

Outline 4

IV. Control of Gene Expression in Eukaryotes

A. Why is it different than in Prokaryotes?

B. Levels of control of gene expression

1. Chromosomal level – gene amplification

2. Chromatin modification

DNA methylation

histone acetylation

3. Transcription

transcription factors

enhancer sequences

4. Post-transcriptional control

5. Control of translation

6. Post-translational control

Patterns of control of gene expression

Negative control - an active regulatory protein turns transcription OFF

Induction - signal molecule makes the regulatory protein active

Repression - signal molecule makes the regulatory protein inactive

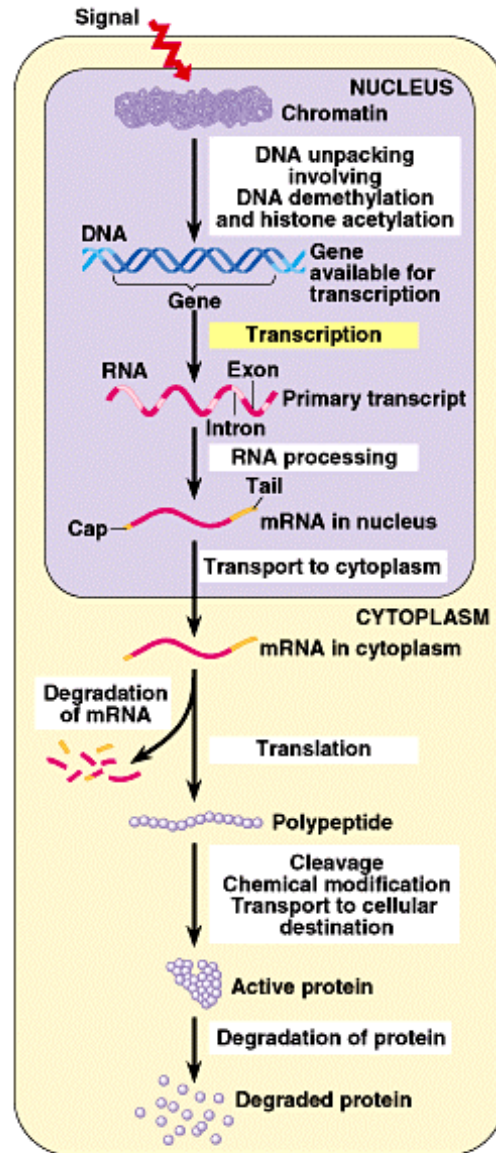
Positive control - an active regulatory protein enhances the rate Of transcription

