



## List of amino acids and their codons

List of amino acids and mrna codons. What are the 20 amino acids and their codons.



TABLE 3-1	20	Primary A	mino Acids in the	Genetic C	ode		
	ABBRE	VIATION		ABBRE	VIATION		
Amino Acid	3-Letter	1-Letter	Amino Acid	3-Letter	1-Lette		
Alanine	Ala	A	Leucine	Leu	L		
Arginine	Arg	R	Lysine	Lys	K		
Asparagine	Asn	N	Methionine	Met	Μ		
Aspartic acid	Asp	D	Phenylalanine	Phe	F		
Cysteine	Cys	С	Proline	Pro	Р		
Glutamic acid	Glu	E	Serine	Ser	S		
Glutamine	Gln	Q	Threonine	Thr	Т		
Glycine	Gly	G	Tryptophan	Trp	W		
Histidine	His	н	Tyrosine	Tyr	V		



Standard RNA Codon table arranged in a circle. The codon table can be used to convert the genetic code to the amino acid order. 1f661e1667c31.pdf The standard genetic code is traditionally displayed as a RNA Codon table, because when proteins are produced in the ribosome cell, it is the rapporteur RNA (MRNA) that guides the synthesis of protein. MRNA sequence is determined by the Genome DNA sequence. [4] In this context, the standard genetic code is called the translation table 1. [3] It can also be displayed in the DNA codons in the following tables occurDNA chains and are arranged in the direction from 5 strokes to 3 beats. Depending on the source of the genetic code, such as the cell nucleus, mitochondria, plastide or hydrogenosomes, various alternative codon tables are used. The genetic code contains 64 different codes in the following tables; Most define the amino acid. [6] Three sequences, UAG, UGA and UAA, called Stop code [1. Note], does not cod the amino acid, but rather a signaling the release of emerging ribosomal polypeptide. In the standard code, Aug (read as methionine) can be used as a starting codon and, in combination with sequences, as the initiation factor initiates the translation. [3] [8] [9] In rare cases, boot codes may also be GUG or UUG standard code; These codons usually mark valine and leucine, but are translated into methionine or formionin as a starting codon. The first table - the standard table - can be used to examine the appropriate amino acid or the appropriate signal if it is the beginning or end of the three nucleotide translation. The second Table, in the opposite way, works in the opposite way, works in the opposite direction: it allows you to set a possible triplet code if the amino acid is known. Because many codons can encode the same amino acid, in some cases the name of the nucleic acid of international and applied chemistry (IUPAC) is given. Translation Table 1 Standard RNA COTON TABLE TABLE BIOKEMIC PROPERTIES OF NON -COLOR AMININE ACCOS - DISCLUSION: STOP CODON \* INIR: Possible code initial genetic code [1] [10] 1. Principle 1. Principle 2. Principle 3. Princi (Tyr/y) Tyrosine UGU (Cys/C) Cysteine UU UCC UCC UUA C GGC C GUA GCA GAA (GLU/E) GLUTAMO R QUEGŠTIS - GGA A GUG GCG GAG GGG G Rnr Inversijos Kodonų Lentelė Standartinė Genetinio Kodo inversijos Lentelė (Kodon's suspicion žumir žminjiMė rnr) Rnr rnr rnr) YRA SPAUSTI ALA, A GCU, GCC, GCA, GCG GCN Ile, I Auu, AUC, AUA AUH Arg, R CGU, CGC, CGA, CGG; AGA, ADD CGN, AGR; o CGY, Mgr Leu, L Cuu, Cuc, Cua, Cug; Uua, uug cun, uur; o cuy, yur asn, n aau, aac aay lys, k aaa, aag aar asp, d gav, gac gay met, mug asn o asp, b aau, aac; Gac ray phe, f uuu, uuc uuc cys, c ugu, ugc ugy pro, p ccu, ccc, ccg, ccg ccn gln, q caa, cag car ser, s ucu, ucc, uca, ucg; Agu, AGC UCN, AGY GLU, E GAA, GAG GAR THR, T ACU, ACC, ACA, ACG ACN GLN or GLU, Z CAA, CAG; GAA, GAA SAR TRP, Z UGG GLY, G GUG, GGK, GGK, GGY, GGY GGN tir, y uau, uak way ego, h ka, sas cai, v guu, guk, gua, gug pushka Ag Stop UAA, UAR Standard DNA DNA Codon Chart Biochemical