Awk Essential Training: Lynda.com

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Awk: named after inventors: simple data filtering and manipulation.

Good for: text files divided into lines, with fields on each line; joining, merging, etc...

Versions: awk, nawk, gawk—> the one we get if we type gawk on command line. (gawk was installed by "brew install". Now your awk is automatically gawk, but it wasn't by default)

- 1. awk '{print \$2, \$1}' names.txt
  - 1. all awk programs start with "awk"
  - 2. the part within single quote is the actual
  - 3. names.txt is the input file
  - 4. {} means apply this command to every **line** in input
  - 5. \$2 is the second field in line, \$1 is the first field in line
  - 6. comma: in print: use a field separator
  - 7. awk 'NR==3' <input>: prints the third record (no need to use the command 'print')
  - 8. command substitution can be understood in awk. However, all awk arguments have to be enclosed in { }. The arguments inside the first { } will not be interpreted as awk arguments, so please don't do that.
- 2. awk looks at each line of text, each field separated by a space
  - 1. \$0 refers to the whole line
  - 2. print \$2 "," \$1 —> concatenation
  - 3. print NF —> print Number of Fields
    - Caution: while NR refers to record number, NF refers to number of fields; it is a numeric variable. To call a particular field, use \$<field\_number>.
  - 4. awk '/up/{print NF, \$0}' name.txt
    - 1. /up/ : a pattern. The action within {} is only applied to lines with this pattern
    - 2. other patterns: NF==6, etc.
    - 3. awk 'NF==6' name.txt: print lines matching this patterN
  - 5. awk '/up/{print "UP:", NF, \$0} /down/{print "DOWN:", NF, \$0}' name.txt: multiple actions in awk
- 3. awk flags:
  - 1. -f: the following program is to be applied to a file
    - 1. exp: awk -f swap.sh name.txt
    - 2. inside swap: only {print \$2, \$1}: no single quotes; no need to be protected from shell

- 2. -F: use what as field delimitor
  - 1. exp: awk -F , '{...}' name.txt
  - 2. -F t : t == tab
- 3. -v: specify the value of a variable
  - 1. exp awk -v hi=Hello '{print \$1, hi}' then type in "hello awk", what to expect is "hello, Hello" (use ctrl+x) to end the program
  - 2. **Caution:** use -v to pass external variables to awk! use -v once for each variable.
- 4. files: can specify multiple file; awk will process one by one
- 5. The input and output can be redirected awk '..... ' <input> > <output>
- 6. Defining record and field:
  - 1. By default: field: any combinations of spaces and tabs
  - 2. -F exp: awk -F , '{print \$2}' <input>
  - 3. other field separators: ABC; "a b" (multi-character separators are not supported in MacOSX...); "" (empty string) —> **Caution:** blank line as RS, no separator (every character is separated) as FS; " "(space) —> space; '[,!]' —> , or ! (single quotes: protect from shell); or just a blank line as RS: "\n\n+".
  - 4. specify within an awk program: with special variable FS
    - 1. exp: awk '{FS=","; print \$2}' <input> semicolon: specify the end of one command
    - 2. caution: awk separate input into fields and records before calling the commands.
    - 3. use BEGIN pattern to let the commands execute first: awk 'BEGIN{FS=","} {print \$2}' <input>
  - 5. specify within an awk program record separator: RS
    - 1. exp: 'BEGIN{RS="!";FS=","} {print \$2}' <input>
  - 6. output field/record separators: OFS ORS
    - exp awk 'BEGIN{RS="\n"; FS=" "; OFS=","; ORS="!"} {print \$2, \$1}' names.txt
  - 7. to specify a range of records: NR==3, NR==8{print}. No quotes. NR==3(or 8) can be any regular expression; they specify the beginning and end.
  - 8. To loop through the records: specify the range of records at the beginning of { }; all commands in { } will be looped through them automatically. If no specification, all records will be looped through. BEGIN{ } happens before looping starts. The END{ } do not loop through any record; commands are just executed.
    - 1. If for loops inside a { } appears to be only applicable to the last record, you have used the wrong syntax...the for syntax is: for (initialization; condition; increment) (new line) body, no { ]!!