Induction - signal molecule makes the regulatory protein active

Repression - signal molecule makes the regulatory protein inactive

Fig. 19.7

Levels of control of gene expression in Eukaryotes

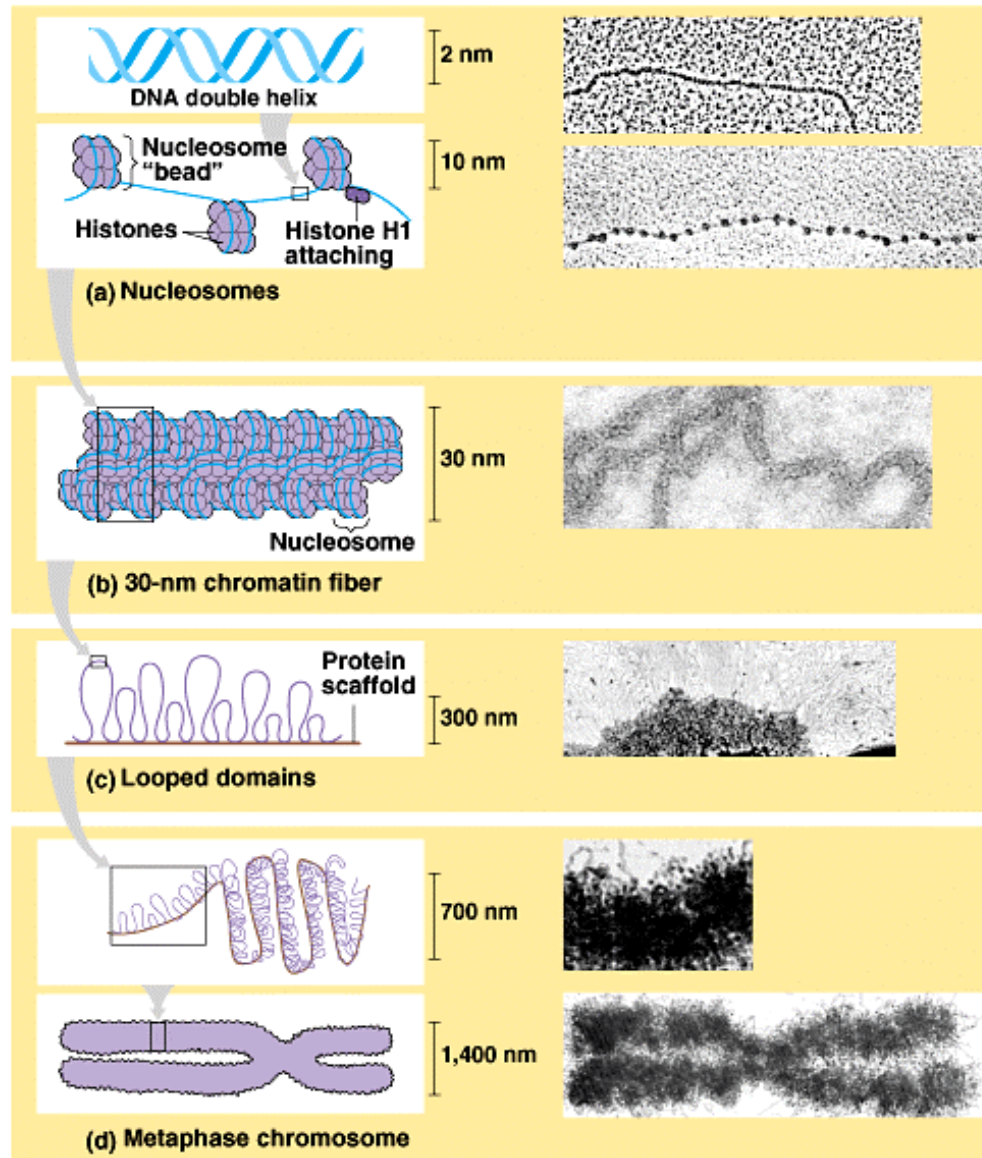


Control of Gene Expression

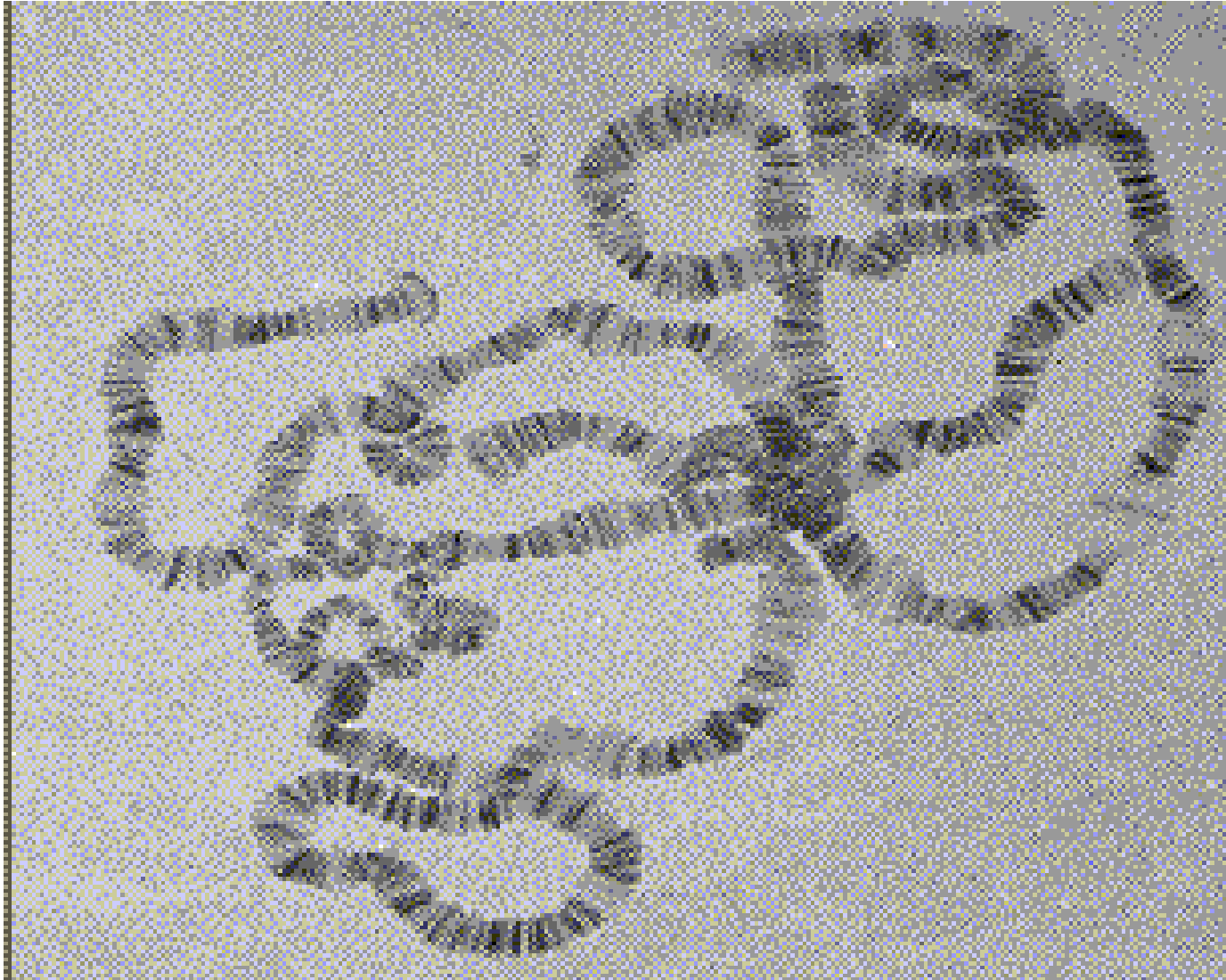
| <u>Level of Control</u> | <u>Control Mechanism</u> |
|-------------------------|---|
| PROKARYOTES | |
| Transcriptional | <ol style="list-style-type: none"> 1. <u>Negative Induction</u>: e.g. lac operon 2. <u>Negative repression</u>: e.g. Trp operon 3. <u>Positive Induction</u>: e.g. lac operon enhanced at low [glucose] 4. <u>Positive repression</u>: e.g. lac operon repression in the presence of glucose |
| EUKARYOTES | |
| Chromosomal/DNA | <ol style="list-style-type: none"> 1. <u>Gene amplification</u>: e.g. polytene chromosomes 2. <u>Chromatin modification</u> <ol style="list-style-type: none"> a. <u>DNA methylation</u>: inactivation of genes by methylation (e.g. Barr bodies) b. <u>Histone acetylation</u>: weakening of histone bonds by addition of acetyl groups |
| Transcriptional | <ol style="list-style-type: none"> 1. <u>Control elements</u>: DNA sequences (e.g. enhancers) that interact with proteins (e.g. activators, transcription factors) to enhance binding of RNA polymerase |
| Post-transcriptional | <ol style="list-style-type: none"> 1. <u>RNA processing</u>: intron removal and exon splicing 2. <u>Nucleotide additions</u>: poly-A tail can prevent mRNA translation or transportation across nuclear membrane 3. <u>mRNA degradation</u>: different mRNAs may have different lifetimes |
| Translational | <ol style="list-style-type: none"> 1. <u>Initiation factors</u>: proteins required to initiate ribosome binding and tRNA binding to mRNA |
| Post-translational | <ol style="list-style-type: none"> 1. <u>Polypeptide modifications</u> 2. <u>Metabolic regulation</u> of gene product: |

