Regular Expressions

Regular Expressions

- 3 types of regular expressions
- Match
- Does the variable contain
 Substitute
- Transliterate

If match, then replace with something

Swap characters

Match - Syntax

- Lets break down the match
- if(){
- This is the block form of if, we'll look at that more in the next section. For now, accept that it is just allowing us to 'visualise' the match

Alms

- Understand what a regular expression is
- Use match, substitute and transliterate
- Know about using i,m,g flags
- Understand some metacharacters
- Capture
- Greedy and non-greedy match
- Count with transliterate

Match

- Problem: I want to see it a sequence contains the codon for Methionine
- So far, we could look to see if there is an index returned from

say index \$seq, 'ATG'

However, that isn't very obvious, and what if the sequence is lower case? bin/01-match.pl

Match - Syntax

\$sequence =~ m/ATG/i

- This is the syntax for a match
- \$sequence
- Our variable to act upon
 The 'true' match operator. The return value of the expression is a boolean, so this sets

(No match to be true would use !~)

match as boolean true, fail as boolean false

Regular Expressions

- Perl's speciality
- Heavily optimised to be as efficient as possible
- Reliable
- Fully featured (as yet, I haven't found anything it is missing)

Match

my \$sequence =
'CCGGATCACTATGACCTGCTTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';
say \$sequence;

if (\$sequence =~ m/ATG/i) {
 say 'Sequence contains Methionine';

 This tests to see if it can find ATG at all in the sequence, and lets us know if true

Match - Syntax

\$sequence =~ m/ATG/i

m/ATG/i

- m tells perl we want to do a match regular expression operation
- /xxx/ delimits the thing we want to match
- i is a flag options

Match

 Obviously, we can just swap ATG for whatever we want to search for

}

Hello World 01223834244

There are also a set of metacharacters we can use, to match out of a set.

Match - Metacharacters

my \$\forall yrics = 'I see a little silhoutte of a man';
if (\$\forall yrics =~ m\s/) {
 say 'Lyrics contain whitespace';
}

- \s matches a whitespace character
- Space
- Line break
- lab

Match - Capture

- How about capturing the first word:
 my (\$first_word) = \$lyrics =~ m/(w+)\s+/;
 say \$first_word;
- Or the last:

my ($\$last_word$) = $\$lyrics = \sim m/s(lw+)ls*lz/m$; say $\$last_word$;

* means zero or more

Match - Metacharacters

```
if ( $sequence =~ m/d+/ ) {
    say 'Sequence contain a number';
}
```

- \d matches a digit
- + matches 1 or more of the preceeding character or metacharacter

```
if ( $sequence !~ m∧d+/ ) {
```

say 'Sequence does not contain a number';

Match - Metacharacters

These are the most common metacharacters you will see along with A – match the start of the string Z – match the end of the string
. - match any character

Match - Greedy

- * means zero or more, + means 1 or more
- However, they like to grab more than less

my \$sequence =
'CCGGATCACTATGACCTGCTTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';
my (\$greedy_match) = \$sequence =~ /(w+ACC)/;

How do we stop this?

say \$greedy_match;

Match - Metacharacters

```
if ( $sequence =~ m/w+/ ) {
    say 'Sequence contains word characters';
}
```

- \w matches a word character
- Alpha
- Underscore
- Digits

Match - Capture

- As well as matching, you can capture part or all of what you are trying to match, by placing brackets around the bit to capture my (\$silhoutte) = \$lyrics =~ m/s(silhoutte)\s/i; say \$silhoutte;
- In this case, if we match the word silhoutte, surrounded by whitespace, then put that word into the variable

Match - Non-Greedy

Use the ? metacharacter

my \$sequence =
'CCGGATCACTATGACCTGCTTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';

my (\$non_greedy_match) = \$sequence =~ /(w+? ACC)/;

say \$non_greedy_match;

 ? tells the regex to stop matching after the first it can match

Match - User defined

 Our user wants to see if their favourite codon is present

my \$sequence =
'CCGGATCACTATGACCTGCTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';

my \$user_wants_to_find_this_sequence = "tt";

How?

Match – i and m

- m makes perl match newlines/end of strings the same as other languages
- In most languages:
- ^ matches the start of a line, but in perl the start of the string
- \$ matches the end of a line, but in perl the end of a string
- This makes it a little confusing, so m changes their meaning
- Example at end of bin/01-match.pl

Substitute - Syntax

Lets look at the syntax, comparing with

\$sequence =~ s/cac/cat/i;

\$sequence =~ m/cat/i

=~ That's the same. We are informing perl on \$sequence that we wish to perform a regex operation

Match – User defined

We can put their variable in the match

if (\$sequence =~ /
\$user_wants_to_find_this_sequence/i) {

say "User sequence \$user_wants_to_find_this_sequence found.";

The regular expression is allowed to contain variables.

