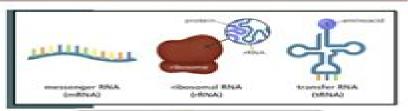
- alternating sequence of exon and intron
- immediate product of transcription
- o composed of two components: Exon and Intron
- Intron- uncoded region or intervening region, spliced, removed to produce mRNA
- · Exon- coded region, coding region, gene
- DNA Helicase- to open or unwind the two DNA strand
- Topoisomerase- to straighten the non-formation, or the wavy/hwist 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



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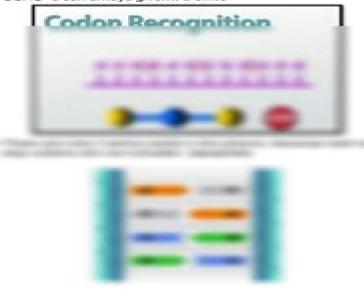
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THE RESIDENCE AND ADDRESS OF THE PARTY OF TH

BASE CODON/ SECONDARY BASES

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v	UUUC } Phe UUA UUG } Leu	UCC UCA UCG	UAC Tyr UAC Stop UAG Stop	UGU Cys UGC Stop UGA Stop	DUAG
c	CUC CUA CUG	CCC CCA CCG	CAU His CAC His CAA Gin	CGC CGA CGG	DOAG
^	AUC AUA Met	ACC ACA ACA ACG	AAC } Asn AAC } Lys AAG } Lys	AGU }Ser AGA }Arg AGG }Arg	2040
o	GUC GUA GUG	GCU GCC GCA GCG	GAC Asp GAA GAG GAU	GGC GGA GGG	0000

- . Secondary bases- used for translation process
- has uracil, 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



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Quick Review Protein Synthesis And The Genetic Code Quick Review Notes:

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