Fig. 19.1

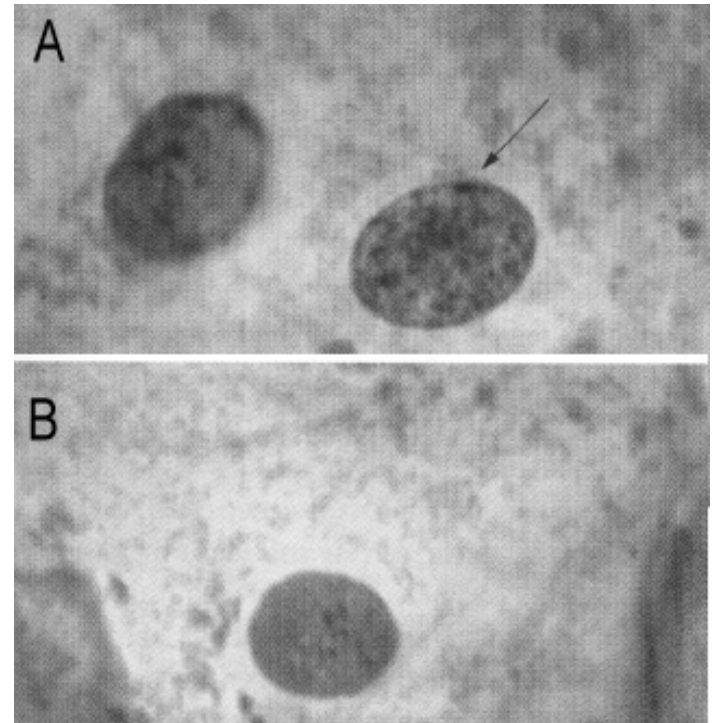
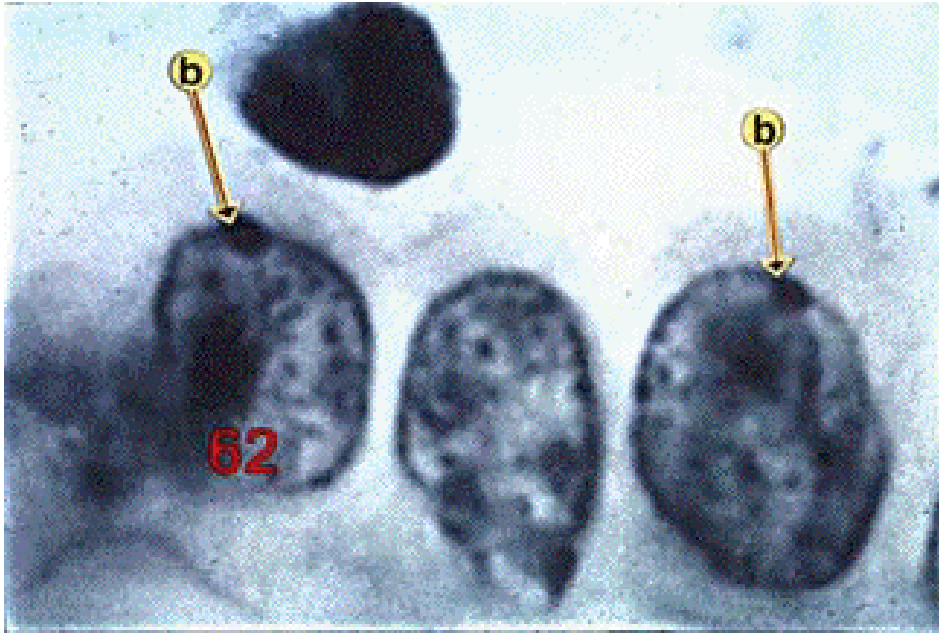
Levels of DNA packing



