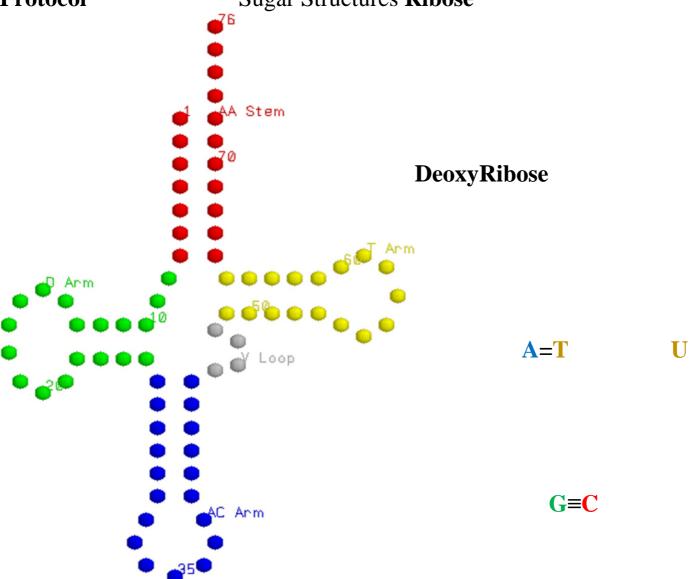
Name Surname:_ Constituents of N	ucleopro	teins (Nucleotides or P	olyNucleotides containing	_	2018.Year ouching Proteins)
	_	·	insAS.doc A Task for		
				)	<b>(33)</b>
using Interactive N	Molecule	viewers: Chemscap	e MDL RasWin G	FireFox 3.5	5.5.v
		professor Marz prep			
	-	es/DnaMarzHTM/II			
-		•	Corey, Pauling, Kolt	un publication	of scientists in
		Menu Stripe) Ball &		=	SA Patent 1965
Atom Name	Symbol	Color	Valence Number		c modeling
Carbon	•	Gray lightly or Black	4		•
	$\mathbb{H}$	White	1		<u>phate</u> backbone <u>PO</u> <sub>4</sub> -△- is ribose △
Hydrogen			1		alent ester bonds
Oxygen	O		onor acceptor ligand up to 4)		es of oxygen.
Nitrogen Sulfur	N S	<b>Bluish 3</b> + 1 (d <b>Yellow</b>	onor acceptor ligand up to 4) -2, +6		RNA bases
Phosphor		Yellow Intensive dark	5 ( & 3 )		nine-Green
Sodium ion	Na <sup>+</sup>	Blue	+1 (coordination up to 6)		ine-Carmine
Magnesium ion		Green	+2 (coordination up to 6)	•	nine-Azure
Calcium ion	$\mathrm{Mg}^{2+} \ \mathrm{Ca}^{2+}$	Gray Dark	+2 (coordination up to 6)		e-Tweety bird
Iron ion	$Fe^{2+}$	Yellow Gray	+2 (coordination up to 6)	•	il-PUrple
Iron ion	<b>Fe</b> <sup>3+</sup>	Yellow Gray	+3 (coordination up to 6)		птетри
1. How many base	e <b>pairs</b> d	•	NA fragment?		
•	_	•	ose <b>DNA</b> of one stra		
2. Which one two	morecun	ar components comp	obe Bivi of one stru	ila buckbolle	•
A 1171 1	C		1 11 . 10	CDNIAG	
3. What net charge	e of one	strand and complete	double strand fragmo	ent of <b>DNA</b> ?.	•••••
<b>4.</b> Which two 2 ki	nd bonds	s-interaction forces (	underline those) suppo	ort stabile stru	icture of <b>DNA</b>
in cellular water ${\mathbb B}$	120 med	ium? Are known fiv	ve <b>5</b> bonds-interaction	n forces in Bi	ochemistry!
			disulfide bridge,5.coordi		•
	molecul	ar units of two chose	en intermolecular bo	nds for <b>DNA</b>	stability:
1			2		•••••
<b>6.</b> Draw on protoc	ol paper	the structural planar	laying of colored ato	oms on comp	uter screen
to symbolic	type of a	toms for two 2 type	base pairs : with two	o hydrogen bo	onds A=T and
<b>G</b> ≡ <b>C</b> with three	e hydrog	en bonds adding hyd	lrogens ⊞ in screen p	oicture using	the button!
-	-	replication of given			<b>————————————————————————————————————</b>
<b>8.</b> Show the forwa	rd direct	G C symbols of <b>bas</b> ion $5' \rightarrow 3'$ markers		  <u>                         </u>	<b></b>
position on ends o and $3' \leftarrow 5'$ anti p		trand <b>fragment</b> ; rection of <b>DNA</b> mar	kers!		
<b>9.</b> Show the difference	ence in <b>r</b>	replication using ba	se symbols A T G C	and	
transcription of g	viven <b>DN</b>	A fragment in to RI	<b>NA</b> →5'		
sequences I and II	using <b>b</b>	<b>ases A U G C</b> symb			<b>□</b> →3'
10. What base pair	-		<b>a</b>		
T base replaced by	y	•••••	3'← -⊞ <mark>0</mark>		
					•

