

Early Genetics

- biochemical group first thought to contain genetic information = proteins

Griffith bacterium experiment~

smooth strain (S)	outer capsule; pathogenic
rough strain (R)	NO capsule; NOT pathogenic

Conclusion~

R cells combined w/ killed S cells transformed into living S cells

Avery bacterium experiment~

- deactivated parts of dead S cells to find what transformed the cells

Conclusion~

DNA transforms the bacteria

Hershey & Chase DNA experiment~

phages reproduced in presence of DNA (not proteins)

Conclusion~

DNA is the genetic material

Chargaff nucleotide experiment~

Conclusions~

Early Genetics (cont)

base composition varies between each species (diff. % nucleotides)

of nitrogenous bases equaled (A=T G=C)

DNA Structure

- x-ray crystallography images of DNA by: Rosalind Franklin

↳ DNA is a helical shape

- construction of the double helix model by: Watson & Crick

- purines (2 rings) A & G

- pyrimidines (1 ring) T & C

- A pairs with T by... 2 H bonds

- C pairs with G by... 3 H bonds

- base pairs present in 1 helix turn = 10

antiparallel: subunits run in opposite directions

DNA Replication Experiment

- experiment done by: Meselson & Stahl

Prediction

replication style	# bands 1st rep.	# bands 2nd rep.
<i>conservative</i>	2	2
<i>semiconservative</i>	1	2
<i>dispersive</i>	1	1

Results

# bands 1st rep.	1
# bands 2nd rep.	2
conclusion =	semiconservative



Replication Process

origin of replication: site where the replication of DNA molecules begins

replication fork: Y-shaped region on the replicating DNA molecule

- E. coli

- ↳ 1 replication origin
- ↳ 500 nucleotides/sec

- human

- ↳ 100s-1000s of replication origins
- ↳ 50 nucleotides/sec

- 2 items required to start replication:

1. primer
2. DNA template strand

- how added nucleotides bring energy:

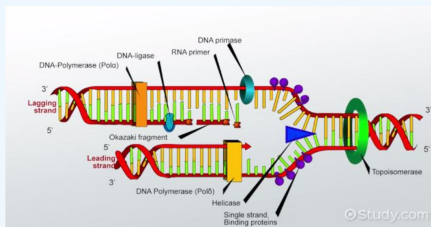
- ↳ nucleotides carried by triphosphate
- ↳ DNA polymerase catalyzes triphosphate
- ↳ 2 phosphates are released

- DNA polymerase adds to 3' end the...

(elongates from 5' to 3')

- lagging strand created from series of... Okazaki fragments

Replication



helicase: enzyme that unwinds & separates the DNA strands

topoisomerase: enzyme that breaks, swivels, & rejoins the DNA

primase: enzyme that synthesizes RNA primers

primer: a short sequence of RNA that starts Okazaki fragments

polymerase III: enzyme that adds nucleotides

polymerase I: enzyme that removes the primer and replaces the nucleotides

ligase: enzyme that forms the final bonds between the fragments and nucleotides

Errors in DNA

- as replication occurs, DNA polymerase finds & corrects any mistakes ---- *reducing the error rate*

- change in the DNA nucleotide is *permanent/mutation* when ---- *the pair is replicated*

- changes in DNA nucleotides due to...

- ↳ 1. replication errors 2. chemicals 3. x-rays 4. spontaneously

telomeres: multiple repetitions of a short nucleotide sequence at the end of a chromosome

