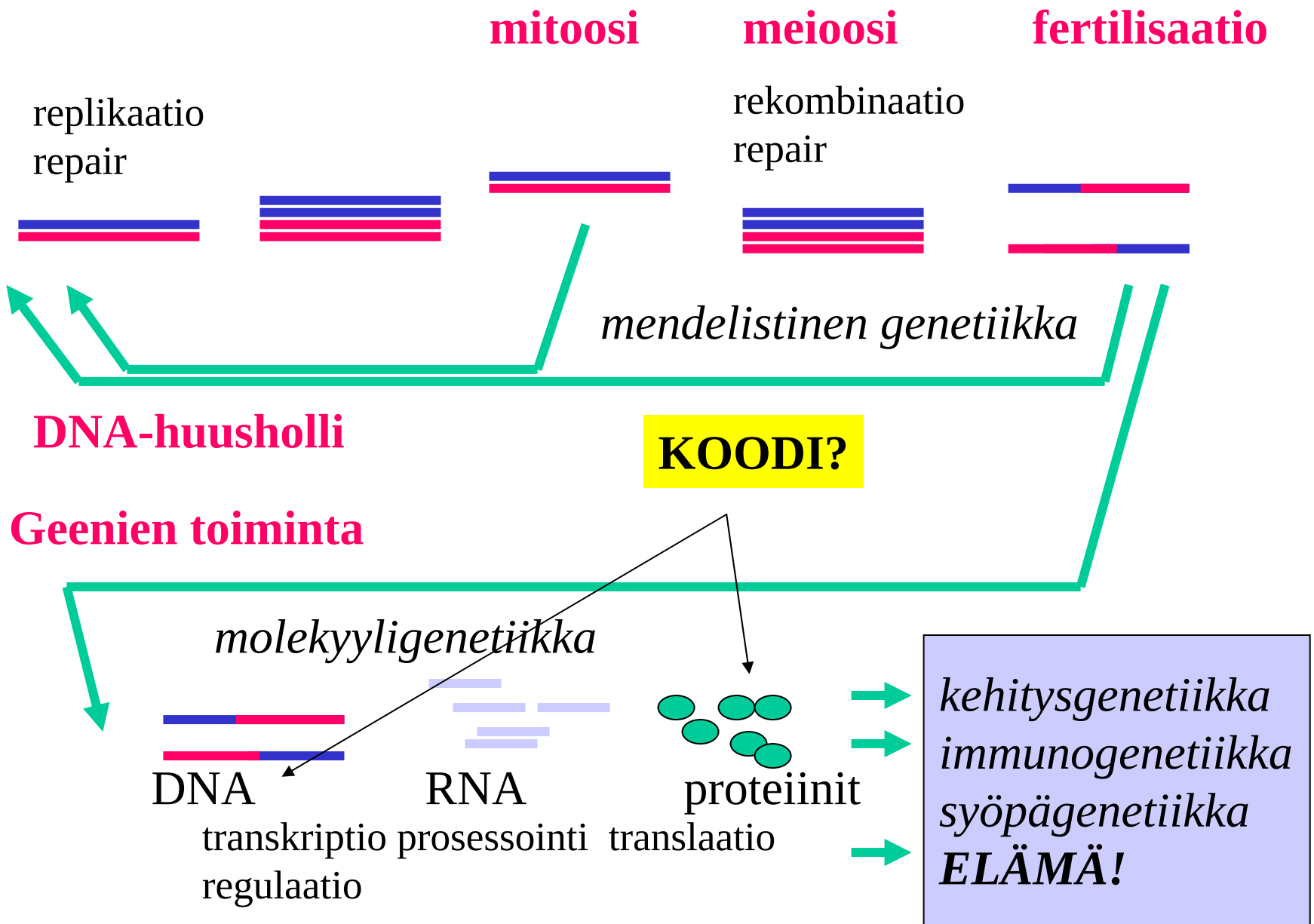




Genetiikan perusteiden miellekartta: geneettinen triplettikoodi

Translaatiossa mRNA:ssa tarvitaan seuraavat *sanomat* :

m⁷G - pipo

Ehkä internal ribosome entry site **IRES**

poly-A -häntä sirkulaarisen polysomin muodostamiseksi

AUG aloituskodoni ensimmäistä metioniinia varten

UAA, UAG tai **UGA** lopetuskodoni RF:ää varten

Sekä ennen AUG-kodonia että STOP-kodonin jälkeen voi olla säätelyyn osallistuvia **UTR**- jaksoja (untranslated region)

UTR koodi **UTR**
m⁷G ---**AUG**----(**IRES**) -----**AUG** -----...-----**UAA**-----**p(AAA)**

Translaatiossa mRNA:ssa tarvitaan seuraavat *sanomat* :

m⁷G - pipo

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m⁷G ---AUG----(IRES) -----AUG-----...-----UAA-----p(AAA)

Tässä tulikin jo pari triplettikoodin sanaa

Aloitusta (= metioniini) **AUG**

STOP: **UAA**, **UGA** tai **UAG**

Koodin murtaminen oli molekyyli­genetiikan historian suuria juttuja: nobelistit Holley (tRNA), Khorana ja Nirenberg (1968)



The Genetic Code

1st position (5' end) ↓	U	C	A	G	3rd Position (3' end) ↓
U	Phe Phe Leu Leu	Ser Ser Ser Ser	Tyr Tyr STOP STOP	Cys Cys STOP Trp	U C A G
C	Leu Leu Leu Leu	Pro Pro Pro Pro	His His Gln Gln	Arg Arg Arg Arg	U C A G
A	Ile Ile Ile Met	Thr Thr Thr Thr	Asn Asn Lys Lys	Ser Ser Arg Arg	U C A G
G	Val Val Val Val	Ala Ala Ala Ala	Asp Asp Glu Glu	Gly Gly Gly Gly	U C A G

