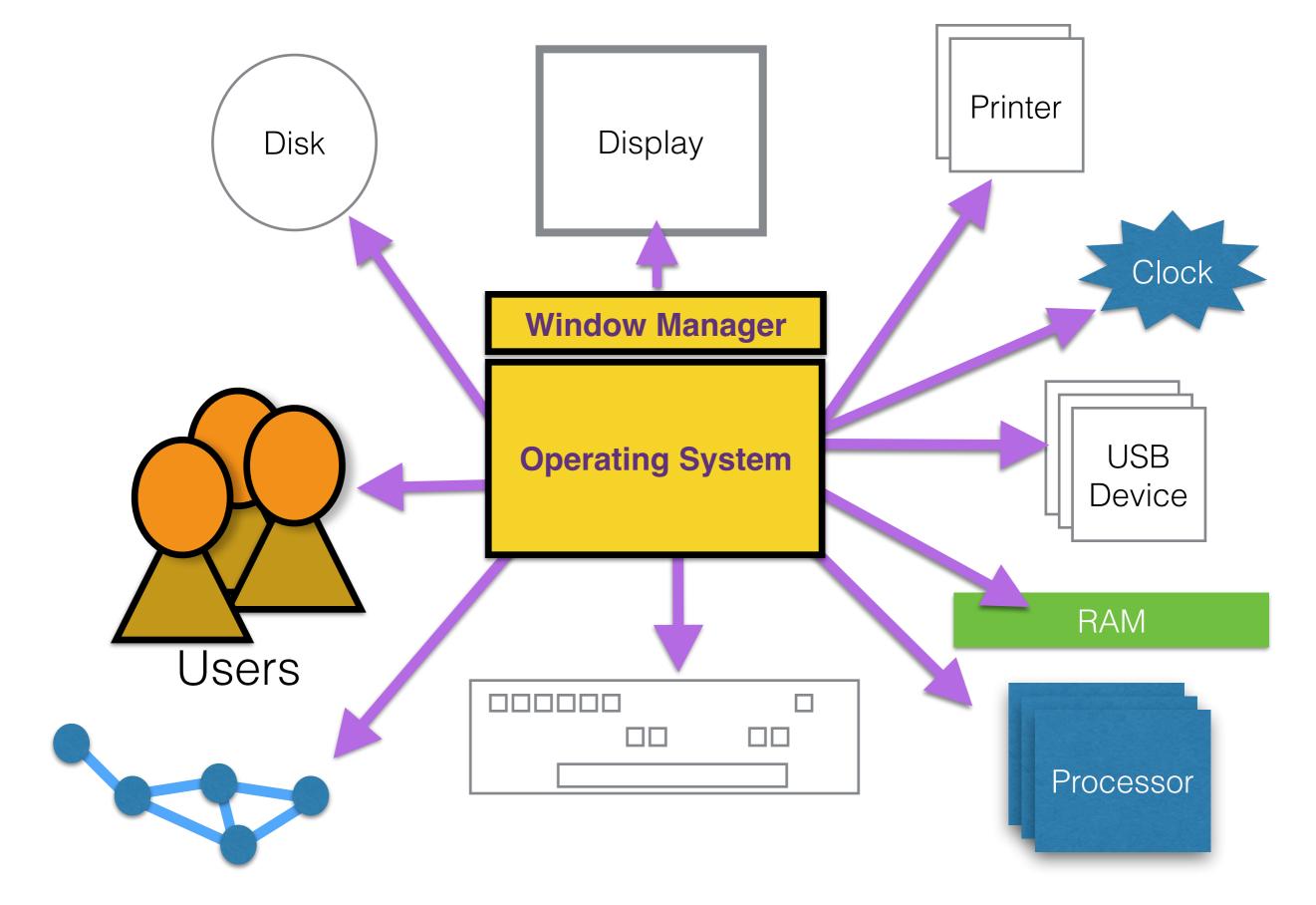
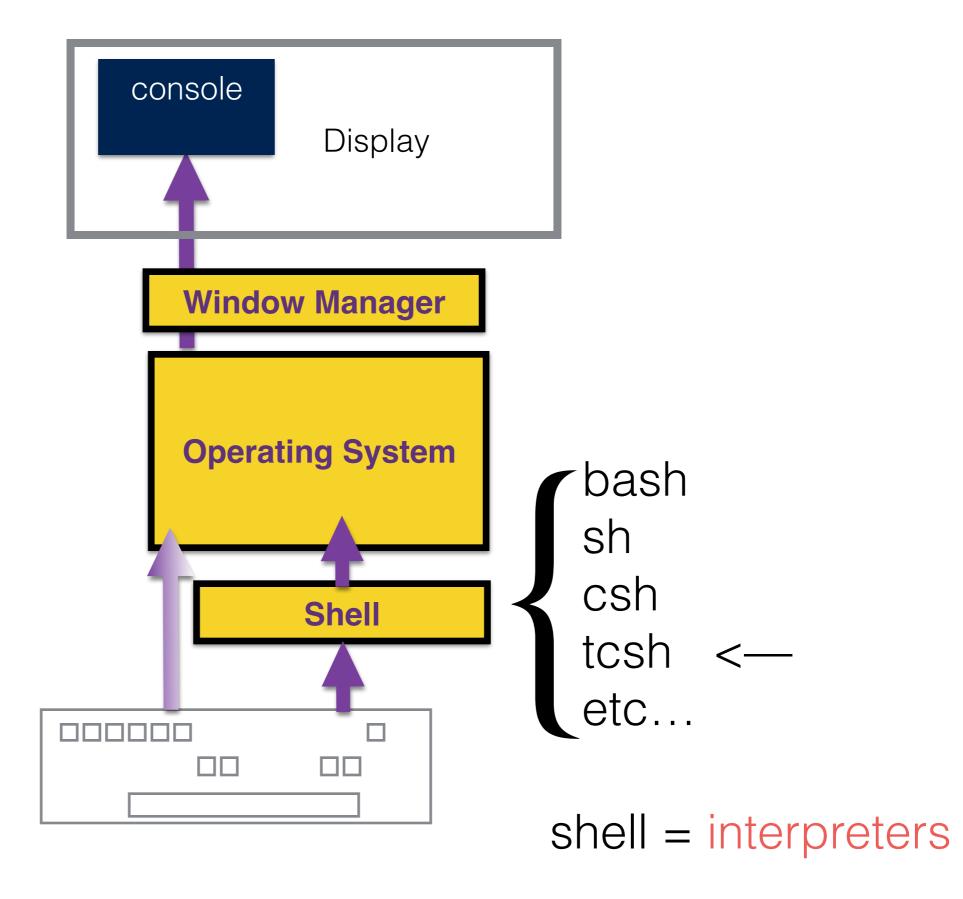
# Introduction to Programming The tcsh Shell