A polytene chromosome from a *Drosophila* salivary gland



Barr bodies - highly methylated mammalian X-chromosomes



Details of transcription in Eukaryotes

Fig. 17.7

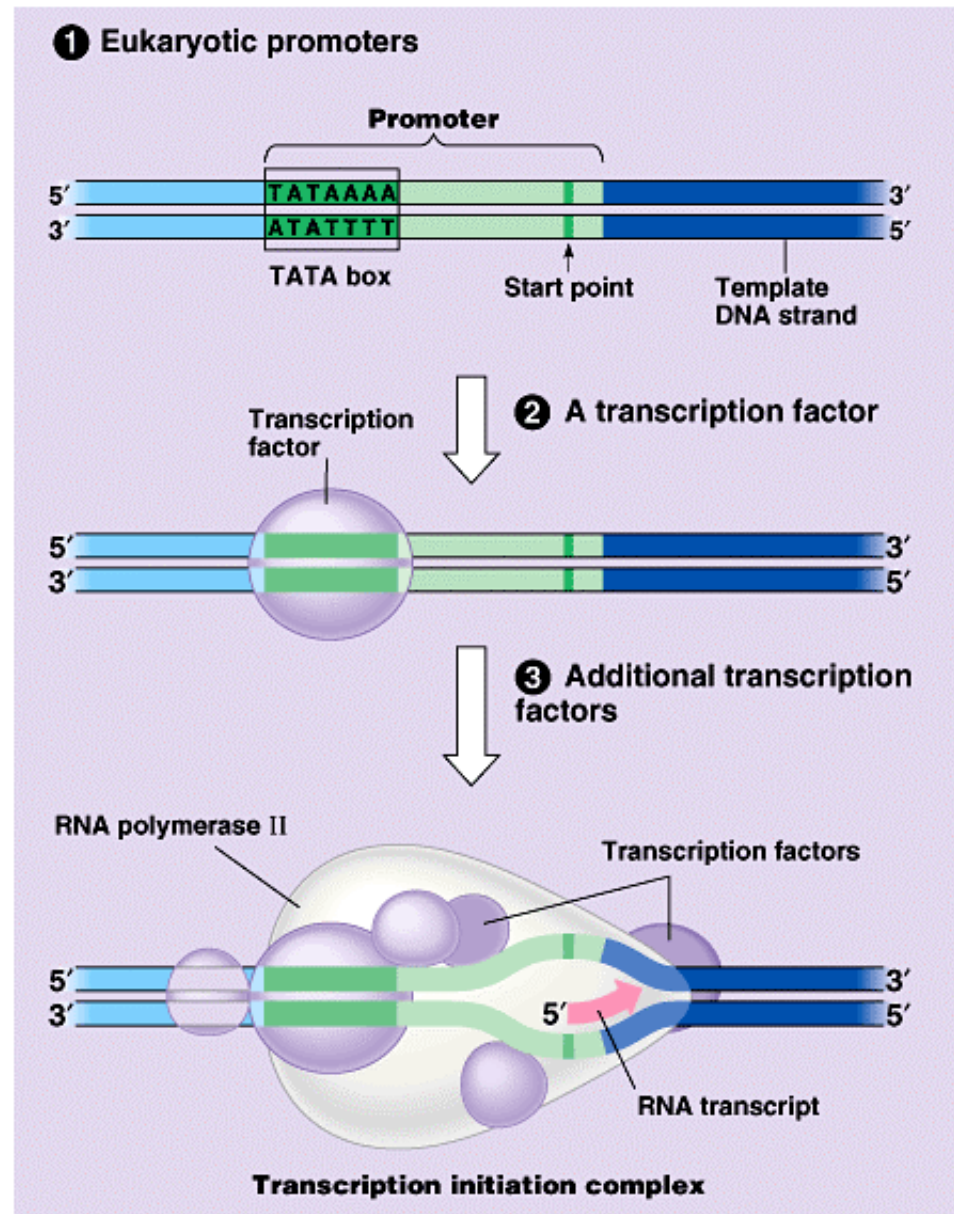


Fig. 19.8

Control elements - DNA sequences that react with proteins to facilitate the binding of RNA polymerase