Amino Acids Codons

A	Ala	Alanine	GCA GCC GCG GCU
C	Cys	Cysteine	UGC UGU
D	Asp	Aspartic acid	GAC GAU
E	Glu	Glutamic acid	GAA GAG
F	Phe	Phenylalanine	UUC UUU
G	Gly	Glycine	GGA GGC GGG GGU
H	His	Histidine	CAC CAU
I	Ile	Isoleucine	AUA AUC AUU
K	Lys	Lysine	AAA AAG
L	Leu	Leucine	UUA UUG CUA CUC CUG CUU
M	Met	Methionine	AUG
N	Asn	Asparagine	AAC AAU
P	Pro	Proline	CCA CCC CCG CCU
Q	Gln	Glutamine	CAA CAG
R	Arg	Arginine	AGA AGG CGA CGC CGG CGU
S	Ser	Serine	AGC AGU UCA UCC UCG UCU
T	Thr	Threonine	ACA ACC ACG ACU
V	Val	Valine	GUA GUC GUG GUU
W	Trp	Tryptophan	UGG
Y	Tyr	Tyrosine	UAC UAU

DNA words are three letters long.

22

DNA FROM THE BEGINNING

The genetic code had to be a "language" – using the DNA alphabet of A, T, C, and G – that produced enough DNA "words" to specify each of the 20 known amino acids. Simple math showed that only 16 words are possible from a two-letter combination, but a three-letter code produces 64 words. Operating on the principle that the simplest solution is often correct, researchers assumed a three-letter code called a codon.

Research teams at University of British Columbia and the National Institutes of Health laboriously synthesized different RNA molecules, each a long strand composed of a single repeated codon. Then, each type of synthetic RNA was added to a cell-free translation system containing ribosomes, transfer RNAs, and amino acids. As predicted, each type of synthetic RNA produced a polypeptide chain composed of repeated units of a single amino acid. Several codons are "stop" signals and many amino acids are specified by several different codons, accounting for all 64 three-letter combinations.



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 - 34 [Genes can be moved between species.](#)
 - 35 [DNA responds to signals from outside the cell.](#)
 - 36 [Different genes are active in different kinds of cells.](#)
 - 37 [Master genes control basic body plans.](#)
 - 38 [Development balances cell growth and death.](#)
 - 39 [A genome is an entire set of genes.](#)
 - 40 [Living things share common genes.](#)
 - 41 [DNA is only the beginning for understanding the human genome.](#)



DNA words are three letters long.

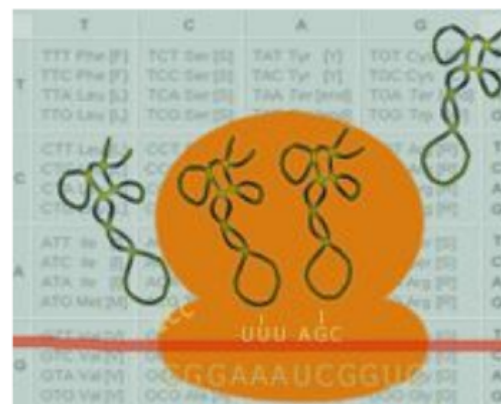
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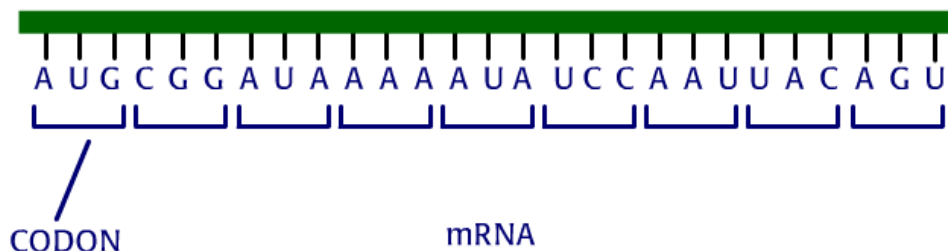
DNA words are three letters long.

22

DNA FROM THE BEGINNING



I'm Marshall Nirenberg. Har Khorana's group and mine "cracked" the genetic code. We figured out how the nucleotide language of mRNA is "translated" into the amino acid language of proteins. Genetic data from Crick and others showed that three nucleotides form a "codon" – an mRNA word that specifies one amino acid.



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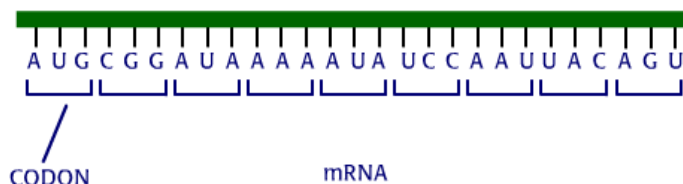
DNA words are three letters long.

22

DNA FROM THE BEGINNING



This made sense, because a codon made from only one or two nucleotides would not produce enough combinations (words) to code for all 20 of the known amino acids.



1 NUCLEOTIDE = 4 POSSIBLE CODONS

2 NUCLEOTIDES = 4 X 4 POSSIBLE CODONS

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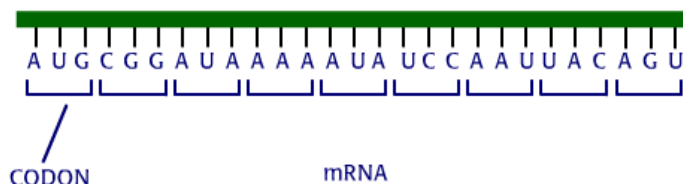
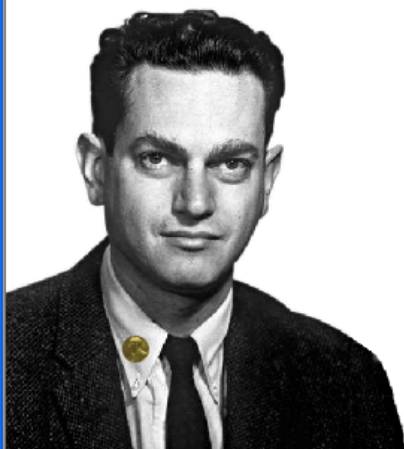
DNA words are three letters long.