## 7. More built-in variables:

- 1. NR: record number exp: use NR==6 as a pattern to extract the sixth records
- 2. FILENAME: filename of file awk is processing
- 3. FNR: record number within that file
- 4. \$0: the whole line
- 5. \$#: the #the field in that record
- 6. \$NF: the last field
- 7. \$(NF-1): next to last field
- 8. as long as \$(a\_number), it specifies a field
- 9. field can also be changed (only in memory; not in input) exp: \$2="hi"
- 10. any field can be specified: \$100="hi", even if original NF=5
- 8. Creating user-defined variables
  - 1. **Caution:** in awk, the variables within the same awk program are valid across { }, even if it is an index in for-statement, etc. Avoid using repetitive indices!
  - 2. In awk, it is unnecessary (and wrong) to do put \$ in front of every variable. \$ is only used to let bash knows that something is a variable; however, when the variable is inside awk, it will be interpreted correctly as a variable, because bash knows awk.
  - 3. exp. awk '{hello=\$1; goodbye=\$2; print hello, goodbye}'
  - 4. all awk inputs are case-sensitive
  - 5. variable type: inter-converted based on context
    - 1. exp: awk ' $\{a=1; b=3; print a/b\}' \longrightarrow 0.333333$
  - 6. awk multiplies first then concatenates
  - 7. parentheses used to specify what happens first
- 9. all awk variables are global variables awk 'BEGIN{a=1} {print a}'
- 10. Math operators:  $+,-,*,/,%,^{\wedge}$  (not exactly the same as bash...)
- 11. increment/decrement operators:
  - 1. a=3; b=++a; print a,b  $\longrightarrow$  4.4 (the value of a is also changed...)
- 12. assignment operators: =, +=, -=, \*=, /=, %=, ^=
- 13. comparison operators: ==, !=, <, <=, >, >= (1 for true, 0 for false...different from bash)
- 14. string operators: concatenation: space. ~,!~ —> whether string value matches regular expression. for null, however, the most convenient way is to use!="" or =="".
- 15. Arrays: [] (different from bash...)
  - 1. awk '{a[1]=\$1; a[2]=\$2; a[3]=\$3; print a[1], a[2], a[3]}'
  - 2. awk only allows 1-dimensional arrays
- 16. Regular expressions

- 1. format: // exp: /abc/; matches "abc", "mabc", etc.
  - 1. watch spaces! no space within / / unless otherwise wanted
- 2. comparison: ~ and !~ : whether something matches a regular expression
  - 1. exp awk '\$4~/up/{print}' <input>
- 3. special patterns for regular expressions:
  - 1. .: matches any single character exp: /a.c/
  - 2. \: escapes . ; \ ; / exp: /a\.c/ matches "a.c"
  - 3. ^: matches the beginning of; \$: matches the end of
    - 1. exp: /^abc/ matches "abcd"
    - 2. exp /abc\$/ matches "dabc"
  - 4. [] matches any character in set
    - 1. exp: /a[xyz]c/ matches "axc"
    - 2. exp: ranges: /a[a-zA-Z]c/ matches "amc"
    - 3. exp: not: /a[^a-z]c/ does not match "abc" but matches "a1c"
  - 5. \*: matches 0 or more occurrences of the previous item
    - 1. exp: /ab\*c/ matches "abbbbbc"
  - 6. +: matches 1 or more occurrences of the previous item
    - 1. exp: /ab+c/ does not match "ac" but "abc"
  - 7. ?: matches 0 or 1 occurrence of the previous item; i.e. the previous item is optional
  - 8. { }: certain number of repeats
    - 1. exp: /ab{3}c/ matches only "abbbc"
    - 2. exp: /ab{3,}c/: matches 3 or more repeats of b
    - 3. exp: /ab{3,4}c/ matches "abbbc" and "abbbbc"
  - 9. (): specify a group to repeat
    - 1. exp: /(ab)+c/ matches "ababc"
- 4. **greediness**: regular expression will match as many items as it can
  - 1. exp: /<.+>/ will match all of "<hi>>....<hi>"
  - 2. to only get the first "<hi>": /<[^>]+>/
- 17. Control Structures: similar to C. Awk is a full-fledged programming language.
- 18. if statement:
  - 1. if (condition) { (new line) if-statements (new line) } else{(new line) elsestatements (new line)}
  - 2. condition: == : numeric equality; ~: regular expressions matching
  - 3. numeric values are considered false if its value is 0; true if its value is something else
  - 4. string values are considered false if it's empty string; true otherwise
  - 5. use semicolon at the end of **each statement** in if-statements, not in conditions
  - 6. the spaces and line breaks are optional; for awk program, any number of breaks and spaces can be used around any element (except assignment)