Name Surname:	9		/D: 1 -		N°201	
http://aris.gusc.lv/ChemFiles/CarnegieMellonU0 prepared in Carnage Melon Univer						
	•		•		-	
1. Find the 5-terminal and 3-terminal						•••
2. Determine nucleotide account on or						
3. Determine net charge of one strand					••••••	•••••
<b>4.</b> What size has <b>tRNA</b> Phe molecule i					11 11	10 20 2
A 5. Which six molecular c	_	_				
↑ structure, <u>call them</u> on re		Uracil a	and D-Ri	bose struct	ture on repo	rt paper
$ \leftarrow \mathring{\mathbf{A}} \rightarrow  $ <b>6.</b> What structure ty	_					
7. Call and explain the	ne five (5)	loops sec	condary 2	2° structur	es for <b>tRN</b> A	Phe
_*	•••••	•••••	•••••	•••••		
0 Cl	1	1.0	11		C ADDIA I	Dha t
<b>8.</b> Show <u>anticodon</u> loop like <b>U</b> four n				_	tor <b>tRNA</b>	ne y
U loop → 5'						
3 nucleotides $\rightarrow$ 5'						
mRNA codon← 3'						
<b>9.</b> Mark on table of <b>genetic codes</b> inv	_				••••	
<b>Table 1. The genetic code</b> . For messe		4 molecu			tic Code	1
Note: that those messenger <b>mRNA</b>	1st			l position		3rd
code begin with U1,C1,A1,G1 but	position (5' and)	U	C	A	G	position
second element with U2 as well	(5' end) ↓		G			(3' end)
C1,G1 with second C2 <u>nucleotide</u>	U	Phe Phe	Ser Ser	Tyr Tyr	Cys Cys	C
tend to specify for translation seven		Leu	Ser	STOP	S-SelCys	A
amino acids on protein chain most		Leu	Ser	STOP	Trp	G
hydrophobic.		Leu	Pro	His	Arg	U
g I		Leu	Pro	His	Arg	C
<b>10.</b> Which three <u>nucleotides</u> set on		Leu	Pro	Gln	Arg	A
codon sequences to specify seven		Leu	Pro	Gln	Arg	G
amino acids having hydrophobic	$\mathbf{A}$	Ile	Thr	Asn	Ser	U
properties!		Ile	Thr	Asn	Ser	C
properties.		Ile <b>Met init</b>	Thr Thr	Lys Lys	Arg Arg	A G
••••••	~	Val	Ala	Asp	Gly	U
•••••	$\mathbf{G}$	Val Val	Ala	Asp	Gly	C
••••••		Val	Ala	Glu	Gly	A
		Val	Ala	Glu	Gly	$\mathbf{G}$
11. Call & depict in short cut symbols	twelve 12	2 tertiary	3° unusu	al structur	res for <b>tRN</b>	<b>A</b> Phe
	•••					
	•••••					
	4.		•-	0.40-	٠	
<b>12.</b> How many $\underline{\mathbf{Mg}}^{2+}$ ionsand What force structural molecular unit of done					coordinativ	/e

