# Data Processing: Formats and Tools

(part 2) a topic in

DM565 – Formal Languages and Data Processing

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sort
uniq
tr
cut
paste
join

head/tail

These are all linux filters, i.e., they do not change their input, but produce output on **stdout**, and can be used in pipes, just like the more complicated grep, sed, and (g)awk.

### Command-Line Tools: sort spu

### **Options for common issues (selected)**

```
ignore blanks
ignore case
sort numerically, alphabetically, by month, version numbers, . . .
specify which field to sort on
specify delimiters
reverse
```

### **Example**

Sort numerically on the 5th column, showing the larger numbers (file sizes) first:

```
> Is -I | sort -n -r - k5
```

# Command-Line Tools: uniq spu -

"filter out adjacent matching lines" - often used after sort

### Options for common issues (selected)

```
ignore case
print only unique or duplicate lines
consider only the first or last some number of characters
consider only some fields
count duplicates
```

#### **Example**

Remove duplicates, ignoring the first field:

```
> cat myfile
41 1 2 3
42 1 2 3
43 3 2 1
> cat myfile | uniq - f1
41 1 2 3
43 3 2 1
>
```

### Command-Line Tools: tr spu ..

"translate or delete characters"

#### **Options for common issues (selected)**

delete characters in a given set delete consecutive duplicates of a character, leaving one occurrence translate by specifying two character sequences of the same length

#### **Example**

Change and delete some characters:

```
> cat myfile
41 Forty + one
42 Forty + two
43 Forty + three
> cat myfile | tr'F + ''f -' | tr -d' [: digit :]'
forty - one
forty - two
forty - three
>
```

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Command-Line Tools: Cut 5DU-5

# "remove sections from each line of files"

## **Options for common issues (selected)**

select numbered bytes
only keep certain characters
select some fields
specify delimiter

specify output delimiter

#### **Example**

Change input/output delimiters and keep columns 2 and 3:

```
> cat myfile
x :41: one
y :42: two
> cat myfile | cut -d : -- output - delimiter = ' ' -f2 ,3
41 one
42 two
>
```

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# Command-Line Tools: paste spu :-

"merge lines of files"

### Options for common issues (selected)

specify delimiter (default is \t)

serial mode (the lines in each file will be concatenated into one line)

#### **Example**

Paste lines using space instead of the default tab:

```
> cat myfile1
41
42
> cat myfile2
forty - one
forty - two
> paste -d ' ' myfile1 myfile2
41 forty - one
42 forty - two
>
```

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# Command-Line Tools: join spu ❖

"join lines of two files on a common (sorted) field" – similar to dbms equi-join Options for common issues (selected)

```
ignore case
specify delimiters
specify join field
```

### **Example**

Join lines on a common field (first field is default):

```
> cat myfile1
42 A
42 C
43 B
> cat myfile2
41 X
42 Y
> join myfile1 myfile2
42 A Y
42 C Y
```

"output the first/last part of files"

### Options for common issues (selected)

```
specify the number of lines
specify start line
specify bytes instead of lines
```

### **Example**

### Print the first two lines of the last 10 (default) lines:

```
> seq 50 | tail | head -2
41
42
```

#### stream-oriented, non-interactive, text editor

Specify patterns (similar to grep),

but change (edit) the matching lines,

not interactively, but via a script – a sequence of commands.

Changes are applied to a line successively, i.e., after one modification, the next change (to the same line) is applied to the modified line.

A command consists of *address* information and an *action*; the address information can restrict the lines affected to some subset.

## Command-Line Tools: sed sou -

Using standard specification syntax, a command has the form

### [address[,address]][!]command[arguments]

(Unfortunately, syntax varies a lot; in manuals and similar documents, [...] is used instead of (...)?, i.e., zero or one occurrence.)

An address can be a *line number* (\$ can be used to mean "the last line") or a *pattern*, which is simply a regular expression (grep-style) surrounded by slashes.

Two addresses can be used to specify an interval and ! negates the address information.

