Programming for Bioinformatics - part 3

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UNIX

- Unix was developped in 1960 at Bell labs by the founders of C
- It was one of the first OS to be multi-tasking, multi-user
- It has a hierarchical file system
- In the root directory we can find
 - /bin contains essential user command binaries
 - /etc configuration files
 - /sbin contains essential system binaries
 - /usr contains binaries and support files for user apps
 - /var contains variable data files
- It is written in C
- This part is about shell scripting
- Unix commands are mostly similar everywhere, but sometimes there are differences
- Alisases can be used for typing frequently used command parameters
 - They can be removed with the unalias command
 - They can be made permanent by putting in the .bashrc

Shell

- The shell is a language interpreter
- When I type a command, it searches for the command in what is in the \$PATH variable
 - /bin /usr/bin /usr/local/bin
- In order to execute commands that are not in \$PATH, i need to give the path
 - ./myscript
- I can write on multiple line by putting before pressing enter

File permissions

- They work for any file (also directories, which are indeed files)
- The fundamental permissions are r, w and x and they can be applied to owner, group and all
- The combination of permission of a file are represented with 3 bits for a single user
 - -000 is no permission, 100 is r-, 010 is -w-, 001 is -x and so on
- I can express a permission status by specifying 3 numbers, and so using octal numbers
 - 0 in octal means 000 in bynary, so it is —
 - 1 means 001, so -x
 - 7 means 111. so rwx
 - 000 is ----- or d----
 - 777 is -rwxrwxrwx or drwxrwxrwx
 - -345 is -wxr-r-x

Some commands

- echo is the Bash way for print
- Print the working directory: pwd
- Create a directory: mkdir
- Create an empty file: touch
 - If I touch an existing file, I change its acces and modification time
- Copy files or directories: cp
 - For doing recursively (for dirs) use cp -r
 - It can overwrite: use cp -i to ask for confirmation!
 - * I can also make an alias cp=cp -i
- Remove files: rm
 - There is no confirmation!
 - rm -r is recursive
 - rm -i asks for confirmation
- Remove empty directories: rmdir
- Move or rename: mv
- Scroll a file:less
 - I can search for words in less with \something
 - I can exit with q
 - more is a primitive version of less
- Search a file: find
 - I write first the directory in which I want to search and then, for instance, the name of the file
 - find . -name myfile.txt
 - I can also search by size, permission (-perm)
- Display the manual: man
- Path of a command: which
- All the paths to a command and associated files: whereis
- Quick one-line info on a command: whatis
- Info on a file: file
 - It tryes to guess the filetype based on its content
- Free disk space: df
- DIsk usage stats: du
- For both df and du the -h option makes the output human-readable
- Reverse a string: rev
- Simple calculations: bc
 - In order to operate on reals instead of integers, I should use bc -1

File compression

- There are many tools and hence formats
- gzip and gunzip are used for .gz files
- tar cfz and tar xfz are used for .tar
- zip and unzip are used for .zip

Network utilities

- Connect to a remote machine: ssh
- Copy remote files : scp
 - It is called secure copy
 - scp user@remotelocation.org:path/to/file /destination/path
- Download from the web: wget
 - It works with http and ftp urls

Globbing

- The Unix shell provides wildcards that can be used to specify filename patterns
 - — ★ matches any number of characters, also none
 - * echo * is equivalent to 1s
 - ? matches a single character
 - [abc] matches a, b or c
 - [!abc] matches not (a, b or c)
 - [a-z] matches any single letter
 - There are some special patterns like [:lower:] or [:digit:]
- I can specify more than 1 pattern in the same line
 - A* T* is equivalent to [AT]*
- Brace patterns can also match non-existing filenames
 - {A,B,C}{A,B,C} is expanded to all the 2 characters combinations of the 2 lists
 - It would be AA AB AC BA BB BC CA CB CC

Redirection

- In Unix devices (printers, screen output, ecc.) are treated as files
 - The stdout and stderr devices are connected to the monitor
 - stdin is connected to the keyboard
- stdout is redirected with >
- stderr is redirected with 2>
- I can append instead of overwrite with >> or 2>>
- I can redirect all the output with &>
 - Be careful, &>> does not work on all systems (!)
- The standard way to append all the output is to redirect stderr to stdout and then append it
 - I can use ls >> file.txt 2>&1
- I can trash an output by redirecting to /dev/null
- stdin can be redirected with <
 - It is almost useless, and it can not work with some commands
- Pipe (|) is used for redirecting the stdout of a command to the stdin of another
 - It is used for building pipelines (!)
- If I want to store an intermediate result in a pipeline, I use tee
 - input command1 | tee output1.txt | command2 > output2.txt

