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# COMP2041/9044

Week 2 - Filtering

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# The core 7 filters

- sort
- wc
- tr
- head / tail
- cut
- uniq
- sed

# sort

- Sorts your input in **ascending line order**
- Lines can be sorted by a specific character within a field

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<b>-r</b>	sort in descending order (reverse sort)
<b>-n</b>	sort numerically rather than lexicographically
<b>-d</b>	dictionary order: ignore non-letters and non-digits
<b>-tc</b>	use character <i>c</i> to separate columns (default: non-blank to blank transition)
<b>-kn</b>	sort on column <i>n</i>

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```
sort -r <filename>
"tab"
1
10
11
12
13

zid|name|phone number

sort -k3 -t"|" < file
```

# WC

(word count)

- Summarises information about the input

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<b>-c</b>	print the number of <b>characters</b>
<b>-w</b>	print the number of <b>w</b> ords (non-white space) only
<b>-l</b>	print the number of <b>l</b> ines only

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```
WC -c < filename           <- Outputs count only
```

or

```
WC -c <filename>           <- Outputs count and file name
```

# tr

(transliterate)

- Replaces a character with another one
  - Can be a range of characters

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**-c** map all bytes *not* occurring in *sourceChars* (**c**omplement)  
**-s** squeeze adjacent repeated characters out (only copy the first)  
**-d** delete all characters in *sourceChars* (no *destChars*)

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```
tr 'abc' '123' < someText

# map all upper-case letters to lower-case equivalents
tr 'z-a' 'a-z' < text
tr '0-9' '9-0' < text

# remove all digits from input
tr -d '0-9' < text
```

# head / tail

- Selects the first / last **n** lines

```
head -n 30 filename      <-- Prints the first 30 lines of file
```

```
tail -n 30 filename      <-- Prints the last 30 lines of file
```

# cut

- Selects specific sections of an input using a “**delimiter**”.
  - Input should be in a certain format
  - Example: “field1 | field2 | field3”

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<b>-f</b> <i>listOfCols</i>	print only the specified fields (tab-separated) on output
<b>-c</b> <i>listOfPos</i>	print only chars in the specified positions
<b>-d</b> <i>c</i>	use character <i>c</i> as the field separator

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```
# print the first column (default delimiter is a tab)
cut -f1 data

# print the first three columns
cut -f1-3 data

# print the first three columns, if '|' -separated
cut -d'|' -f1-3 data

# print the first five chars on each line
cut -c1-5 data
```

# uniq

(unique)

- Removes all **adjacent** lines that are not unique

- 
- c also print number of times each line is duplicated
  - d only print (one copy of) duplicated lines
  - u only print lines that occur uniquely (once only)
- 

A  
B  
C  
D  
D  
C  
B  
B  
A

```
uniq filename      -> A B C D C B A
```

```
uniq -c filename   -> returns 1 A, 1 B, 1 C, 2 D, 1 C, 1 B, 1, A
```

```
uniq -d filename   -> returns D
```

```
uniq -u filename   -> returns A B C C B A
```



# sed

(stream editor)

- The most complicated filter you'll see this term 🤪

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**-n** do not print lines by default - applies all editing commands as normal but displays no output, unless p appended to edit command

**-E** extended regular expressions  
like grep you often want this

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<b>p</b>	print the current line
<b>d</b>	delete (don't print) the current line
<b>s/regex/replace/</b>	substitute first occurrence of string matching <b>regex</b> by <b>replace</b> string
<b>s/regex/replace/g</b>	substitute all occurrences of string matching <b>regex</b> by <b>replace</b> string
<b>q</b>	terminate execution of sed

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<i>line_number</i>	selects the specified line
<i>start_line_number,end_line_number</i>	selects all lines between specified line numbers
<i>/regex/</i>	selects all lines that match <b>regex</b>
<i>/regex1/,/regex2/</i>	selects all lines between lines matching regex1 and regex2

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# sed examples (from lectures)

```
# print all lines
sed -n 'p' < file

# print the first 10 lines
sed '10q' < file
sed -n '1,10p' < file

# print lines 81 to 100
sed -n '81,100p' < file

# print the last 10 lines of the file?
sed -n '$-10,$p' < file # does NOT work
```

```
# print only lines containing 'xyz'
sed -n '/xyz/p' < file

# print only lines NOT containing 'xyz'
sed '/xyz/d' < file

# show the passwd file, displaying only the
# lines from "root" up to "nobody" (i.e. system accounts)
sed -n '/^root/,/^nobody/p' /etc/passwd

# remove first column from ':'-separated file
sed 's/[^:]*://' datafile

# reverse the order of the first two columns
sed -E 's/([[:^:]]*)([[:^:]]*)(.*)$/\2:\1:\3/'

# replaces all occurrences of a,e,i,o,u with ""
sed 's/[aeiou]/g' story.txt
```

# Combine multiple filters

Using “|” aka the pipeline

```
# extract first field, sort, and tally  
cut -f1 data | sort | uniq -c
```

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# Tutorial Questions

Question 2-4, 6

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# More filters

The less commonly used filters

- join
- find

# join

- Allows you to merge lines from two files based on a common field, effectively combining related data.

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<b>-1 <i>k</i></b>	key field in first file is <i>k</i>
<b>-2 <i>k</i></b>	key field in second file is <i>k</i>
<b>-a <i>N</i></b>	print a line for each unpairable line in file <i>N</i> (1 or 2)
<b>-i</b>	ignore case
<b>-t <i>c</i></b>	tab character is <i>c</i>

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# find

- Find a file in your directory using regex
- More commonly used than using:
  - “ls | grep -E <regex>”
- Refer to manual to learn more about this

```
home
  week1
    lecture1
    lab1
    lecture2
  week2
    lecture3
    lab2
    lecture4
  week3
    lecture5
    lab3
    lecture6
```

```
[terence@Terences-MacBook-Air home % find . -name "lab*"]
./week1/lab1
./week2/lab2
./week3/lab3
[terence@Terences-MacBook-Air home % find * -name "lab*"]
week1/lab1
week2/lab2
week3/lab3
```

“.” current directory

“\*” expands to a list of everything in folder (not advised)

# Regular Expression Resources

Cheat Sheet: (Search “regex cheat sheet” and find the one by David Child)

<https://cheatography.com/davechild/cheat-sheets/regular-expressions/>

Online Regex Tester:

<https://regex101.com/>

! Disclaimer: This website uses a different method to run regular expressions, so there will be some cases where the output differs



# Tutorial “Slides”

<https://2041.terencelim.dev/26t1-w09a> (will be uploaded during the lab)



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# LAB Time!

Room: Ainsworth 302 (String Lab)

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