# Command-Line Tools: sed SDU &

#### **Examples: On Address Specification**

- d delete all lines
- 42d delete line 42
- 1,10!d delete all lines except lines 1-10
- 1,/^\$/d delete from line 1 through first blank line
- /**^\$/,\$d** delete from the first blank line through the last line **/42/,42d** delete from the first line containing the number 42 through line 42

/^Proof/,/qed\$/d

delete from the first line starting with "Proof"

### through the first line ending with "qed"

So, on command-line, one writes, for instance,

cat myfile | sed ' 42! d '

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# Command-Line Tools: sed spu ₽

### **Useful Options**

A filename can be given as argument to **sed**; if omitted, input comes from **stdin**.

- -n suppress output, unless an explicit print command is issued (see later) -f the next argument is a file name containing a script
- -e the next argument is a command

(if necessary to avoid confusion with a file name argument

-E use regular expression syntax as for grep -E

#n as the first line of a script is an alternative to -n

The print command **p** is used exactly like **d**.

Unless -n is used, printed lines will come out twice.

However, **p** together with **-n** can be useful when

there are several commands. Kim Skak Larsen (IMADA) DM565

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### Command-Line Tools: sed spu -

#### **Substitute**

The substitute command  ${\bf s}$  takes arguments and optionally flags. Leaving out the address specification, the syntax is

### s/pattern/replacement/[flags]

A flag can be a number i, indicating that it is the ith occurrence that should be replaced. The flag g (global) indicates that all occurrences should be replaced, and p prints.

The pattern is just a regular expression (grep-style).

The replacement string can contain special characters:

\d the dth group from the pattern (grep-style)

& the entire string matched by the regular expression

\\ backslash

**\&** ampersand

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#### **Examples**

We can fix a spelling error by sed -E 's/iether/either/g'

We can translate the second occurrence of "datalogi" on each line by sed -E 's/datalogi/computer science/2'

We can change "datalogi" to "datalogistudiet" by sed -E 's/datalogi/&studiet/g'

If a file contains two columns, separated by one colon, we can switch the two columns using sed -E 's/(.\*):(.\*)/\2:\1/'

Regular expressions match strings as long as

possible, left to right. Kim Skak Larsen (IMADA) DM565 topic: Data

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## Command-Line Tools: sed SDU &

#### **Transform**

The sed-equivalent of **tr** is the command **y**, performing one-to-one character-to-character replacement (can be prefixed by addresses).

#### **Example**

sed 'y/123/234/' will increment all the digits 1, 2, and 3.

#### Quit

The quit command **q** stops processing when (if) the single address specification is reached.

#### **Example**

sed '10q' will terminate after the first 10 lines

have been processed. Kim Skak Larsen (IMADA) DM565 topic: Data

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# Command-Line Tools: sed

#### Other Commands

There also command for inserting and appending before or after a line, changing lines in a fixed manner, and more, but the syntax becomes more cumbersome and it might be nicer to use other tools.

### **Multiple Commands**

**sed** is most convenient for simple changes, possibly by piping into another **sed**. However, multiple commands are possible, with the following somewhat odd requirements.

#### Format:

```
[ address [ , address ]][!]{
command [ arguments ]

command [ arguments ]
}
```

where the opening brace must be last on a line and the closing alone on a line. Kim Skak Larsen (IMADA) DM565

# Command-Line Tools: awk

Aho, Weinberger, Kernighan

We will use gawk (GNU awk), but usually just say "awk".

a full programming language
can be used on command-line or via script
handles fields nicely (not just lines)
understands numbers (not just text)
C-like syntax, but also grep-like patterns
an awk script is a sequence of pattern {action}

# Command-Line Tools: awk I

#### **Patterns**

**BEGIN** and **END** are special patterns that only match at the begining or end of a file, respectively, used for initialization and announcement of results regular expressions enclosed in *I...I* 

C-like conditionals with comparisons and logical connectives: **&&**, ||, ! **s** ~ **r** is true if the string **s** 

matches the regular expression  ${\bf r}$  (asymmetric!); use double quotes around the arguments if they contain special symbols arithmetic and built-in mathematical functions