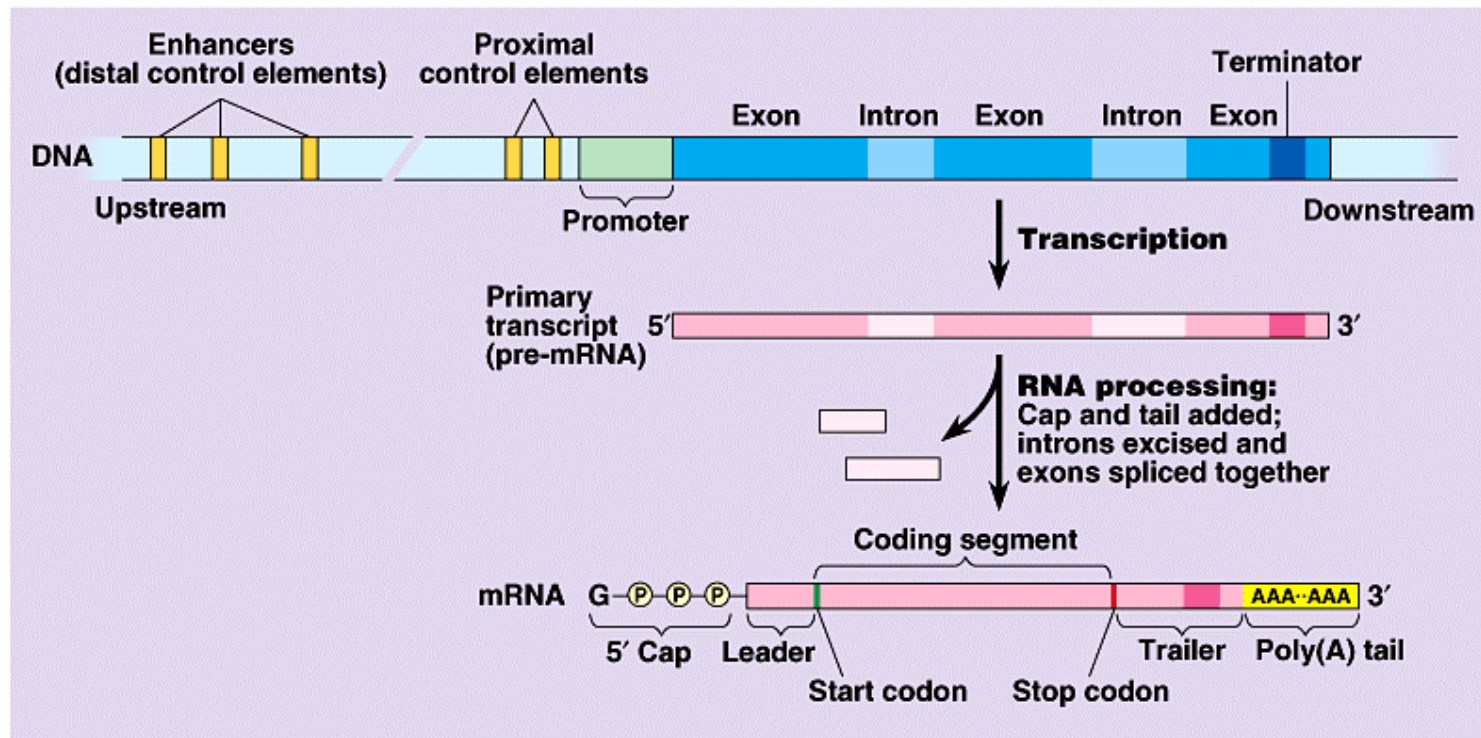
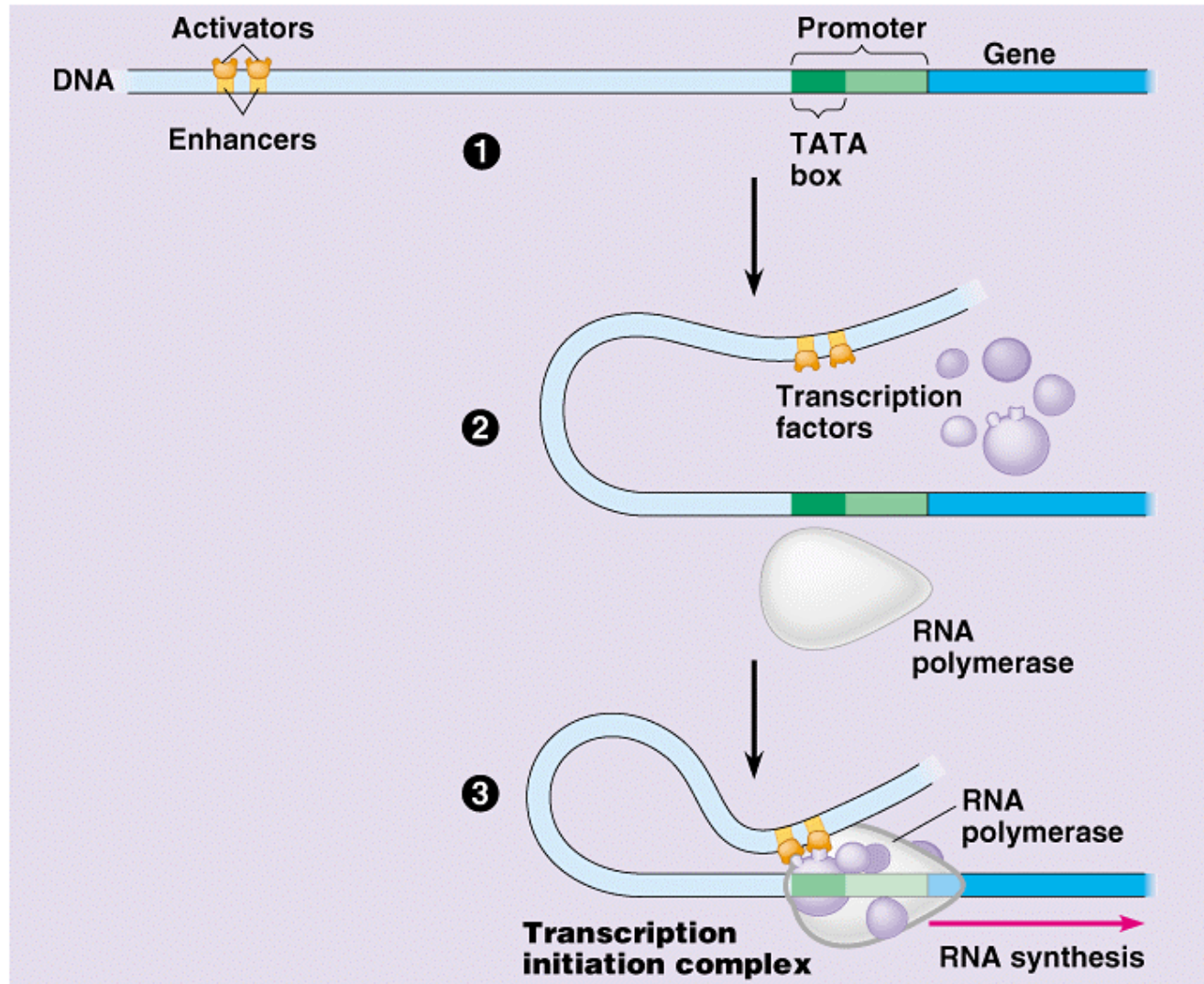


Fig. 19.9

Interaction between activator proteins, enhancer sequences, and transcription factors



Control of Gene Expression

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Fig. 19.7

Levels of control of gene expression in Eukaryotes