properties of amino acids do not sting basal acid - end: stop codon \* inir: possible start standard genetic code [14] [14] [Note 3] [ (Leu/L) Leucine TCA TAA STOP (OCHRA) \*[Note 2] TGA STOP (OPAL OPAL) \*[Note 2] A TTG TCG TAG STOP (Yellow) \*[Note 2] TGG (TRP/W) TRPTOFAN G CTT CCT (Pro/P) Prolin Cat (His/H) histidine CGT (arg/R)) Arginine) Arginine T CTC CCC CGC C CA CAA CAA (GLN/Q) Glutamine CGA A CGA CG CG CGG GG G ATT ( Ile/i) Isoleucina Act (thr/t) Treonin Aat (asn/n) asparagin agt (ser/s))) serinin t atc Acc aac ac c ata aaa (lys/k) lysine aga (arg/r) Arginine A (Met/M) MessininAAG AGG G G GTT (Val/V) Valine GCT (Ala/A) Alanine GAT (Asp/D) Aspartic acid â GGT (Gly/G) Glycine T GTC GCC GAC GGC C GTA GCA (Glu/E) Glutamic acid â GGA A GTG GCG GAG GGG G Table of Inverted Standard Genetic Codes (compressed using IUPAC notation)[13] Truncated Amino Acid DNA Codons Truncated Amino Acid DNA Codons Ala, A GCT, CGC, CGA, CGG; AGA, AGG, CGN, EGR; or CGY, MGR Leu, L CTT, CTC, CTA, CTG; TTA, TTG, CTN, TTR; or CTY, YTR Asn, N AAT, AAC AAY Lys, K AAA, AAG AAR Asp, D GAT, GAC GAY Met, M ATG Asn or Asp, B AAT, AAC; GAT, GAC RAY Phe, F TTT, TTC TTY Cys, C TGT, TGC, TCA, TCG; AGT, AGC TCN, AGY Glu, E GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN Gln or Glu, Z CAA, CAG; GAA, GAG, GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn or Glu, Z CAA, CAG; GAA, GAG, GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn, Q CAA, CAG CAR Ser, S TCT, TCC, TCA, TCG; AGT, AGC TCN, AGY Glu, E GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn or Glu, Z CAA, CAG; GAA, GAG, GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn or Glu, Z CAA, CAG; GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn or Glu, Z CAA, CAG; GAA, GAG GAR Thr, T ACT, ACC, ACA, ACG ACN GIn or Glu, Z CAA, CAG SAR Trp, W TGG Gly, G GGT, GGC, GGA, GGG GGN Tyr, Y TAT, TAC TAY His, H CAT, CAC CAY Val, V GTT, GTC, GTA, GTG GTN START ATG STOP TAA, TGA and the same start and start at a sta amino acid regardless of the organism or the source. However, there is consensus that the genetic code evolves[17] so that a codon is translated differently depending on the genetic source. [16][17] For example, in 1981, it was found that the usage of AUA, UGA, AGA, and AGG codons in the mammalian mitochondrial coding system differed from the universal code,[16] Stop codons may also be affected: the ciliated universal stop codons UAA and UAG code for glutamine,[17][4. Note] The following table lists these alternative codons. Biochemical properties of amino acids Polar nonpolar basic acid Termination: stop codon \*Comparison of codon translations using alternative and standard genetic codes[3]The standard code for standard 1 is Table 8 (plant chloroplasts). Mitochondrial circle 2 Aga aga stop \* arg (r) agg agg stop \* arg (r) agg agg stop \* arg (r) ata aua Met (M) ILE (I) (i) ctt cuu thh (t) leu (l) ctc cuc throp (t) leu (l) cta cua thu (t) leu (l) ctg cug Thr (Stop \* CGA CGA CGA No Arg (R) CGC CGC No ARG (R) Form, produced and mitochondrial + Mycoplasma 4 TGA UGA TRP (W) Stop \* includes Translation Table 7 (kinetoplasty) Mitochondria -vertebrate 5 Aga Aga Sers (S) Arg (R) Ata Aua Met (M) Ile (I) TGA UGA ZG (W) Stop \* ciliatansln (q) Stop \* Tag Uag Gln 1 Translation Table Alternate Nuclear Yeast 12 CTG CUG SERS (S) LEU (L) ASCIDIC MITOCHondria 13 Aga Aga Gly (G) Arg (R) Agg Agg Gly (G) Arg (R) Ata Au Met (M) ILE (I) TGA UG (UG (G) Arg (R) Ata Au Met (M) ILE (I) TGA UG (UG (G) Aga (R) Au Met (M) ILE (I) TGA UG (I) ILE (I) TGA UG (I blekarisma 15 day UAG GLN(q)stop\*as of November 2016 18: Not in NCBI update. As for Translation Table 6 Table 6