#### **Actions**

C-like statements if there is no action, lines matching the pattern

are printed (sed-style) Kim Skak Larsen (IMADA) DM565 topic: Data

### Command-Line Tools: awk

### Example

```
Is | gawk '
BEGIN { print " List of tex files :" ; count = 0}

\( \lambda \text{ tex } / \{ \text{ print ; count } += 1 \}
END \{ \text{ print " Total :" , count , " files " }
\( \lambda \text{ text} \)
```

### gives output (in my example)

```
List of tex files:
2023 lecture . tex
def - colors . tex
lecture . tex
preamble . tex
Total : 4 files
```

Alternatively, one can print with C's printf.

## Command-Line Tools: awk

### **Field Manipulation**

**RS** is the record separator (default newline)

**FS** is the field separator (default other maximal whitespace sequences) **OFS** is the output field separator (default space)

**NR** is the number of the current record (line)

NF is the number of fields in the current record

\$1, \$2, ..., \$0 are the fields and the entire record

### **Examples**

cat myfile | gawk '{ print NR, \$2 \* \$3, \$(NF-2)}' prints the line number, the value of fields 2 and 3 multiplied together, and the third to last field.

```
We can assign to the field variables as in cat myfile | gawk '{ $1 = $2; $2 = ""; print }'
```

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### Command-Line Tools: awk

#### **Built-In Functions**

The linux command **wc** can be realized as follows:

```
gawk '
BEGIN { OFS = "\ t " }
{ chars += length ($0 ) + 1; words = words + NF }
END { print NR , words , chars , FILENAME }
' myfile
```

+1 since \$0 does not include newline.

Variables are initialized to the empty string or zero as

string concatenation – placing strings next to each other separated by blanks **substr(s, m, n)** – **n** characters from position **m** in **s** advanced **split** operations **system** calls and **exit** 

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Command-Line Tools: awk

dictionaries, including ARGV (ARGC for the length of ARGV)

#### **Control Structures**

appropriate. Also

if-then-else

while and do-while

for-loops C-style and for (key in array) { ... }

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# Command-Line Tools: awk

### **Example**

Print input in reverse order:

```
cat myfile | gawk '
  { line [ NR ] = $0 }
   END {
      for (i = NR; i > 0; i = 1)
        print line [ i ]
```

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### **Useful Options**

Filenames can be given as argument to **gawk**; if omitted, input comes from stdin.

**-f** the next argument is a file name containing the program **-F** the next argument is to be used as input field separator -v var=val initialize a variable prior to execution

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# JSON-Like Formats: JSON

JavaScript Object Notation

JSO N

JSON-Like Formats: XML

eXtensible Markup Language

L

```
< animals >
  < animal >
  < name > Panda </ name >
  < cuteness > 1.0 </ cuteness >
  < color > white </ color >
  < color > black </ color >
  </ animal >
  < name > Panther </ name >
  < cuteness > 0.7 </ cuteness >
  < color > black </ color >
  </ animal >
  < name > Panther </ name >
  < cuteness > 0.7 </ cuteness >
  </ animal >
  </ animal >
  </ animals >
```

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# JSON-Like Formats

There is more to both formats.

The essence is that it is named parentheses structures expressing records (attribute/value pairs) and sequences (arrays, lists).

There are many variants of XML (HTML) with similar structure. Command-Line tools can to some extent be used for data discovery, and possibly simple code execution.

To get full power, use a programming language with an appropriate package. Packages read json/xml files and deliver data in native formats.

# JSON-Like Formats: Python Example \_\_\_\_\_

```
# Prints
# " d ": 4 ,
# " e ": 5
# Panda
import json
# Data in program for testing
json_data = ' {" c ": 3 , " d ": 4 , " a ": 1 , " b ": 2 , " e ": 5} '
parsed ison = ison . loads ( ison data )
print ( json . dumps ( parsed json , indent =4 , sort keys = True ))
# It is just dictionaries and lists
with open ('animals.json','r') as f:
      animals dict = json . load (f)
print ( animals_dict [ " animals " ][0][ " name " ])
```

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