HW 1: Write a Perl program that implements DNA translation to amino acit sequence

Your program should read the sequence from a data file and write the results into another data file

Submission deadline: 24 March 2019

PS: HWs submitted after the deadline will not be accepted

Email your HW to: nizamettinaydin@gmail.com You should place **I2B-HW1** in the "Subject" field.

You must name your file by using your matriculation number (for example, 07011068-hw1.pl)

Reminder:

Example 1.

Transcribe the following DNA to RNA, then use the genetic code to translate it to a sequence of amino acids.

${\tt TCATAATACGTTTTGTATTCGCCAGCGCTTCGGTGT}$

Answer 1.

To transcribe the DNA, first substitute each DNA for it's counterpart (i.e., G for C, C for G, T for A and A for T): TCATAATACGTTTTGTATTCGCCAGCGCTTCGGTGT

AGTATTATGCAAAACATAAGCGGTCGCGAAGCCACA

Next, remember that the Thymine (T) bases become a Uracil (U). Hence our sequence becomes: AGUAUUAUGCAAAACAUAAGCGGUCGCGAAGCCACA

Using the genetic code is also easy – just split the RNA sequence into triplets: :

AGU AUU AUG CAA AAC AUA AGC GGU CGC GAA GCC ACA

then look each triplet (codon) up in the genetic code table. So AGU becomes Serine, which we can write as Ser, or just S. AUU becomes Isoleucine (Ile), which we write as I. Carrying on in this way, we get:

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			Seco	nd letter			
		U	С	Α	G		
First letter	U	UUU Phe UUC Leu	UCU UCC UCA UCG	UAU Tyr UAC Stop UAG Stop	UGU Cys UGC Stop UGG Trp	UCAG	Third letter
	O	CUU CUC CUA CUG	CCU CCC CCA CCG	CAU His CAA GIn CAG	CGU CGC CGA CGG	U C A G	
	A	AUU AUC AUA lle AUG Met	ACU ACC ACA ACG	AAU Asn AAC AAA AAG Lys	AGU AGC Ser AGA AGG Arg	U C A G	
	G	GUU GUC GUA GUG	GCU GCC GCA GCG	GAU GAC Asp GAA GAG Glu	GGU GGC GGA GGG	UCAG	

A=Ala=Alanine
C=Cys=Cysteine
D=Asp=Aspartic acid
E=Glu=Glutamic acid
F=Phe=Phenylalanine
G=Gly=Glycine
H=His=Histidine
I=Ile=Isoleucine
K=Lys=Lysine
L=Leu=Leucine
M=Met=Methionine
N=Asn=Asparagine
P=Pro=Proline
Q=Gln=Glutamine
R=Arg=Arginine
S=Ser=Serine
T=Thr=Threonine
V=Val=Valine
W=Trp=Tryptophan
Y=Tyr=Tyrosine