↳ buffer zone to delay erosion of the genes as they get replicated

telomerase: enzyme that catalyzes the lengthening of telomeres (restore original length)

histone: protein responsible for the first level of packing of chromosomes

nucleosome: segment of DNA wound around a protein unit

Gene Expression Background

gene: region of DNA expressed to produce a functional product (polypeptide/RNA molecule)

transcription: synthesis of RNA from DNA template

translation: synthesis of proteins from encoded mRNA

primary transcript: initial RNA transcript from any gene (pre-mRNA)

codon: 3 nucleotide sequence that specifies a particular amino acid

- *eukaryotes*~

from nucleus to ribosome

- *prokaryotes*~

from cytoplasm to ribosome



Transcription

RNA polymerase: enzyme that controls the transcription of DNA to RNA

↳ pries DNA strands apart & joins the RNA nucleotides

↳ moves 3' to 5' (strand formed 5' to 3')

↳ attaches at the **promoter**

3 STAGES OF TRANSCRIPTION

1. Initiation

- *transcription factors:* protein that allows for polymerase to attach to DNA and transcribe

- 3 items to make up transcription initiation complex = transcription factors, RNA polymerase, & promoter

- *TATA box:* promoter that is 20-25 nucleotide from the starting point

* prokaryotes have NO transcription factors

2. Elongation

a. 10-20 nucleotides exposed at a time

b. nucleotides added to the 3' end of the RNA molecule

- difference between RNA & DNA nucleotides = different sugars

- nucleotide RNA that DNA doesn't have... *uracil*

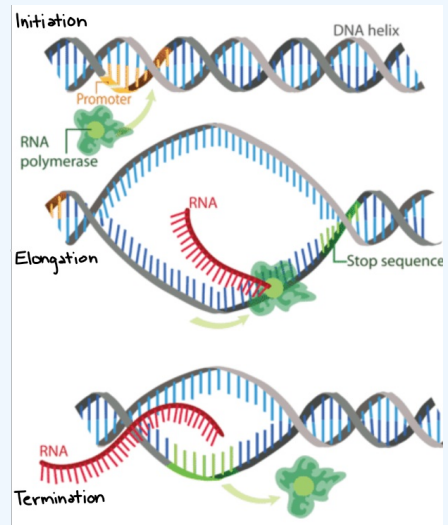
- RNA & DNA nucleotides held together by... *hydrogen bonds*

3. Termination

a. transcription of the *polyadenylation signal* adds nucleotides of AAUAAA to RNA

b. protein cuts the pre-mRNA from polymerase = end of process!

Transcription Diagram



Pre-mRNA Modification

- 5' end receives **5' cap**

- 3' end receives **poly-A tail** (enzyme adds 50-250 more A nucleotides)

↳ *facilitate export* from nucleus

↳ *protect mRNA* from hydrolytic enzymes

↳ *help ribosomes attach* to end of mRNA

RNA splicing: process of removing RNA sections from pre-mRNA

splicing:

- *introns:* noncoding sequences of pre-mRNA

- *exons:* sequences of pre-mRNA used for translation

- 3 benefits of introns:

↳ make many different polypeptides

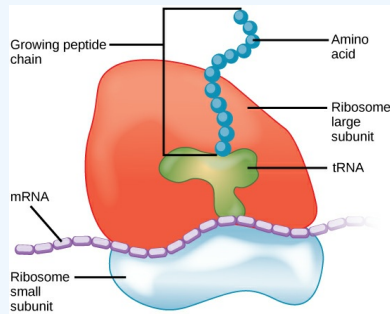
↳ discrete structural/functional regions

↳ increase exon shuffling (new protein function)

ribozymes: RNA molecule that functions as an enzyme

(create 3D structure; contain functional groups; H bond w/ DNA or RNA)

Ribosome Structure



tRNA: transfers amino acids from cytoplasm to ribosomes (& contain anticodon)

↳ **anticodon:** nucleotide triplet on tRNA molecule

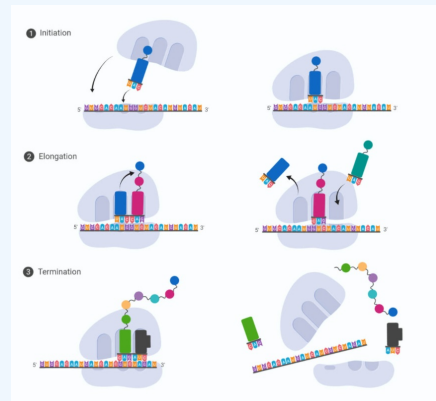
↳ **'wobble':** flexible base pairing at the 3rd codon position