22

DNA FROM THE BEGINNING



However, a three-nucleotide codon produces 64 combinations. This would produce a redundant, or degenerate, code where several different codons specify the same amino acid. The parsimony principle – that the simplest solution is often right – ruled out a four-nucleotide codon.



3 NUCLEOTIDES = 4 X 4 X 4 POSSIBLE CODONS
3 NUCLEOTIDES = 64 codons for 20 amino acids

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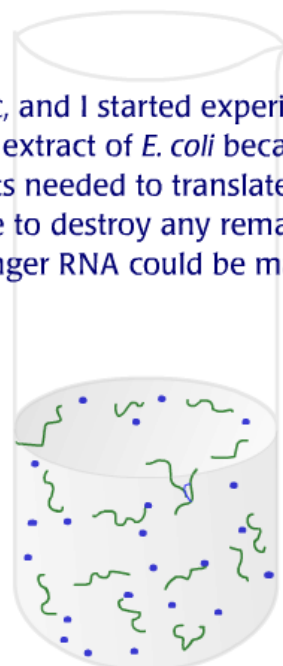
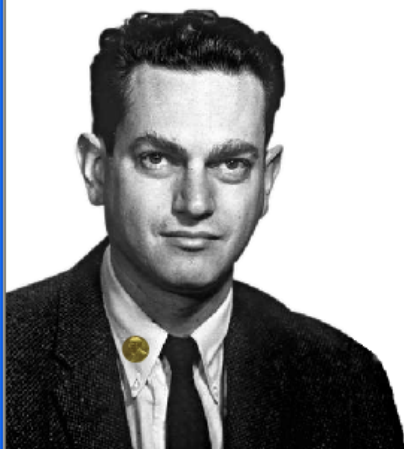
DNA words are three letters long.

22

DNA FROM THE BEGINNING



In 1961, Heinrich Matthaei, a visiting post-doc, and I started experiments to test the triplet codon hypothesis. We used a cell-free extract of *E. coli* because we believed that this extract should contain all the components needed to translate mRNA into proteins. The extract was treated with DNase to destroy any remaining *E. coli* DNA – so there was no template from which messenger RNA could be made.



CELL-FREE EXTRACT

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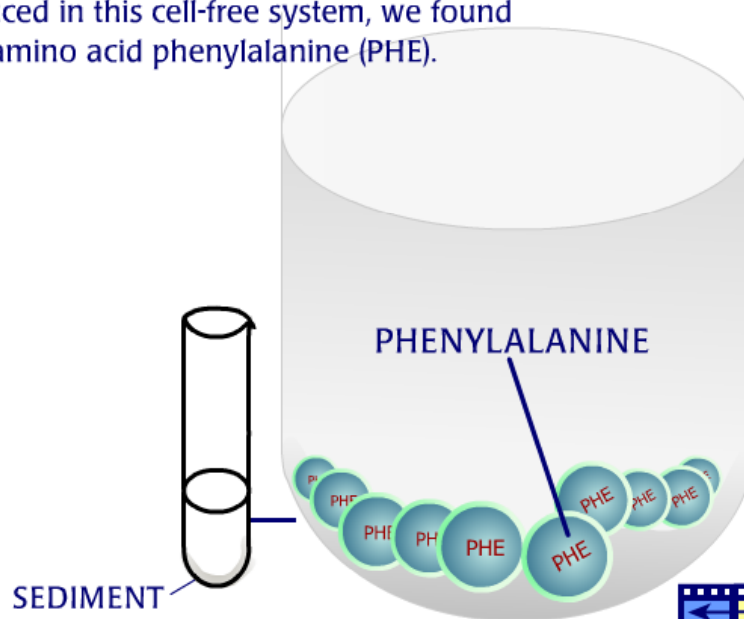
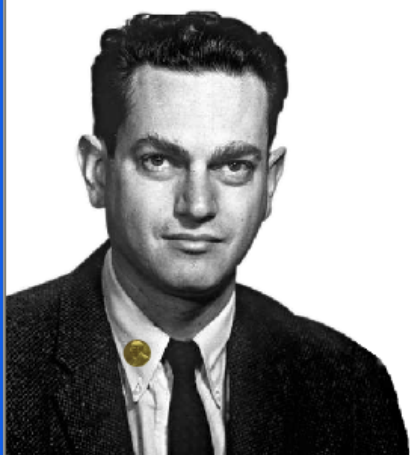
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DNA FROM THE BEGINNING



When we examined the products produced in this cell-free system, we found polypeptides composed entirely of the amino acid phenylalanine (PHE).



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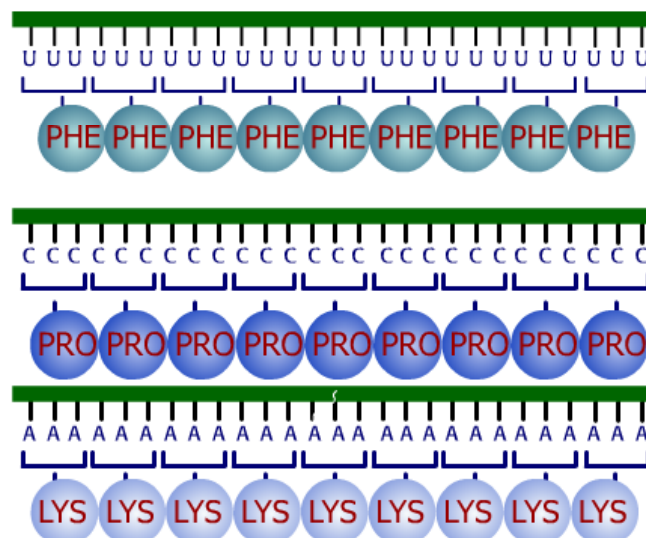
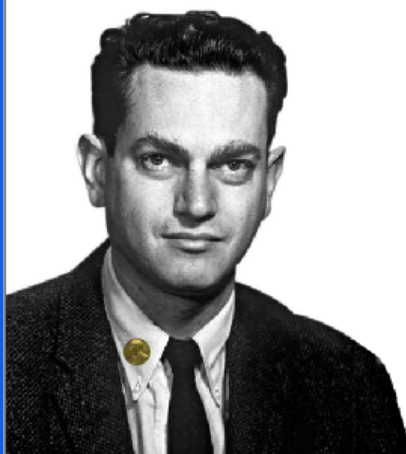


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NA words are three letters long.