- 7. exp: awk '{if (NF < 8) {print "hi"} else {print "good"}}'
- 8. in an awk file:
  - 1. {if (NF < 8) {print "hi"} else {print "good"}}
  - 2. execute the file awk -f 1.awk <input>
- 9. in a shell file:
  - 1. exp: awk '{if (NF < 8) {print "hi"} else {print "good"}}'
  - 2. execute: sh 1.sh
  - 3. or: chmod +x 1.sh then: ./1.sh

## 19. for statement:

- 1. {for (initialization (=, not ==!); condition; increment) {(new line) body}. The { } is optional for single-line statements but always a good idea to use. (Since unlike bash, awk does not have do...done pair to enclose a statement.)
  - 1. **Caution:** all awk statements need to be enclosed in { }, even for, while, if, etc.!
  - 2. If you want to put a for loop inside a while loop, be careful! The for loop will finish as long as the initial condition for the while loop is true!
- 20. BEGIN: at the beginning of the program, do this this and that; END: at the end of the program, do this this and that.
  - 1. exp: BEGIN{ beginning of file) } { (main body) } END{ (end of file)}
  - 2. END is **very** important! if something is written in END, it will not be looped through every line. if it is in { } with no patterns ahead, it will be looped through every line of input!
- 21. printf(): same syntax as C
  - 1. printf (format, value...)
  - 2. exp: awk -F '{printf("%s\t%s\t%d\n", \$1,\$2,\$3}
  - 3. %#s: the field is # length %-#s: left justified
  - 4. %f: six digits after decimal point by default. %#1.#2f: #1: width (including decimal; #2: #digits after decimal
  - 5. %d: integer

## 22. String manipulation:

- 1. the first letter is letter 1, not 0
- 2. string functions: none of these is going to alter the input file
  - 1. length([string])
  - 2. index(string,target): return the **first** find of target in string; indexed by the first letter of target
  - 3. match(string,regexp==regular expression): looks for the first find like index; returns index; also sets RSTART to index, RLENGTH to length of the match
  - 4. substr(string, start, [,length]): looks for the substring starting at start with length length. If length is not specified, it's the substring starting

- at start till the end.
- 5. sub(regexp, newval[, string]): replaces the **first** match of regexp with newval
- 6. gsub(regexp, newval[, string]):replaces **all** matches of regexp with newval
- 7. split(string, array[, regexp]): break fields into **subfields** and store in array; default: regexp = FS.

## 23. Arrays

- 1. associative arrays: a["hi"]=1, a["good"]=2...etc. To iterate through, use for (index in array) {body}
- 2. sort can be used to sort arrays. exp: <output> | sort -rn -k 2
  - 1. -rn: reverse, numerical
  - 2. -k: key is field 2 of output
  - 3. in awk program: format: asort(a, a, "@val\_num\_desc"). See gnu. The second a is destination.
- 3. simulate a multi-dimensional arrays
  - 1.  $\exp \arctan [i "," j] = 2, etc.$
- 4. delete (see gnu)
- 5. **Caution:** watch the indices!!! tempDistance[,2] is not a valid index!
- 24. Math functions:
  - 1. int(x): integer part of x
  - 2. rand() 0-1; never 1
  - 3. srand([x]): seed random
  - 4. sqrt(x)
  - 5. sin(x), cos(x)
  - 6. atan2(y,x): atan2(0,-1) = Pi
  - 7. log(x)
  - 8. exp(x)
- 25. Pipes: pipe the output of a command as the input of another command 1. can be used to preprocess the input, etc.
- 26. CSV file: refer to the course
- 27. Awk scripts: use #!/bin/bash then type awk -Ft 'BEGIN {...} NR > 1{#Skip header.....} END { ....} > <output.txt> use chmod +x <file> to make it executable