- # of amino acids used= 20

- **makeup of a ribosome:**

- *large & small subunit*- made of proteins and rRNAs (eukaryotes in nucleolus & prokaryotes in cytoplasm)

Translation Diagram



polyribosomes: series of ribosomes moving over an mRNA at the same time

chaperone protein: proteins that assist polypeptides in forming 3D structures

signal peptides: sequence of amino acids at beginning of polypeptide tagging it to where it will go

Translation

3 STAGES OF TRANSLATION

1. Initiation

a. small subunit binds to mRNA & initiator tRNA

b. *translation initiator complex*= attachment of large subunit (& initiation factors)

2. Elongation

a. codon recognition- anticodon of tRNA pairs w. mRNA codon

b. peptide bond formation- removes polypeptide from tRNA by forming peptide bond

c. translocation- empty tRNA released

* ribosome moves 5' to 3'

3. Termination

a. stop codon- "release" factor accepted

b. hydrolysis of bond- freeing polypeptide

c. subunits dissociate- mRNA can be used again

Nucleotide Mutations

Nucleotide-pair sub.:

AACCA~~G~~TT
AACCA~~A~~TT

Insertion:

TACTTCAA~~A~~
TAC~~A~~TTCAA

Deletion:

TCAAACCG~~G~~
TCAA~~_~~CCGG

point mutation: change in a single nucleotide

frameshift mutation: change in nucleotide # to not be a multiple of 3

↳ may still code for same amino acid

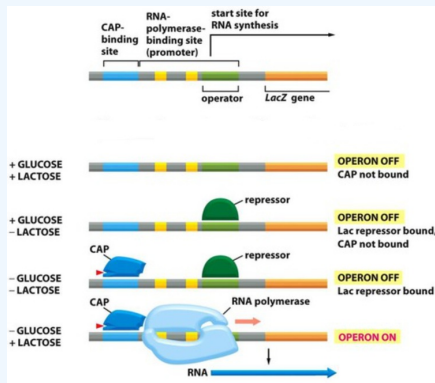
↳ may code for stop codon early

↳ may result in protein not functioning properly

Regulation of Gene Expression

- responds to changes in environmental conditions
- either adjusts **activity** of enzymes present or **production** of enzymes
- 3 things to make up an *operon*: operator; promoter; genes
- operator**: segment of DNA within promoter that controls the access of RNA polymerase to the genes
- repressor**: protein that binds to operator to block attachment of RNA polymerase
- ↳ made by activity of *regulatory gene*
- repressible operon**: transcription is inhibited by small molecule binding to regulatory protein
- inducible operon**: stimulated when small molecule binds to regulatory protein

Lac Operon



- *high lactose* = allolactose bind to repressor to change shape & no longer attach
- *low glucose* = high levels of cAMP combine with CAP

Differential Gene Expression

- differential gene expression = different cell types
- 3 processes of development:
 1. cell division
 2. cell differentiation
 3. morphogenesis
- cytoplasmic determinants**: substances in the egg that influence the course of early development
- induction**: embryonic cells influence the development of another (change in gene expression)
- homeotic genes**: genes that control pattern formation as an organism develops

Biotechnology

Gel electrophoresis

- separates DNA by size and charge
- DNA negatively charges
- ↳ smaller segments = farther to bottom

Polymerase Chain Reaction (PCR)

- create many copies of DNA segment
- ↳ DNA denatured → primers added → DNA replicated

Recombinant DNA

- DNA segment put into plasmid to be reproduced

DNA Sequencing

- establish the order of nucleotides
- ↳ labeled with dye

