## **CONCEPT:** EUKARYOTIC POST-TRANSCRIPTIONAL REGULATION

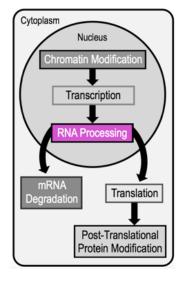
<ul><li>Eukary</li></ul>	votes regulate de	ene expression a	at the post-trans	scriptional level in	ways:

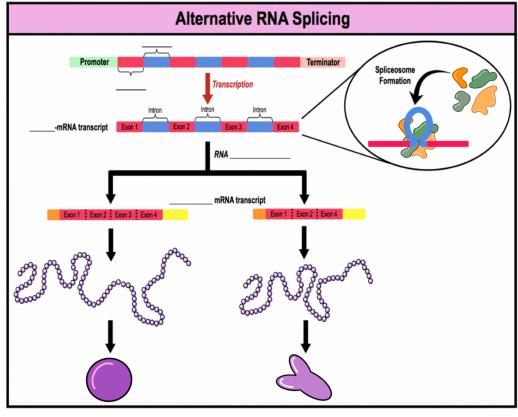
- 1 Alternative RNA \_\_\_\_\_ results in different protein products from the same mRNA transcript.
  - 2) RNA processing adds a 5' \_\_\_\_\_ & poly-\_\_\_\_ tail to mRNA for protection from RNA degrading enzymes.
  - 3 mRNA can be tagged for degradation or transcription is blocked by small noncoding \_\_\_\_\_ molecules.

# 1) Alternative RNA Splicing

- Recall: Eukaryotes require \_\_\_\_\_-transcriptional modifications like RNA splicing which can alter gene expression.
  - □ **Alternative splicing**: when different mRNA molecules are produced from the \_\_\_\_\_ premature RNA.
  - □ The \_\_\_\_\_: the RNA-protein complex that removes introns from premature RNA.

**EXAMPLE:** Alternative RNA splicing results in different protein products from the same premature RNA.





**PRACTICE** Alternative RNA splicing has been estimated to occur in more than 95% of multi-exon genes. Which of the following is **not** an evolutionary advantage of alternative RNA splicing?

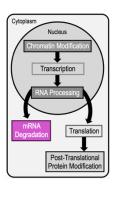
- a) Alternative RNA splicing increases diversity without increasing genome size.
- b) Different protein variants can be expressed by the same gene in different tissues.
- c) Alternative RNA splicing creates shorter mRNA transcripts.
- d) Different protein variants can be expressed by the same gene during different stages of development.

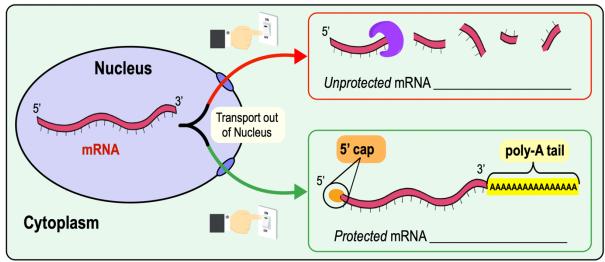
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## 2) mRNA Protection in the Cytoplasm

- •mRNA transcripts must be transported to the \_\_\_\_\_\_ where they can be translated by ribosomes.
  - ☐ The cytoplasm has many RNA degrading enzymes destroy *foreign* viral RNA molecules.
- •The \_\_\_\_\_' cap & poly-A \_\_\_\_\_ of mRNA molecules \_\_\_\_\_\_ the mRNA from degradation by enzymes.

**EXAMPLE:** mRNA is protected from degradation by cytoplasmic enzymes with a 5' cap and poly-A tail.

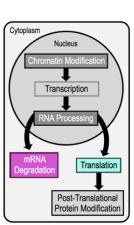


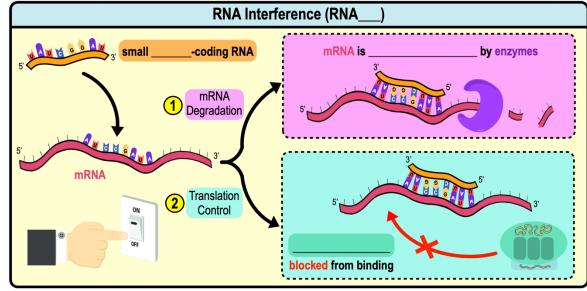


#### 3) RNA Interference

- •RNA \_\_\_\_\_\_ (RNA\_\_): process of small noncoding RNAs blocking translation of target mRNA molecules.
  - □ **Small noncoding RNA:** short strands of RNA that have a *complementary sequence* to their mRNA target.
- •There are \_\_\_\_\_ possible scenarios that turn gene expression OFF:

**EXAMPLE:** RNA Interference can block ribosome binding or recruit cellular enzymes for mRNA degradation.





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PRACTICE: Which of the following statements best describes the function of RNAi?

- a) Small RNA molecules interfere with translation by targeting ribosomes for degradation.
- b) Small DNA molecules interfere with mRNA molecules by blocking their ability to bind to a ribosome.
- c) Small RNA molecules interfere with translation by targeting specific tRNA molecules
- d) Small RNA molecules interfere with translation by blocking a target mRNA's ability to bind to a ribosome.

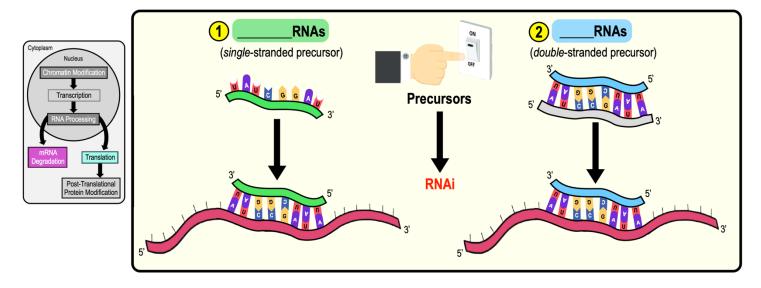
## **Types of Small Noncoding RNAs**

•There are \_\_\_\_ classes of RNAs involved in RNAi:

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- □ BOTH types bind to a target mRNA by complimentary base paring & turns \_\_\_\_\_ expression of that gene.
- •The only difference between microRNAs & siRNAs is the structure of their precursor form:
- □ mircoRNAs have a \_\_\_\_\_\_-stranded precursor & siRNAs have a \_\_\_\_\_-stranded precursor

**EXAMPLE:** RNA interference by two types of small noncoding RNAs (microRNAs & siRNAs).



## **PRACTICE**: Which of the following best describes siRNA?

- a) A short double-stranded RNA with one strand that can complimentarily bind to and inactivate an mRNA.
- b) A single-stranded RNA with internal complementary base pairs that allow it to fold into a cloverleaf pattern.
- c) A portion of rRNA which is a component of the large and small ribosomal subunits.
- d) A molecule, known as Dicer, that can degrade or cut RNA sequences.