Genetic Code

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Genetic Code

- A triplet sequence of nucleotides on the mRNA is the codon for each amino acid.
- 64 (4³) different codon
- Nirenberg got Nobel price in 1968 for this
- 31 tRNA carrying 20 amino acids which translate
 61 codons

First nucleotide 5' end	Second nucleotide				Third nucleotide 3' end
	U	C	Α	G	
U	Phe	Ser	Tyr	Cys	U
	Phe	Ser	Tyr	Cys	C
	Leu	Ser	Stop	Stop	Α
	Leu	Ser	Stop	Trp	G
C	Leu	Pro	His	Arg	U
	Leu	Pro	His	Arg	C
	Leu	Pro	Gln	Arg	A
	Leu	Pro	Gln	Arg	G
A	lle	Thr	Asn	Ser	U
	lle	Thr	Asn	Ser	C
	lle	Thr	Lys	Arg	Α
	Met	Thr	Lys	Arg	G
G	Val	Ala	Asp	Gly	U
	Val	Ala	Asp	Gly	C
	Val	Ala	Glu	Gly	Α
	Val	Ala	Giu	Gly	G

Features of Genetic Code

- Triplet Codons
 - Codons are on mRNA
- Non-overlaping
 - The codons are read one after another in a continuous manner
 - Starting point is important
- Non-punctuated
 - There is no punctuation between codon they are continuous

Degenerate

- 61 codons and 20 amino acids
- So one amino acid has more than one codon
- If amino acid has more than one codon, first two bases in the codon will be the same, only third one is different, this reduces the effect of mutation.

Unambiguous

One codon stands only for one amino acid

Universal

Same in all species

Wobbling phenomenon

- The reduced stringency between the third base of the codon and the complementary nucleotide in the anticodon is called wobbling
- The pairing of codon and anticodon can wobble at the third letter
- GGU, GGC, GGA for glycine can pair with CCI of glycine tRNA
- Reduce possibility of mutation effect

Terminator codons

- Three codons do not code for any amino acids
- Nonsense codons
- Punctuator codon
- End of protein synthesis
- UAA, UAG, UGA
- UGA some time code for seleno-cysteine (21st amino acid)

Initiator codon

- AUG in most cases. Also code for metheonine
- GUG in some case

Mitochondria

- Only 22 tRNA
- UGA (standard stop codon) is read as Trp
- AGA and AGG (standard codons for Arg) are read as stop codons

Thank You