Much more useful than forcing only to match developer designated expression

Substitute

another in the sequence. Problem: We want to replace a codon with

bin/02-substitute.pl

my \$sequence =
'CCGGATCACTATGACCTGCTTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';

say 'we do not have a cat'

unless (\$sequence =~ m/cat/i) {

Substitute - Syntax

\$sequence =~ s/cac/cat/i; \$sequence =~ m/cat/i

- s instead of m
- We tell perl that we want to do a substitution type of regex
- i as before we want to do case insensitive

Match – i and m

- What do the i and m mean?
- They are flags which tell the regex to work completeness, not because you need to in particular ways (note, these are here for
- i this tells the regex to ignore case

Substitute

Lets replace cac with cat

\$sequence =~ s/cac/cat/i;

if (\$sequence =~ m/cat/i) { say 'we now have a cat'

How many 'cat's?

say \$sequence;

1, but there were 3 'cac's

Substitute - Syntax

\$sequence =~ s/cac/cat/i; \$sequence =~ m/cat/i

- /match/replace/ instead of /match/
- match is identical, and you can use all the same matching tech
- But, perl needs to know what to replace the match with, so we put that in a second 'column'

Substitute 'g'lobally

- How many 'cat's?
- say \$sequence;
- but thorough
- 1, but there were 3 'cac's
- It only substituted the first one it found!
- How do we replace all of them? \$sequence =~ s/cac/cat/gi;
- say \$sequence;
- The g flag tells perl to keep matching and substituting until the end

Substitute – User defined

You can have variables as the match, and the replacement

The return value is the number of

Substitute - How many

Substitutions made?

substitutions made, or undef if 0

say \$sequence =~ s/aaa/cac/i; # 1 (we didn't specify

- my \$favourite_codon = 'aaa'; # it's all the first letter of my name;
- my \$least_favourite_codon = 'cat'; # I don't actually like cats
- \$sequence =~ s/\$least_favourite_codon/
 \$favourite_codon/gi;
- say \$sequence;

This means you can use it as a boolean

say \$sequence =~ s/cat/aaa/gi; # undef, we had already cleared away all the cats

say \$sequence =~ s/aaa/cac/gi; # 2 (the remaining

Substitute – Don't be greedy!

- Because the match is exactly the same as with a match regex, + and * are still greedy my \$clone = \$sequence;
- \$sequence =~ s/g.*cac/aaa/i;
- say \$sequence;
- \$clone =~ s/g.*?cac/aaa/i;
- say \$clon
- You would not have been able to do multiple substitutions, as the first match ate everything between the first g and last cac

Match and Substitute

- There are other metacharacters, and ways of constructing really powerful short regexes with other flags
- However, everything you have seen here will form the basis of any match you do, and you should be able to construct virtually any match you want
- 90% of my regexes use nothing more than shown here

Transliterate - Syntax

- Let's look at the syntax
- reverse_strand =~ tr/ACGT/TGCA/;
- It's quite similar to substitute, but there are important differences

my Ssequence =
'CCGGATCACTATGACCTGCTTTCGCACCTGCTCG
CGCCGTCACGCTCGCAGTC';

bin/03-transliterate.pl

Iransliterate

- =~ We now know what this means
- tr tells perl to perform a transliterate operation on *\$reverse_strand*

In-place substitution of the dna letters

say \$reverse_strand;

say \$sequence;

my \$reverse_strand = \$sequence; \$reverse_strand =~ tr/ACGT/TGCA/

Transliterate

- The 3rd type of regex is transliterate. It stands a little aside from match and substitute, as it is more literal (i.e. less metacharacter and flag driven)
- Problem: I want to know the reverse strand of my DNA sequence
- You can't use substitute easily
- s/T/A/gi
- s/A/T/gi

Transliterate - Syntax

reverse_strand =~ tr/ACGT/TGCA/;

- /ACGT/TGCA/
- This is the important bit. Whereas before we had match 'patterns' and substitution strings, we don't here.
- As it reads \$reverse_strand, each time it finds a letter in the first part, it replaces with the equivalent letter in the second
- Find an A, replace with T
- Find a G, replace with a C

Transliterate – Return Value

 As with substitute telling you the number of substitutions made, transliterate returns the number of transliterations made (or undef)

\$reverse_strand = \$sequence;

say \$bees =~ tr/ACGT/TGCA/;

Transliterate – GC percentage

With a bit of playing, you could work out the percentage GC content

say "percentage GC = "

((\$sequence =~ tr/CGcg/CGcg/) * 100 / (\$sequence =~ tr/ACGTacgt/ACGTacgt/));

say \$sequence;

Again, unchanged, but we know what the GC percentage is

Transliterate - Coding a Message

 A bit of fun, lets use this to apply a simple cypher

my \$original_message = 'The quick brown fox jumped over the lazy dog.';

say \$original_message;

my \$coded_message = \$original_message; \$coded_message =~ tr/a-zA-Z ./N-ZA-Mn-za-m. /;

say \$coded_message;

In regexes, we can shorten sequences with a dash, they get expanded by the parser.

Quick play with regexes

 Lets just spend a few minutes playing with regexes.

Produce a regex which will capture a name and a phone number into variables from the following:

Andy Brown: 01234 567890

- Bonus points for
- Using metacharacters
- name parts and phone parts

Transliterate – Counting Characters

You can replace a character with itself, to get a count of that character in the string, without changing the string

say \$sequence;

say "The number of A's in above = " . (sequence = tr/Aa/Aa/);

say \$sequence;

It is unchanged, but we know how many A's are present