Chlorophycean mitochondrial 16-day uag leu (l) stop \* trematodo mitochondrial 21 tga UGA suffer tca uca stop \* cheese (s) tag uag leu (l) stop \* thrustochytrium mitochondrial 23 TTA UUA Stop \* leu (l) Similar to translation table 11. Pterobranchia Mitochondrial 24 Aga Aga ser (S) Arg (R) Agg Agg Lys (K) Arg (R) TGA UGA TRP (W) Stop \* Candidate Section SR1 and 25 TGA UGA Gly (G) Bond \* Nuclear Tanophilic Pachyzolene 26 CTG CUG Ala (A) Leu (L) Nuclear Caryotene 27 TAA UAA Gln (Q) Bond \* TG UGA Stop \* or Trp (W) ) Alloy \* Condylostoma Nuclear Alloy 28 TAA UAA \* or Gln Alloy (Q) \* TAG UAG Stop \* or Gln Alloy (Q) \* TGA UGA Alloy \* or Trp Alloy (W) \* Nuclear Mesodinium 29 TAA UAA Tyr Alloy (Y) \* TAG UAG Tyr (Y) Alloy \* Nuclear Peritrich 30 TA SAU Glu (E) - Stop \* TAG UAG Glu (E) - Stop \* TAG UAG Stop \* or Glu (E) - Stop \* TAG UAG Stop \* or Glu (E) - Stop \* TGA UGA Trp (W) Stop \* Cephalodiscidae Mitochondrial Code 33 AGA AGA Ser (S) Arg (R) TAA UAA Tyr (R) TGA UGA Trp (W) Stop \* Cephalodiscidae Mitochondrial Code 33 AGA AGA Ser (S) Arg (R) TAA UAA Tyr (Y) Stop \* TGA UGA Trp (W) Stop \* TGA UGA Glu (E) - Stop \* TAG UAG Glu (E) - Stop \* TGA UGA Glu

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I-Letter Abbreviation	<b>3-Letter Abbreviation</b>	Amino Acid	Numerical Codons
A	Ala	Alanine	421, 422, 423, 424
В	-	_	_
С	Cys	Cysteine	141, 142
D	Asp	Aspartic acid	431, 432
E	Glu	Glutamic acid	433, 434
F	Phe	Phenylalanine	111,112
G	Gly	Glycine	441, 442, 443, 444
Н	His	Histidine	231, 232
I	lle	Isoleucine	311, 312, 313
J	Stop	Opal	143
K	Lys	Lysine	333, 334
L	Leu	Leucine	113, 114, 211, 212, 213, 214
M	Met (Start)	Methionine	314
N	Asn	Asparagine	331, 332
0	-	-	_
Р	Pro	Proline	221, 222, 223, 224
Q	Gln	Glutamine	233, 234
R	Arg	Arginine	241, 242, 243, 244, 343, 344
S	Ser	Serine	121, 122, 123, 124, 341, 342
Т	Thr	Threonine	321, 322, 323, 324
U	_	_	
V	Val	Valine	411, 412, 413, 414
W	Trp	Tryotophan	144
X	Stop	Ochre	133
Y	Tyr	Tyrosine	131, 132
Z	Stop	Amber	134

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