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# OS = manager of everything!





#### tcsh, the tee-cee-shell

http://en.wikipedia.org/wiki/Tcsh

# History

- The built-in history command displays the previously-entered commands
- Use of 1 / 1 at the command line to allow the user to select a command from the history to edit/execute
- Invoking previous commands using command history
  - !! executes the previous command
  - !n executes the nth command that was previously executed
  - !-n executes the command that was executed n commands ago
  - !string executes the most recently executed command that starts with string
  - !?string executes the most recently executed command that contains string
- Using history in new commands
  - !\* refers to all of the arguments from the previous command
  - . ! \$ refers to the last argument from the previous command
  - !^ refers to the first argument from the previous command
  - !:n refers to the nth argument from the previous command
  - !:m-n refers to the mth through nth arguments from the previous command
  - !:n-\$ refers to the nth through the last argument from the previous command

#### Variables

```
● ● ●
                                misc - ssh - 73 \times 21
          ssh
beowulf2:~/handout> set x=100
beowulf2:~/handout> echo $x
100
beowulf2:~/handout> set name=Sophia
beowulf2:~/handout> echo $name
Sophia
beowulf2:~/handout> set fullName="Smith College"
beowulf2:~/handout> echo $fullName
Smith College
beowulf2:~/handout> set files=`ls`
beowulf2:~/handout> echo $files
file1.txt file2.txt file3.txt fulghum generateDNAFiles.py generateDNA.py
last.txt README temp/ ulysses.txt
beowulf2:~/handout>
```

# Loops

```
beowulf2:~/handout> foreach x ( one two three )
foreach? echo $x
foreach? end
one
two
three
beowulf2:~/handout>
```