Text manipulation

- Concatenate and print to stdout: cat
 - cat file1 file2 > file3 creates file3 containing the concatenation of file1 and file2
- Print the first/last n lines: head -n and tail -n
 - head -4 myfile.txt prints the first 4 lines
 - I can print a specific line by piping head and tail
 - * head -4 myfile | tail -1
- Sort a file content: sort
 - The default sorting behaviour is lexicographic: 10 comes before 1
 - If I specify to sort according to a specific column I specify -k
 - columns are defined by whitespaces
 - sort -k 2 myfile sorts according to the second column
 - I can sort numerically with sort -n
 - I can remove duplicated lines with sort -u
 - * This works only if the lines are next to each other ofter sorting (!)
- Report or omit repeated lines: uniq

- It detects only adiacent duplicates (!)
 - * It is algortimically complex to detect unsorted duplicates
- uniq -d prints only duplicated lines
- Extract a column from a file: cut -f
 - -f specifies the field separator, that defaults to Tab
- Count stuff: wc
 - wc -1 counts the lines in a file
 - wc -c counts the bytes (carachters including the newline)
- Compare files line by line: diff
 - It prints the lines that are missing from the respective files
 - It is similar to the NW algorithm
- Find common lines in sorted files: comm

Regex and grep

- They are more powerful than globe patterns
- grep means global regular expression print
 - It searches in a file and prints all the lines that match the pattern
- egrep can handle extended regex
- fgrep is fast but does not deal with regex, it searches fixed patterns
- Regex searches for a string containing the expression
 - abc matches ANY string containing abc
- Standard regex
 - . matches any single carachter
 - * matches any number of times (also 0) the previous carachter
 - ^b matches anything that begins with b
 - [ABC] matches A, B or C
 - ^ and \$ match beginning and end
 - A{3,5} matches AAA, AAAA and AAAAA, A{2,} matches 2 or more A
 - \ turns of a metacarachter
- Extended regex
 - ? matches the preceding 0 or 1 time
 - + matches the preceding 1 or more times
 - (exp1|exp2) matches either exp1 or exp2
- I can invert the matching with grep -v
- I can search for words using grep -w

awk

- awk is a programming language designed for text processing
- The name comes from the initials of the authors
- The awk synthax can be used for text processing, arithmetic operations, string operations, and others
- It reads from STDIN or from a file
- The input is read as a set of records divided into fields
- A record is a line by default
- A field is a word by default
- The field and record separators can be changed, also to regex
- The specified operation is performed for every record, or to records that match a pattern
- The syntax is awk 'pattern {action}' [filename]
- If I use regex, it is awk '/regex/ {action} [filename]
- If action is not specified, it defaults to echo to STDOUT, essentially behaving like grep
- I can reference single fields of a record in the action
 - \$1,\$2 refer to the first and second field

- \$0 refers to the entire record
- I can specify multiple pattern with multiple actions
- There are optional blocks
 - BEGIN{action} and END{action} are executed before and after the parsing
- Variables can be built-ins or user-defined
 - Variables do not need to be declared
 - It is good practice to initialize them in the BEGIN block
- Some built-ins
 - NR is the number of the current record (line)
 - NF is the field number (word)
 - FS is the field separator
 - RS is the record separator
 - \$n represents the nth field
- I can change the field and record separators by assigning FS and RS (!)
- Some standard operators
 - Standard arithmetic operators: '+ * / % ++ -'
 - Relational operators: == != < >=
 - Logical operators: &&(AND) ||(OR) !(NOT)
- There are 2 regex operators
 - str ~ /regex/ returns true if str matches regex
 - str !~ /regex/ returns true if str does not match regex
- Some string operators
 - They can change with awk version (!)
 - length(str) returns the lenght
 - sub(regex,repl,str) replaces the regex in str with repl
 - substr(str,pos,len) extracts a substring
 - index(str,match) returns the index of match in str, if it exists
 - tolower(str) and toupper(str) convert cases
- The print function
 - print str1,str2 prints the strings separated by the built-in OFS
 - * If , is omitted the strings are printed without separator
 - OFS (output field separator) can be assigned and defaults to space
 - A newline is always added at the end of the print
- Commands can be separated in a single block by {cmd1; cmd2}
- getline is used to read the next line from input
 - It is like file.readline() in python
 - Once read, the line is not read again (!)
- printf takes in input a format string followed by a comma-separated list of arguments
 - printf means formatted printing
 - It is similar to printf() in C, since it ask was written by the same author of C (!)
 - I can put special symbols in my string, that refer to datatypes
 - * %d %f %s refer to int, float, string
 - I can then refer to the symbols after the print
 - * {printf "%d is an integer",1}
 - I can remove newlines by
 - * awk '{printf "%s ",\$0; getline; print \$0}
- Associatives arrays are basically dictionaries
 - awk '{array[key]=value}'
 - The key can also be in the form [key1,key2,keyn]
 - * In this case it just concatenates the keys as strings and forms a complex key
 - Non-existent values default to 0 (!)
- The if statement is done like
 - awk {if(test)codeblock;else codeblock}
- The for loop is done like in Perl

- awk {for(i=0,i<10,i++) codeblock}</pre>
- But I can also iterate on an array
 - awk {for(key in arr) print arr[key]}
- The while loop
 - awk {while(test) codeblock}