22

DNA FROM THE BEGINNING



Other researchers and I quickly recognized the power of this approach. Using RNA templates containing different nucleotide combinations, we assigned amino acids to about 50 triplet codons.



		SECOND LETTER								
		U	C	A	G					
U	UUU	PHE	UCU	UAU	UGU	CYS	U			
	UUC		UCC				UAC	?	C	
	UUA		UCG				?	?	A	
	UUG		?				?	UGG	TRP	G
C	CUU	LEU	CCU	CAU	CGU	ARG	U			
	CUC		CCC				CAC	CGC	C	
	?		CCA				CAA	CGA	A	
	?		CCG				?	?	G	
A	AUU	ILE	ACU	AAU	?	?	U			
	AUC		ACC				AAC	AGC	SER	C
	AUA		ACA				AAA	AGA	ARG	A
	AUG		MET				?	?	?	G
G	GUU	VAL	GCU	GAU	GGU	GLY	U			
	GUA		GCC				GAC	GGC	C	
	?		GCA				GAA	GGA	A	
	?		?				?	?	G	

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DNA FROM THE BEGINNING



Hi, I'm Phil Leder. I helped Marshall Nirenberg with the rest of the genetic code.



		SECOND LETTER					
		U	C	A	G		
FIRST LETTER	U	UUU } PHE UUC } UUA } LEU UUG }	UCU } SER UCC } UCG } UCA ?	UAU } TYR UAC } ? } ? }	UGU - CYS UGC ? ? } UGG - TRP	U C A G	THIRD LETTER
	C	CUU } LEU CUC } CUA ? CUG ?	CCU } CCC } CCA } CCG }	CAU } HIS CAC } CAA - GLN CAG ?	CGU } CGC } CGA } CGG ?	U C A G	
	A	AUU } AUC } AUA } AUG - MET	ACU } ACC } ACA } ACG ?	AAU } ASN AAC } AAA - LYS AAG ?	AGU ? AGC - SER AGA - ARG AGG ?	U C A G	
	G	GUU } VAL GUA } GUC } GUG ?	GCU } GCC } GCA } GCG ?	GAU } ASP GAC } GAA - GLU GAG ?	GGU } GGC } GGA } GGG ?	U C A G	



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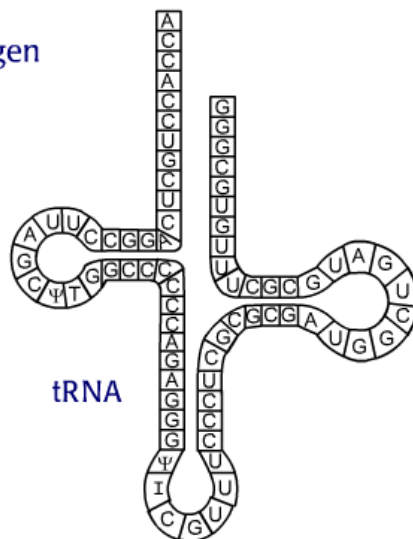
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22

DNA FROM THE BEGINNING



tRNA is the molecule that carries amino acids to the ribosomes for protein synthesis. In 1962, Robert Holley solved the structure of tRNA. Although tRNA is single stranded, stretches of complementary nucleotides hydrogen bond to form short double-stranded regions, which bend the tRNA into a characteristic cloverleaf shape.



tRNA



MOLECULES OF GENETICS

GENETIC ORGANIZATION AND CONTROL

- 6 [Genes are real things.](#)
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- 12 [Evolution begins with the inheritance of gene variations.](#)
- 13 [Mendelian laws apply to human beings.](#)
- 14 [Mendelian genetics cannot fully explain human health and behavior.](#)

- 15 [DNA and proteins are key molecules of the cell nucleus.](#)
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- 17 [A gene is made of DNA.](#)
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- 19 [The DNA molecule is shaped like a twisted ladder.](#)
- 20 [A half DNA ladder is a template for copying the whole.](#)
- 21 [RNA is an intermediary between DNA and protein.](#)
- 22 [DNA words are three letters long.](#)
- 23 [A gene is a discrete sequence of DNA nucleotides.](#)
- 24 [The RNA message is sometimes edited.](#)
- 25 [Some viruses store genetic information in RNA.](#)
- 26 [RNA was the first genetic molecule.](#)
- 27 [Mutations are changes in genetic information.](#)
- 28 [Some types of mutations are automatically repaired.](#)

- 29 [DNA is packaged in a chromosome.](#)
- 30 [Higher cells incorporate an ancient chromosome.](#)
- 31 [Some DNA does not encode protein.](#)
- 32 [Some DNA can jump.](#)
- 33 [Genes can be turned on and off.](#)
- 34 [Genes can be moved between species.](#)
- 35 [DNA responds to signals from outside the cell.](#)



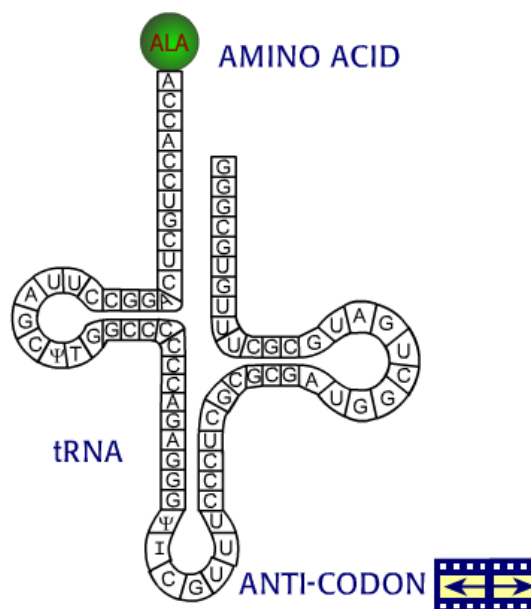
DNA words are three letters long.

