

- alternating sequence of exon and intron
- **immediate product of transcription**
- composed of two components: **Exon and Intron**
- **Intron- uncoded region** or intervening region, spliced, removed to produce mRNA
- **Exon- coded region**, coding region, gene
- **DNA Helicase-** to open or unwind the two DNA strand
- **Topoisomerase-** to straighten the non-formation, or the wavy/twist in the strand during the process of transcription, to prevent topographical non-formation (buhol)
- **snRNA-** small nuclear RNA, aka **spliceosome ribonuclear**, it removes the intervening or splice the introns.
 - snRNA + small ribonucleic protein= **spliceosome**
- **Ligase-** enzyme that joins the Exons after removing the introns
 - ligates the Exon with one another
 - molecular glues
- **Post-transcriptional-** removal of introns

TYPES OF RNA → PROTEINS TRANSLATION

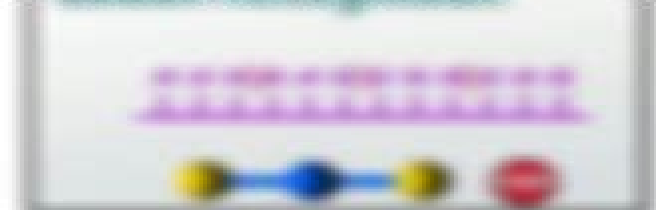


BASE CODON/ SECONDARY BASES

	U	C	A	G	
U	UUU } Phe UUC } UUA } Leu UUG }	UCU } Ser UCC } UCA } UCG }	UAU } Tyr UAC } UAA } Stop UAG } Stop	UGU } Cys UGC } UGA } Stop UGG } Trp	U C A G
C	CUU } Leu CUC } CUA } CUG }	CCU } Pro CCC } CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } Arg CGC } CGA } CGG }	U C A G
A	AUU } Ile AUC } AUA } AUG } Met	ACU } Thr ACC } ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }	U C A G
G	GUU } Val GUC } GUA } GUG }	GCU } Ala GCC } GCA } GCG }	GAU } Asp GAC } GAA } Glu GAG }	GGU } Gly GGC } GGA } GGG }	U C A G

- **Secondary bases-** used for translation process
- has **unacil**, this is what mRNA contains in order to be translated into protein, that is why it is called **secondary** (primary- DNA to mRNA)
- 64 base codons
- **start codon-** AUG (Met)
- **stop codon-** UGA, UAG, UAA (no corresponding amino acids)
- UCAG- u can always give.... a smile

Codon Recognition



Quick Review Protein Synthesis And The Genetic Code

Quick Review Notes

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