**Protocol** Sugar Structures **Ribose** 



Ribose 
$$-\mathbf{0}$$
— $\mathbf{P}$ — $\mathbf{0}$ — Ribose  $-\mathbf{0}$ — $\mathbf{P}$ — $\mathbf{0}$ — $\mathbf{0}$ 

Name Surname: \_\_\_\_\_\_, \_\_\_\_\_ Group  $N^o$  \_\_\_\_2018.Year

Table 1. The genetic code. For messenger RNA molecule  $mRNA \ \Psi$  Genetic Code

Sets of three 3 nucleotides (codons) in an mRNA molecule are translated into amino acids AA in the course of protein synthesis according to the rules shown. The codons G U G and GAG, for example, are translated into valine and glutamic acid, respectively.

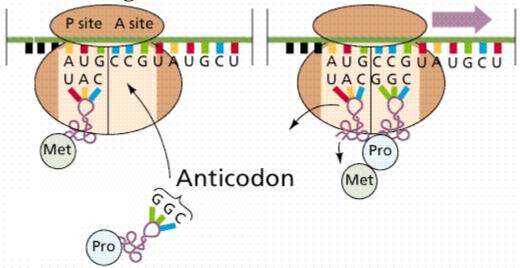
Note: that those messenger mRNA code begin with U1,C1,A1,G1 but second element with U2 as well C1,G1 with second C2 nucleotide tend to specify for translation on protein chain seven amino acids most hydrophobic.

lessenger RNA molecule HIRNA 4 Genetic Coue						
<b>1</b> st		3rd				
position		position				
(5' end) 🗆	U	C	A	$\mathbf{G}$	(3' end) 🗆	
4	Phe	Ser	Tyr	Cys	U	
	Phe	Ser	Tyr	Cys	C	
	Leu	Ser	<b>STOP</b>	S-SelCys	A	
	Leu	Ser	STOP	Trp	G	
2	Leu	Pro	His	Arg	U	
( '	Leu	Pro	His	Arg	C	
	Leu	Pro	Gln	Arg	A	
	Leu	Pro	Gln	Arg	G	
	Ile	Thr	Asn	Ser	U	
$\mathbf{A}$	Ile	Thr	Asn	Ser	C	
<u> </u>	Ile	Thr	Lys	Arg	A	
	Met init	Thr	Lys	Arg	G	
	Val	Ala	Asp	Gly	U	
	Val	Ala	Asp	Gly	C	
	Val	Ala	Glu	Gly	A	
	Val	Ala	Glu	Gly	$\mathbf{G}$	
	1st position	1st position (5' end)   U  Phe Phe Leu Leu Leu Leu Leu Leu Leu Leu Lou Leu Leu Leu Leu Leu Leu Leu Leu Leu Le	1st position (5' end) U C  Phe Ser Phe Ser Leu Ser Leu Ser Leu Pro Leu	1st position (5' end)     U	1st position (5' end) U C A G  Phe Ser Tyr Cys Tyr Cys Leu Ser STOP S-SelCys Leu Pro His Arg His Arg Leu Pro Gln Arg Gln Arg Leu Pro Gln Arg Leu Pro Gln Arg Leu Pro His Arg Leu Pro His Arg Leu Pro His Arg Leu Pro His Arg H	

Translation in ribosome start with methionine:

Met init, Pro, Phe, Leu, Ile, Val, Ala. 1, 2, 3, 4, 5, 6, 7

## How do we go from mRNA to Protein?



8. encodes transfer tRNAPhe ribonucleic acid for phenylalanine Phe amino acid transport.

4 nucleotides U loop:  $\rightarrow$  5'U33-O2'MG34 - A35 - A36 3'  $\rightarrow$  anticodon loop

3 nucleotides Phe anticodon: → 5' O2'MG34 – A35 – A36 3' → anticodon sequence

3 nucleotides codon on mRNA:  $\leftarrow$  3' C3 - U2 - U1 5'  $\leftarrow$  codon sequence ant parallel

Incoming AA Pro tRNA with←3'GGC5' anticodon complementar

to mRNA →5'CCG3' codon

Initiation of the Translation begins by AminoAacid Met init tRNA  $\leftarrow$ 3' UAC 5' anticodon complementar to mRNA  $\rightarrow$ 5' AUG 3' codon