22

DNA FROM THE BEGINNING



Using their cell-free translation system, Zamecnik and his group showed that tRNA becomes activated when the amino acid binds to the tRNA's stem. This step requires chemical energy in the form of ATP.



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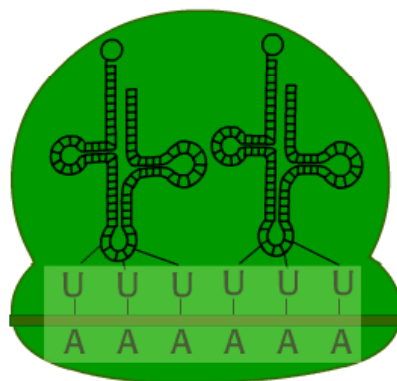
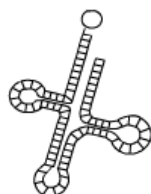
DNA words are three letters long.

22

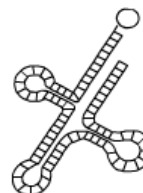
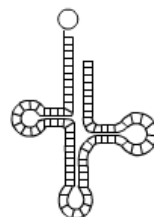
DNA FROM THE BEGINNING



I first made short RNA chains composed of three or six nucleotides. When added to a cell-free extract, an activated tRNA will read these trinucleotide or hexanucleotide sequences.



HEXANUCLEOTIDE SEQUENCE



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DNA FROM THE BEGINNING



After that, all I had to do was make specific trinucleotide and hexanucleotide chains to confirm the genetic code.



		SECOND LETTER					
		U	C	A	G		
FIRST LETTER	U	UUU } PHE UUC } UUA } LEU UUG }	UCU } UCC } SER UCG } UCA }	UAU } TYR UAC } UAA } UAG }	UGU } CYS UGC } UGA } UGG } TRP	THIRD LETTER	U
	C	CUU } CUC } LEU CUA } CUG }	CCU } CCC } CCA } PRO CCG }	CAU } HIS CAC } CAA } ASN CAG }	CGU } CGC } ARG CGA } CGG }		C
	A	AUU } AUC } ILE AUA } AUG } MET	ACU } ACC } ACA } THR ACG }	AAU } ASN AAC } AAA } LYS AAG }	AGU } AGC } SER AGA } ARG AGG }		A
	G	GUU } GUA } VAL GUC } GUG }	GCU } GCC } GCA } GCG }	GAU } ASP GAC } GAA } GLU GAG }	GGU } GGC } GGA } GGG }		G

GENETIC CODE TABLE

- Genes are real things.
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DNA words are three letters long.

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DNA FROM THE BEGINNING



We also found that translation starts with a specific codon AUG – the only unique codon – and there are stop codons that end translation. So, in addition to words, the genetic code also has punctuation points.



		SECOND LETTER					
		U	C	A	G		
FIRST LETTER	U	UUU } PHE UUC } UUA } LEU UUG }	UCU } UCC } SER UCG } UCA }	UAU } UAC } UAA } STOP UAG } STOP	UGU } UGC } UGA } STOP UGG } TRP	U C A G	THIRD LETTER
	C	CUU } CUC } LEU CUA } CUG }	CCU } CCC } CCA } PRO CCG }	CAU } CAC } CAA } CAG }	CGU } CGC } CGA } CGG }	U C A G	
	A	AUU } AUC } AUA } AUG } MET	ACU } ACC } ACA } ACG }	AAU } AAC } AAA } AAG }	AGU } AGC } AGA } AGG }	U C A G	
	G	GUU } GUA } GUC } GUG }	GCU } GCC } GCA } GCG }	GAU } GAC } GAA } GAG }	GGU } GGC } GGA } GGG }	U C A G	

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(Mouseover table to enlarge)

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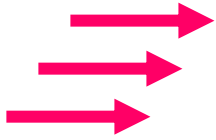
Geneettinen koodi ja avoimet lukukehykset: polypeptidiä koodaavan geenin “ennustamisen” päävälineet

On tapana kirjoittaa ja tallettaa geenipankkiin vain toinen juoste, alkaen 5' päästä, esimerkiksi näin:

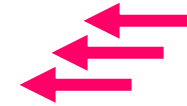
5' - TTATTTTATTTTCGATAATTCGACCTTAAACGCGAAACTTCACTTAAC - 3'

Geenin viesti voi kumminkin olla kummassa tahansa juosteessa, ja sen jako tripleteihin voi alkaa molemmissa kolmesta eri paikasta

Tässä voisi siis olla peräti kuusi erilaista lukutapaa, joista vain yksi lienee oikea (aloituskodonia ei etsitä, vaan pelkästään stoppeja)



5' - TTATTTTATTTTCGATAATTCGACCTTAAACGCGAAACTTCACTTAAC - 3'
3' - AATAAAATAAAGCTATTAAGCTGGAATTTGCGCTTTGAAGTGAATTG - 5'



