

Handling Data using Regular Expressions Lecture #4 - COMP 364 January 11, 2010, updated 2012 Derek Ruths

Pipelines

- The | operator will take the output of a command and send it to another command
 - curl http://en.wikipedia.org/wiki/Pipeline_(Unix) | head -n 10
 - cat /usr/share/dict/words | less
 - cat *.fasta | grep AAA
 - tail and head can be combined together with a pipeline!

Regular Expressions: Motivation

- Create a data set from a subset of a data file
 - Extract protein interactions for one organism from the STRING database
- Count specific items in a data set
 - How many genes code for ribosomal proteins in the human genome?
- Extract data from a file
 - Get all the gene locations from an NCBI genome file

All of these tasks involve selecting a subset of entries from a larger (textual) data set.

Regular expressions

- Purpose: precisely define a class of words/character sequences that have some parts that "look the same"
 - Words containing the string "ba"
 - Words that start with a capital letter
 - Character sequences containing only the characters A, C, T, and G
- We have already seen simple regular expressions: wildcards (* and ?)

egrep: selecting lines from files

- Select all lines containing a specific regular expression:
 - egrep "<regular expression>" <file to select lines from>
- Count the number of lines containing a specific regular expression:
 - egrep -c "<regular expression>" <file to select lines from>

- egrep -c "[[:alnum]:]*[ba][[:alnum:]]*" words.txt all words containing "ba"
- egrep "[ACGT]{5,}" e_coli.txt all sequences of ACTG longer than 5 characters

The process of writing a regular expression

- The process has three steps:
- 1. Knowing what it is you want to match and how it might appear in the text
- 2. Writing a pattern to describe what you want to match.
- 3. Testing the pattern to see what it matches

Regular Expressions (in grep)

- Specifying a specific string:
 - egrep "hello" foo.txt find all lines containing the word "hello"
- Specifying a variable position: "." and bracketed expressions ("[...]")
 - egrep "hs00." foo.txt find all lines containing the string "hs00<anything>"
 - egrep "hs00[0123456789]" foo.txt find all lines containing the string "hs00#"
 - egrep "hs00[0-9]" foo.txt shorter way of writing the above
 - egrep "hs00[^0-9]" foo.txt find all lines containing the string "hs00<anything but a number>"
- Exercises
 - Find all lines in foo.txt containing the string "hs00<alphabetical character>"
 - Find all lines in foo.txt containing the string "hs00<alphanumeric characters>"

Character classes

- Specified by the brackets [], a character class is a regular expression that matches exactly one character, and the possibilities for that character are specified in the brackets.
- For example, to match "What" and "what", a good regular expression would be "[Ww]hat", which says that the first character can be either 'W' or 'w' (but not both!) and that the following characters must be exactly "hat"
- Other examples: hs00[0-9], [ACGT]+, COMP[34][56]4

Backslash magic

- "." means "anything"... how do we specify that we want a period?
- "[" is the beginning of a variable position... how do we specify that we want a left square brace?
- "-" indicates a range of characters... how do we specify that we want a dash?

When in doubt, backslash the character!

Repetition operators

- ? the preceding item is optional and matched at most once
- * the preceding item will be matched zero or more times
- + the preceding item will be matched one or more times
- {n} the preceding item is matched exactly n times
- {n,} the preceding item is matched n or more times
- {n,m} the preceding item is matched at least n times, but not more than m times.

Regular expressions: specifying longer variable regions

- What if I wanted to find all sequences in which there were several variable positions?
 - Find all lines containing an email address:
 - egrep "[A-Za-z0-9.-]+@[A-Za-z0-9.-]+\.[A-Za-z]+" emails.txt
 - Find all lines containing DNA sequences longer than 5 nucleotides:
 - egrep "[ACTG]{6,}" genome.txt

Grouping regular expressions

- Parentheses group expressions: repetition operators can act on these groups
 - $(TA)+A{3,} = TATA box$
- The "|" character indicates an "or" either expression can match
 - N[^P](S|T)[^P] = N-glycosylation site motif

Exercises

- Write a regular expression for each of the following:
 - A telephone number
 - A telephone number with optional dashes
 - A telephone number with an optional extension
 - A sequence of DNA containing an exon (ensure that the coding component has a correct coding region)

Useful shortcuts

- [:digit:] = 0-9
- [:alnum:] = A-Za-z0-9
- [:alpha:] = A-Za-z
- [:blank:] = tab or space
- [:punct:] = punctuation symbols
- [:space:] = any whitespace
- [:graph:] = anything EXCEPT whitespace
- [:upper:] = A-Z
- [:lower:] = a-z

Exercises

- A telephone number
- A UNIX path
- The scientific name of an organism

Anchors

- ^ = the beginning of a line
- \$ = the end of a line

- What do these regular expressions correspond to?
 - ^(Hello|Greetings) [[:upper:]][[:lower:]]+!
 - ^(100|[1-9][[:digit:]])[[:space:]](T|F)[[:space:]][[:alpha:]]+\$