

Solution:

									G	T	G	G	G	T	G	A	A	A	A	T	G	A	A	A	C	T	A	T	T	T	G	T	T	C	C	T	G	A	
C	C	T	T	G	C	C	G	C	G	T	G	G	G	T																									
A	C	T	T	G	C	C	G	C	G	T	G	G	C	T																									
				G	C	C	G	C	G	T	G	G	G	T	G	A	A	A																					
					A	C	T	C	G	T	G	T	G	T	G	A	A	A	A																				
							G	C	G	T	G	G	G	T	G	A	A	A	A	T	G																		
										T	G	A	G	T	G	A	A	A	A	T	G	A	A	A															
												G	G	T	G	A	A	A	A	T	G	A	A	A	C	T													
												G	G	T	G	A	A	A	A	T	A	A	A	A	C	T													
													G	T	A	A	A	A	A	T	G	A	A	A	C	T	G												
													G	G	G	A	A	A	A	T	G	A	A	A	C	T	C												
																A	A	A	A	T	G	A	A	A	C	T	A	T	T	T									
																	A	A	A	T	G	A	A	A	C	T	A	T	T	T	G								
																					G	A	A	A	C	T	A	T	T	T	G	G	T	C	C				
																						A	A	A	C	T	C	T	T	T	G	T	T	C	C	T			
																									C	T	A	T	T	T	G	T	T	C	C	T	G	A	C
																									C	G	A	T	T	T	G	T	T	C	C	T	G	A	C

Amino acid sequence in each three reading frames:

Nucleotide sequence:																																		
C	C	T	T	G	C																								C	C	T	G	A	C

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[illegible]

Reads - cut out

A	C	T	T	G	C	C	G	C	G	T	G	G	C	T
A	A	A	C	T	C	T	T	T	G	T	T	C	C	T
C	C	T	T	G	C	C	G	C	G	T	G	G	G	T
C	G	A	T	T	T	G	T	T	C	C	T	G	A	C
G	C	C	G	C	G	T	G	G	G	T	G	A	A	A
G	G	T	G	A	A	A	A	T	A	A	A	A	C	T
G	G	G	A	A	A	A	T	G	A	A	A	C	T	C
G	C	G	T	G	G	G	T	G	A	A	A	A	T	G
C	T	A	T	T	T	G	T	T	C	C	T	G	A	C
A	A	A	T	G	A	A	A	C	T	A	T	T	T	G
C	T	A	T	T	T	G	T	T	C	C	T	G	A	C
G	A	A	A	C	T	A	T	T	T	G	G	T	C	C
C	C	T	T	G	C	C	G	C	G	T	G	G	G	T
A	A	A	A	T	G	A	A	A	C	T	A	T	T	T
A	C	T	C	G	T	G	T	G	T	G	A	A	A	A
G	T	A	A	A	A	A	T	G	A	A	A	C	T	G
G	G	T	G	A	A	A	A	T	G	A	A	A	C	T
T	G	A	G	T	G	A	A	A	A	T	G	A	A	A

Amino Acid translation table

		Second base in codon												
		T			C			A			G			
First base in codon	T	TTT	Phe	F	TCT	Ser	S	TAT	Tyr	Y	TGT	Cys	Y	T
		TTC	Phe	F	TCC	Ser	S	TAC	Tyr	Y	TGC	Cys	Y	C
		TTA	Leu	L	TCA	Ser	S	TAA	Stop	*	TGA	Stop	*	A
		TTG	Leu	L	TCG	Ser	S	TAG	Stop	*	TGG	Trp	W	G
	C	CTT	Leu	L	CCT	Pro	P	CAT	His	H	CGT	Arg	R	T
		CTC	Leu	L	CCC	Pro	P	CAC	His	H	CGC	Arg	R	C
		CTA	Leu	L	CCA	Pro	P	CAA	Gln	Q	CGA	Arg	R	A
		CTG	Leu	L	CCG	Pro	P	CAG	Gln	Q	CGG	Arg	R	G
	A	ATT	Ile	I	ACT	Thr	T	AAT	Asn	N	AGT	Ser	S	T
		ATC	Ile	I	ACC	Thr	T	AAC	Asn	N	AGC	Ser	S	C
		ATA	Ile	I	ACA	Thr	T	AAA	Lys	K	AGA	Arg	R	A
		ATG	Met	M	ACG	Thr	T	AAG	Lys	K	AGG	Arg	R	G
	G	GTT	Val	V	GCT	Ala	A	GAT	Asp	D	GGT	Gly	G	T
		GTC	Val	V	GCC	Ala	A	GAC	Asp	D	GGC	Gly	G	C
		GTA	Val	V	GCA	Ala	A	GAA	Glu	E	GGAGly	Gly	G	A
		GTG	Val	V	GCG	Ala	A	GAG	Glu	E	GGGGly	Gly	G	G

Third base in codon

Standard genetic code. The table shows how triplets of nucleic acid bases correspond to different amino acids. Besides the codon ATG with always codes for methionine, alternatively TTG, CTG, ATT, ATC, ATA and GTG can serve as initiation codons, in which case they are translated as methionine rather than the amino acid indicated.