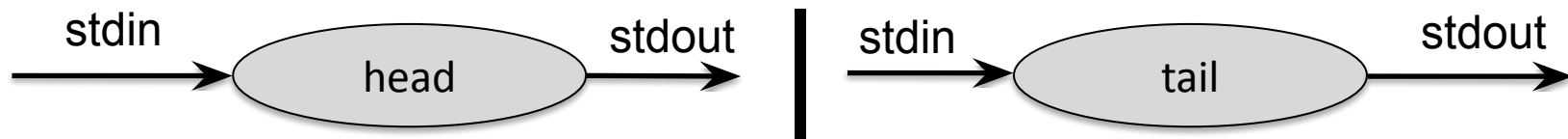
Exercise

Prepare FASTQ data file:

```
# go to home directory
cd
# make a new dir
mkdir projects/fastq && cd projects/fastq
# copy a fastq tarball to the new dir
cp /data-shared/fastq/fastq.tar.gz .
# decompress files
tar -zxvf fastq.tar.gz
# list files
ls -sh
```


Pipes '|'

Chaining standard input and output:



```
head -8 HRTMUOC01.RL12.00.fastq | tail -4 | less
```

Neater way to structure pipelines

```
< HRTMUOC01.RL12.00.fastq head -8 | tail -4 | less
```

Globbing & wildcards (*, ?, [class])

What if I need to choose multiple files?

```
cd ~/projects/fastq
```

```
ls *.fastq # choose all fastq files
```

```
ls HRTMUOC01.RL12.0?.fastq # one character
```

```
ls HRTMUOC01.RL12.0[1-9].fastq # one numerical character
```

Exercise

How many reads are in all fastq files?

```
cd ~/projects/fastq  
cat *.fastq | wc -l
```

```
expr XXXX / 4 ## Or  
echo $( (XXXX/4) )
```

Variables

Variable: storage location paired with an associated symbolic name

```
CPU=4
```

```
echo $CPU
```

```
FILE=~/projects/fastq/HRTMUOC01.RL12.00.fastq
```

```
echo $FILE
```

Loops

Loop over set of parameter values:

```
PARAM=$(echo {0..9})
```

```
for v in $PARAM
```

```
do
```

```
    echo $v;
```

```
Done
```

```
# one line syntax
```

```
For v in $PARAM; do echo $v; done
```

Installing software in Unix

- *The easiest way is to use package manager (apt-get)*

```
sudo apt-get install htop
```

```
sudo apt-get install tmux
```

Installing software in Unix

- *Otherwise we have to download the source code and compile it on its own (canonical way in Unix):*

```
# Downloading compressed source code
```

```
wget -O - ..url.. | tar xvz
```

```
# Cloning from Git repository
```

```
git clone ..url..
```

```
# Compilation of binaries
```

```
cd ...directory..
```

```
./configure
```

```
make
```

```
sudo make install
```

bedtools2

- *See our website*

```
# Download the compressed source code
```

```
wget https://github.com/arg5x/bedtools2/releases/download/v2.25.0/bedtools-2.25.0.tar.gz
```

```
tar -zxvf bedtools-2.25.0.tar.gz
```

```
# Or clone Git repository
```

```
git clone https://github.com/arg5x/bedtools2
```

```
# Compile binaries
```

```
cd bedtools2
```

```
make
```


What have we learned today?

- Move around directory structure
- Create directories
- List and explore content of directories
- View plain text files
- To copy, move and rename files
- Read compressed files
- Use variables and lists
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Why should we learn unix shell? ChatGPT...

- You are welcome to use it as it is common these days among programmers but be aware of errors - it is a good servant requiring instant double-check

That's all for today...