## More Loops

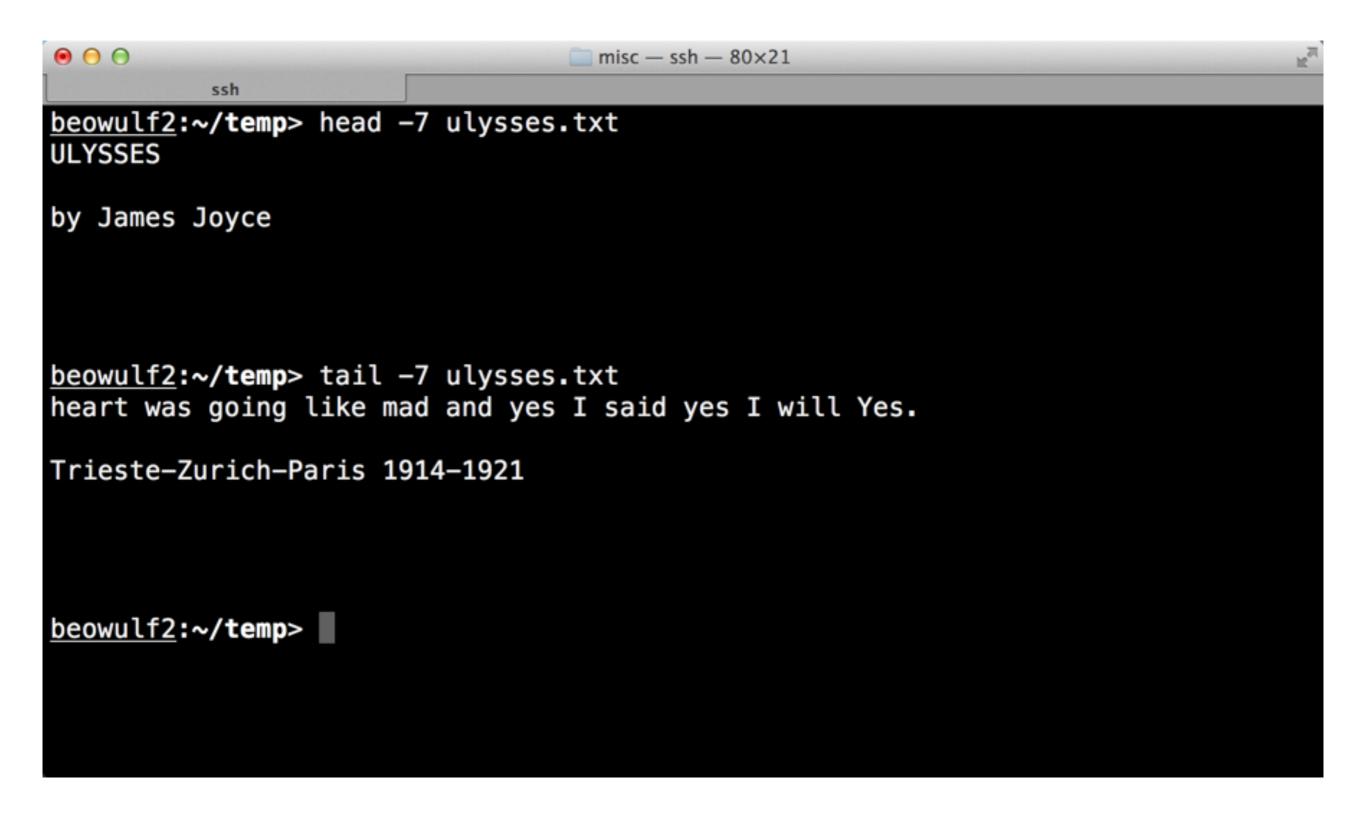
```
\Theta \cap \Theta
                                       misc - ssh - 80 \times 21
            ssh
beowulf2:~/handout/temp> foreach n ( `seq 0 9` )
foreach? echo "DNA_$n.txt"
foreach? end
DNA_0.txt
DNA_1.txt
DNA_2.txt
DNA_3.txt
DNA_4.txt
DNA_5.txt
DNA_6.txt
DNA_7.txt
DNA_8.txt
DNA_9.txt
beowulf2:~/handout/temp>
```

