

b. transcription factors (general and specific)

c. Enhancers (distal control elements)

d. Activators

i. What structural features seem to be common to the hundreds of activators identified?

e. mediator proteins

f. repressors

9. How do activators and repressors act indirectly to regulate gene expression? What are the effects of these indirect actions?

10. How do eukaryotes coordinate gene expression without operons? GIVE ALL POSSIBLE METHODS!!!!

11. What advantage(s) does post-transcriptional regulation have?

12. Describe alternative RNA splicing.

13. Prokaryotic mRNA is degraded rapidly. What does this allow?

a. How does this contrast to eukaryotic mRNA?

14. What may be responsible for the degradation of mRNA?

15. Initiation of translation presents another opportunity to regulate gene expression. Explain each of the following.
- a. Regulatory proteins and UTRs
 - b. short poly-A tails
 - i. What is so amazing about this?!?!?!
 - c. “global” control
16. Post-translational opportunities for regulation include protein processing and degradation.
- a. In what ways are proteins processed?
 - b. Explain “selective degradation.”
 - i. How are proteins marked for destruction? (include ubiquitin)

Noncoding RNAs play multiple roles in controlling gene expression

17. Study figure 15.13. How are microRNAs and/or small-interfering RNAs involved gene regulation?

Researchers can monitor expression of specific genes

18. How can you use to nucleic acid hybridization to identify whether the bacterial colonies contain the gene of interest?
- a. Briefly explain how the procedure works (study figure 15.14)
19. Explain how reverse transcriptase is used to produce cDNA.
20. What is a genomic library?
21. How can you detect the presence of a gene of interest without using DNA libraries?