Ensimmäinen *kodoni* voi siis olla

>>> UUA, UAU tai AUU >> TEMPLAATTINA ALEMPI JUOSTE
tai

<<< GUU, UUA tai UAA <<< TEMPLAATTINA YLEMPI JUOSTE

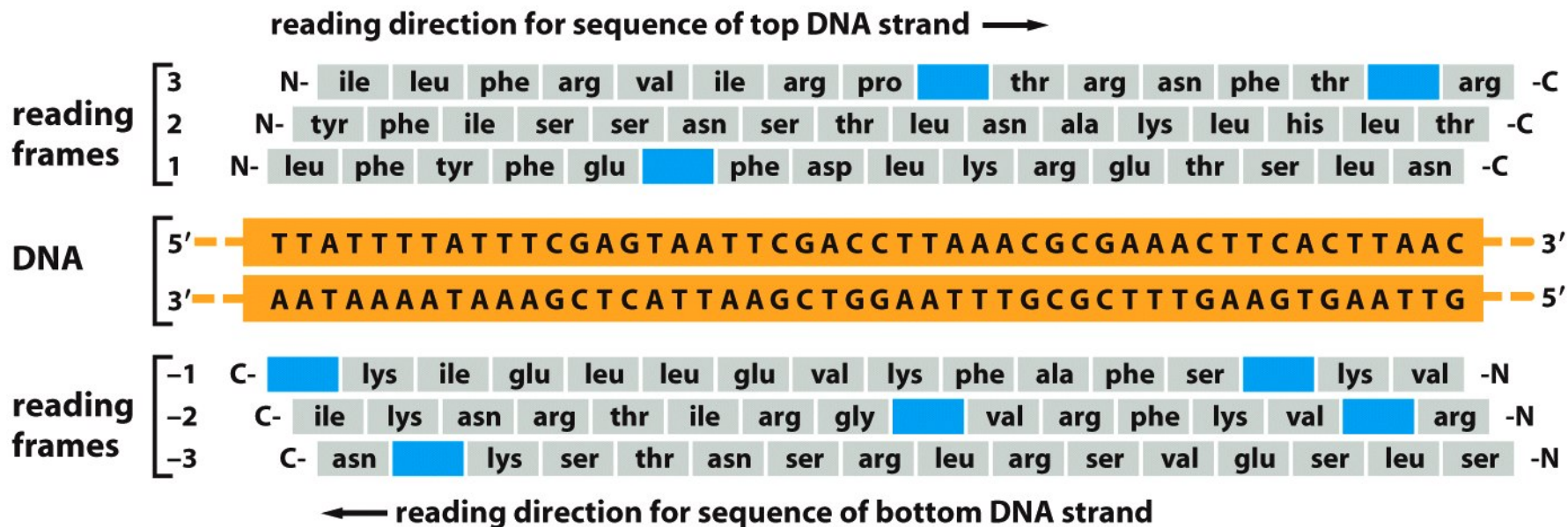


Figure 8-52a Molecular Biology of the Cell 5/e (© Garland Science 2008)

STOP on merkitty sinisellä laatikolla (UAA, UAG, UGA).

Avain lukemiseen:

leu UUA, UUG, CUU, CUC, CUA, CUG

Mikä siis on geeni? Alempi vai ylempi juoste?

CELL 550

180

190

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

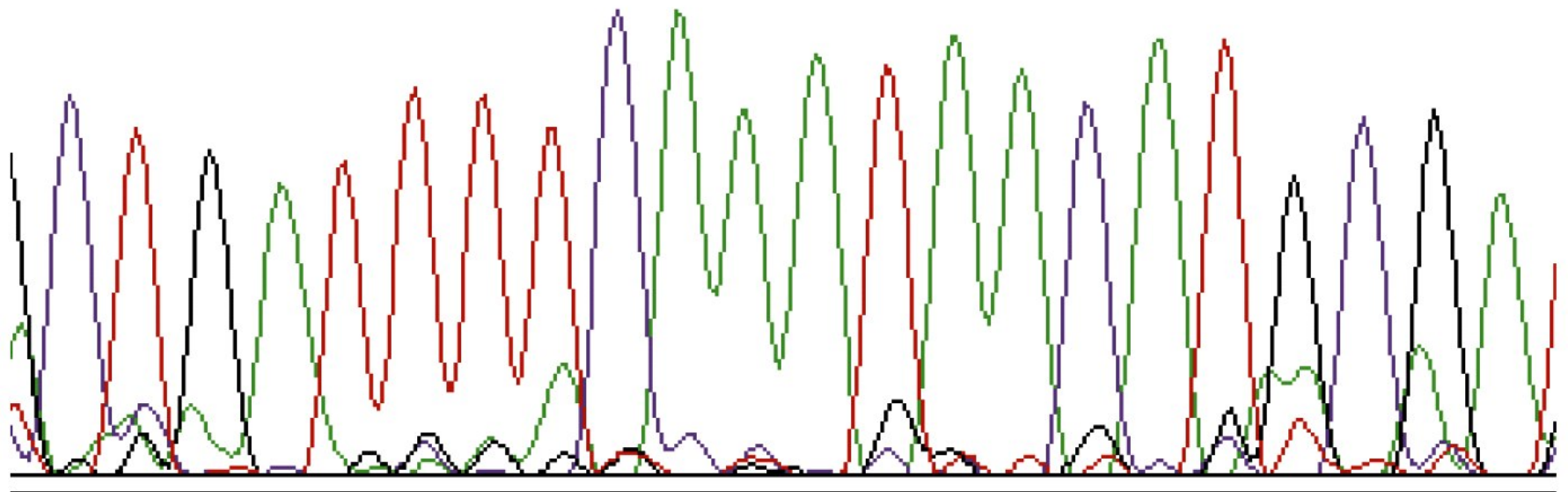


Figure 8-51 Molecular Biology of the Cell 5/e (© Garland Science 2008)