# Some Linux Commands

# Linux Command: grep

```
misc - ssh - 80 \times 21
beowulf2:~/temp> getcopy ulysses.txt
beowulf2:~/temp> grep magenta ulysses.txt
of an elder in Zion and a smokingcap with magenta tassels. Horned
beowulf2:~/temp>
beowulf2:~/temp> grep -i magenta ulysses.txt
of an elder in Zion and a smokingcap with magenta tassels. Horned
beowulf2:~/temp> wc ulysses.txt
  32663 264965 1520798 ulysses.txt
beowulf2:~/temp> wc -l ulysses.txt
32663 ulysses.txt
beowulf2:~/temp>
```

#### Linux Commands: tail & head



## Redirection & Pipes

# (No) Redirection



#### java HelloWorld

Hello World!

Is -1
file1.txt
file2.txt
file3.txt

#### Redirection



#### java HelloWorld > outputFile.dat

ls -1 > listFiles.txt
cat listFiles.txt

file1.txt file2.txt file3.txt

#### Repeated Redirection



java HelloWorld > outputFile.dat java OtherProg >! outputFile.dat

```
head -10 ulysses.txt > sample.txt #(create)
echo "..." >> sample.txt #(append)
tail -10 ulysses.txt >> sample.txt #(append)
```

#### Redirection



```
cat > data.in

1
2
3
^D
java MyProg < data.in
```

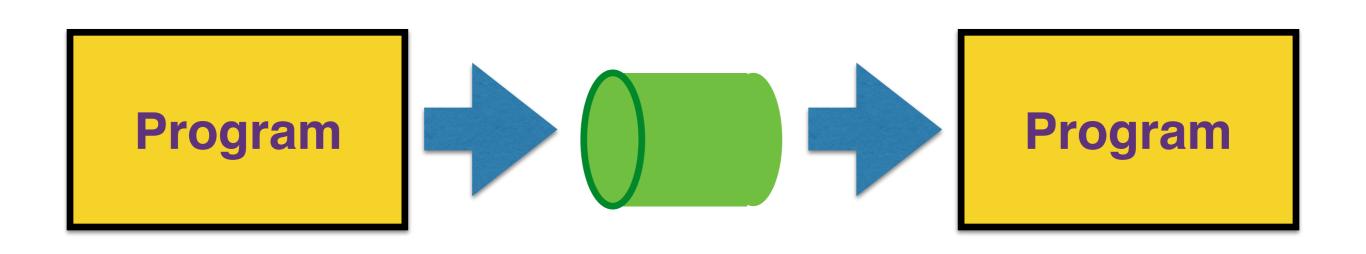
#### Redirection



wc < ulysses.txt > ulysses.info

**cat ulysses.info** 32663 264965 1520798

### Pipes



grep orange ulysses.txt | wc -l

grep orange ulysses.txt | tail -3

#### Problem:

- Get several DNA files from <a href="http://cs.smith.edu/">http://cs.smith.edu/</a>

   ~212a/DNAFiles/, named DNA\_0000.txt to

   DNA\_0099.txt
- Find how many files have the special string "CGTGACTCAA" in them, aka TPA-response element.
- Identify the files.

### Summary

- Using programming languages, one writes programs where the *library functions* allow one to manipulate **data**.
- Using shell commands, one can write programs, called *scripts*, where the functions available are operating-system commands, allowing one to manipulate operating system objects, such as files, Web data, devices, network information, or user information.

```
#! /bin/tcsh
# findDNA.sh
# D. Thiebaut
# Demo script for tcsh lecture, CSC212
#
# URL where DNA files are residing
set url=http://cs.smith.edu/~212a/DNAFiles
# pattern to search in DNA files
set pattern=CGTGACTCAA
# download all files DNA 000x where x is 0 to 9
foreach n ( `seq 0 9` )
   wget -q $url/DNA_000$n.txt
end
# download all files DNA_00x where x is 10 to 99
foreach n ( `seq 10 99` )
   wget -q $url/DNA_00$n.txt
end
# count number of files downloaded
set num=`ls DNA* | wc -l`
echo "Downloaded $num DNA files"
# identify files containing pattern and save to file
grep -l $pattern DNA* >! $pattern.txt
# remove all DNA files
unalias rm
rm DNA_*
```