COMP 364 Lecture 24 Wednesday, March 5th

Announcements:

- Quiz 1 graded
- HW 2 not yet graded
- HW 3 is supposed to be posted completely by Friday
 - Should be due **March 14th** (late by march 17th)
- Quiz 2 material includes everything from Quiz 1 until the next lecture
 - Arrays, Hashes, Pattern Matching.
 - Will take place on March 19th (Notes are allowed)
 - Material from Quiz 1 won't be explicitly tested on.
- Today: Pattern Matching:
 - Anchors, Getting "all" matches, Substitution & Translation

Note:

 $\$str = \sim /./ = >$ period matches any character except new line unless: $\$str = \sim /./s = >$ s at the end will allow matching of ANY character including new line

<u>Problem:</u> Does an amino acid sequence begin with a certain pattern (for example AXV) and end with another pattern(e.g. YYD)?

 $\$protSeq = \sim /AXV.*YYD/ = >$ Is **incorrect!** The match can occur anywhere in the and we are interested in the start and the end of the sequence

string,

Anchors:

- ^ matches a pattern at the start of a string
- \$ matches a pattern at the end of the string

 $protSeq = \ /\ AXV. *YYD / =$ Is the correct solution for the above problem.

There are more anchors in the book, but won't be covered in class.

Getting "all" the matches:

@ $All = \$protSeq = \sim /A/g; =>$ @All gets all matches in \$protSeq g – stands for global.

Sample Code:

\$protSeq = "ADSASAQHDSAHUAHOY" @All = \$protSeq =~ /A/g; print "@All\n"; #will print the letter A 5 times

 $@All = \$protSeq = \sim /A/H/g;$

print "@All\n"; #will print the letter A 5 times and the letter H 3 times

Caveats to "all":

each successive match must start after previous ends

$$'EAAB(1)' = \sim /AA/AB/g$$

 $/A(A/B)/g$

Will find the AA but not the AB

Could be solved by searching for AA and AB separately(on a separate line).

By default * and + take as much as they can"

$$@All = 'BAAAAAA BC' = \sim /A + /g;$$

- A, AA, AAA, AAAA, AAAAA will all match the above pattern, but in fact only the longest pattern (AAAAA) will be returned. It will return as many A in a row as it can find. 'BABABABC' =~ /(AB*)./g => will give ABABABC
- This default behavior can be changed by adding a ? after the * or +.

@
$$All = 'BAAAAABC' = \sim /A + ?/g \rightarrow ('A', 'A', 'A', 'A', 'A')$$
 an array that contains a single A five times

$$@ALL = "BABABABC' = \sim /(AB+?)./g \rightarrow$$

? -- will match the desired pattern only once(or as few times as possible - zero or one)

AUG (...)*?UAA – finds first start codon & first subsequent stop codon