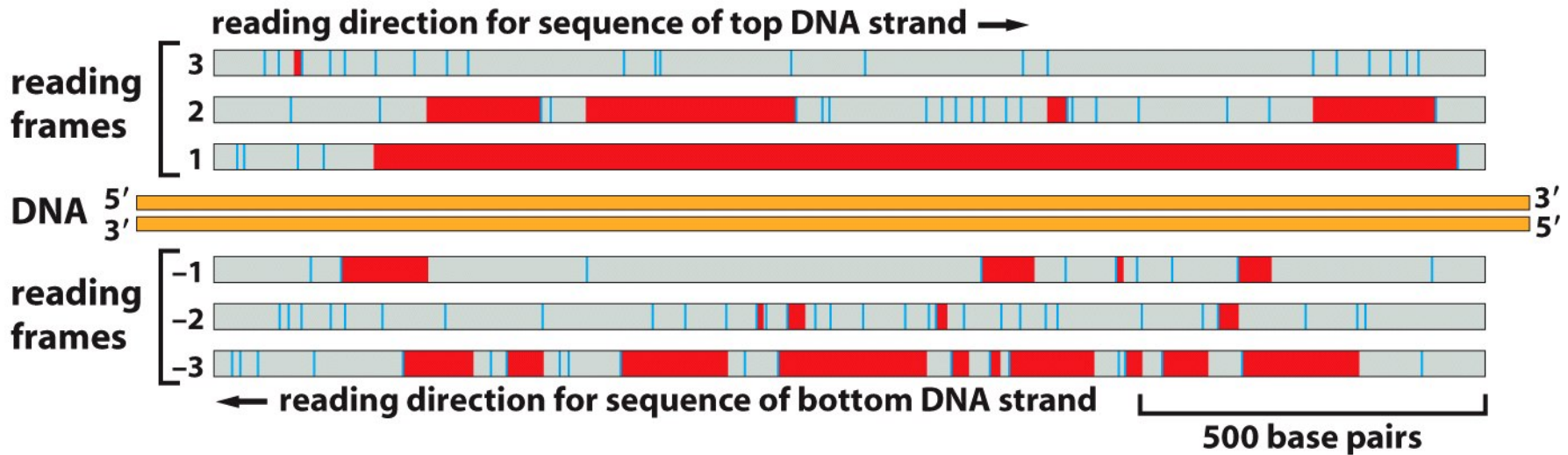


Figure 8-52b Molecular Biology of the Cell 5/e (© Garland Science 2008)

Tässä esimerkissä on tarkasteltu 1700 nukleotidia. STOP on merkitty vihreällä viivalla, ja mahdolliset START-STOP -välit on väritetty punaisella.

Ainoastaan FRAME 1 näyttää sisältävän pitkän jakson, joka *voi* olla polypeptidiketjun rakennusohjetta.

Tätä sanotaan **ORF**: iksi (open reading frame) ja se on geenin ennustamisen ensimmäinen askel.

ORF:in ennustaminen jos matriksia on vähän enemmän

***Gyrodactylus arcuatus*in mtDNA (CO1)**

Ensimmäinen TÄYSI kodoni ATT

Ensimmäinen TÄYSI kodoni ATT

Huonolta näyttää. Harmaat tähdet on stoppeja

[illegible]

M4: Alignment Explorer (C:\Documents and Settings\lumme\Desktop\MANUSCRIPTS\mtDNA phylogeography of arcuatus\arcuatus mtDNA EDITED.mas)

Data Edit Search Alignment Web Sequencer Display Help

DNA Sequences Translated Protein Sequences

1LV16 BtA1 (L-Varjakka) DQ078701	-	Y	S	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
V2 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
10LV1 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
18LV11 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
1LV18 BtA2 (L-Varjakka) DQ078702	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
2LV23 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
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3LV25b BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
4LV30 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
6LV41 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
24LV65 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
41LV101 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
41LV102 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
42LV108 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
42LV110 BtA2 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
V6 BtA4 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
12LV2 BtA4 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
12LV3 BtA4 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
20LV13 BtA4 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
2LV21 BtA4 (L-Varjakka) DQ078704	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
4LV29 BtA7 (L-Varjakka) DQ078707	-	Y	L	E	F	D	*	V	V	V	D	E	H	S	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	A	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H
4LV31 BtA8 (L-Varjakka) DQ078708	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
3LV25a BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
3LV27 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
4LV32 BtA9 (L-Varjakka) DQ078709	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
5LV34 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
5LV35 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
5LV37 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
7LV46 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T	A	E	Q	V	*	T	C	Y	C	S	L	Y	I	Y	Q	V	F	L	E	Y	L	V	L	Y	L	Y	V	L	Y	L	V	L	E	V	Y	L	L	M	L	V	I	D	D	*	M	Y	E	H	I	Y	L	H	
40LV100 BtA9 (L-Varjakka)	-	Y	L	E	F	D	*	V	V	V	D	E	H	F	I	H	H	Y	Q	G	L	H	I	T</																																																					

[illegible]

Fenylalaniinilla alkaa, stoppeja ei ole, variaatiota vähän. Tämä kehys OK.

tagctatgactaccaccctaattacccacattattaacccccctcgcatatatcggtccccg ttcttttagcagtcgcttttcctca
ccttac tcgaacgaaa agtccttgga tatatgcaacttcggaaagggccaaacatcgctcggtccatacggattgcttca
acctatcgcgagcggcctaactattcattaaagaaccagttcgaccgtccacctcctcccccttcctatttctcgctaca
cccatacttgcccttacgcttgactaaccctgtgagcccccatgcccatcccttaccaccattacagacctaataatctcggg
gtactatttgtcctcgcactttccagcctagccgtgtattctattttaggctcagggtagcttcaaattctaaatatgcccta
tgagctctacgagcagtggcacaaaccatttctacgaagtcagccttgactaatcttactcagcgtaattatctttacgg
gggatttacactacaaaccttcaatgtagcccaagaaagcatctgactactcgtagcagcctgaccccttgccgccatat
ggta tatctctact ctagctgaaa

4441 caaacgtgc accctttgac cttacagaag gagaatcaga attagtctcc ggatttaatg
4501 tagaatacgc cggagggccc ttcgccctct tctttctagc cgaatacgt aatatccttc
4561 taatgaacac actctcggcc attctatttt taggcgcac ccacatcccc gccttccccg
4621 aattaacagc cgtaaacctataaacaagg ccgccctcct ctccgttgta tttttatgag
4681 tacgagcctc ctaccacga tttcgctacg accaactcat acatttagtt tgaaaaagct
4741 tcttaccttt gacactagcc cttgtcctat gacacctagc acttccaacc gcaatggcag
4801 gcctccctcc ccaacttta gcccgaagg attgtgcctg aatgtttaag gaccaccttg

[NCBI homepage](http://www.ncbi.nlm.nih.gov/)

Jonkun otuksen jonkun geenin aminohapposekvenssiä??

MTTTLITHIINPLAYIVPVLLAVAFLTLLERKVLGYMQLRKG
PNIVGPYGLLQPIADGLKLFIKEPVRPSTSSPFLFLATPMLALT
LALTLWAPMPIPYPITDLNLGVLFVLALSSLAVYSILGSGWAS
NSKYALIGALRAVAQTISYEVSLGLILLSVIIFTGGFTLQTFN
VAQESIWLLVPAWPLAAMWYISTLAETNRAPFDLTEGESEL
VSGFNVEYAGGPFALFFLAEYANILLMNTLSAILFLGASHI
PAFPELTAVNLMTKAALLSVVFLWVRASYPRFRYDQLMHLV
WKSFLPLTLALVLWHLALPTAMAGLPPQ L

????????????????

[NCBI homepage](https://www.ncbi.nlm.nih.gov/)

Geneettinen koodi on “universaali”

Oleellisesti samaa koodia, *pienin* muutoksin, käyttävät arkit, bakteerit, eukaryootit ja vielä mitokondriot ja kloroplastitkin

Eukaryootin tuman geenit ja mitokondrion geenit luetaan eri koodilla, siksi mitokondriossa on oma rRNA (2) ja 22 tRNA:ta (**TRANSLAATIOKONEISTO**)

Esim. UGA on universaalisti STOP, mutta mitokondriossa se on tryptofaanin koodi, ja AGA ja AGG taas ovat yleensä arginiinin koodina, mutta nisäkäsmitokondriossa ne luetaan STOP.

The Genetic Code

1st position (5' end) ↓	U	C	A	G	3rd Position (3' end) ↓
U	Phe Phe Leu Leu	Ser Ser Ser Ser	Tyr Tyr STOP STOP	Cys Cys STOP Trp	U C A G
C	Leu Leu Leu Leu	Pro Pro Pro Pro	His His Gln Gln	Arg Arg Arg Arg	U C A G
A	Ile Ile Ile Met	Thr Thr Thr Thr	Asn Asn Lys Lys	Ser Ser Arg Arg	U C A G
G	Val Val Val Val	Ala Ala Ala Ala	Asp Asp Glu Glu	Gly Gly Gly Gly	U C A G

Phe	171 UUU	AAA 0	Ser	147 UCU	AGA 10	Tyr	124 UAU	AUA 1	Cys	99 UGU	ACA 0
	203 UUC	GAA 14		172 UCC	GGA 0		158 UAC	GUA 11		119 UGC	GCA 30
Leu	73 UUA	UAA 8	Ser	118 UCA	UGA 5	stop	0 UAA	UUA 0	stop	0 UGA	UCA 0
	125 UUG	CAA 6		45 UCG	CGA 4	stop	0 UAG	CUA 0	Trp	122 UGG	CCA 7
Leu	127 CUU	AAG 13	Pro	175 CCU	AGG 11	His	104 CAU	AUG 0	Arg	47 CGU	ACG 9
	187 CUC	GAG 0		197 CCC	GGG 0		147 CAC	GUG 12		107 CGC	GCG 0
	69 CUA	UAG 2		170 CCA	UGG 10	Gln	121 CAA	UUG 11		63 CGA	UCG 7
	392 CUG	CAG 6		69 CCG	CGG 4		343 CAG	CUG 21		115 CGG	CCG 5
Ile	165 AUU	AAU 13	Thr	131 ACU	AGU 8	Asn	174 AAU	AUU 1	Ser	121 AGU	ACU 0
	218 AUC	GAU 1		192 ACC	GGU 0		199 AAC	GUU 33		191 AGC	GCU 7
	71 AUA	UAU 5		150 ACA	UGU 10	Lys	248 AAA	UUU 16	Arg	113 AGA	UCU 5
Met	221 AUG	CAU 17		63 ACG	CGU 7		331 AAG	CUU 22		110 AGG	CCU 4
Val	111 GUU	AAC 20	Ala	185 GCU	AGC 25	Asp	230 GAU	AUC 0	Gly	112 GGU	ACC 0
	146 GUC	GAC 0		282 GCC	GGC 0		262 GAC	GUC 10		230 GGC	GCC 11
	72 GUA	UAC 5		160 GCA	UGC 10	Glu	301 GAA	UUC 14		168 GGA	UCC 5
	288 GUG	CAC 19		74 GCG	CGC 5		404 GAG	CUC 8		160 GGG	CCC 8

Figure 34 The human genetic code and associated tRNA genes. For each of the 64 codons, we show: the corresponding amino acid; the observed frequency of the codon per 10,000 codons; the codon; predicted wobble pairing to a tRNA anticodon (black lines); an unmodified tRNA anticodon sequence; and the number of tRNA genes found with this anticodon. For example, phenylalanine is encoded by UUU or UUC; UUC is seen more frequently, 203 to 171 occurrences per 10,000 total codons; both codons are expected to be decoded by a single tRNA anticodon type, GAA, using a G/U wobble; and there are 14

tRNA genes found with this anticodon. The modified anticodon sequence in the mature tRNA is not shown, even where post-transcriptional modifications can be confidently predicted (for example, when an A is used to decode a U/C third position, the A is almost certainly an inosine in the mature tRNA). The Figure also does not show the number of distinct tRNA species (such as distinct sequence families) for each anticodon; often there is more than one species for each anticodon.

Kaikki kodonit eivät ole yhtä suosittuja