

Cell cycle $G1 \rightarrow S \rightarrow G2 \rightarrow M$

Cyclin present during TRANSITIONS, specificity \rightarrow regulation + then 1) phosphorylation
 2) proteasomes and ubiquitin
 & temperature sensitive sometimes
 CDKs always there

- temperature sensitive mutations

- switches b/w active 3D conformation and inactive conformation

• checkpoints help regulate and fix mistakes

Science things

transgene organisms

CANCER

Concepts

- Contact inhibition prevents cancer (can't form colonies)
- Ames

- ~~amplifier~~ oncogenes turn on growth, inhibit cell death \rightarrow normal = proto-oncogene \rightarrow only one homologue needed for a gain of function to increase function for cancer.
- Tumor suppressors - protects cell from cancer \rightarrow recessive mutation needed for cancerous type

Ames Test - can this chemical cause mutations?

Indirect Acting Mutagen { adding liver extract to make things reactive

Direct

- (1) His⁻ bacterial cells \rightarrow plate \rightarrow plate + mutagen
- (2) if mutagen + plate has colonies b/c His⁻ \rightarrow His⁺, then it has higher mutagenic potential

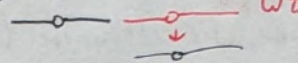
Things

- RAS } oncogenic
- MYC }

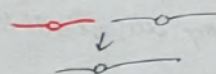
Ex - Retinoblastoma and loss of heterozygosity

$Rb^{+/-} \rightarrow Rb^{-/-}$
 $\rightarrow Rb^{+/+}$

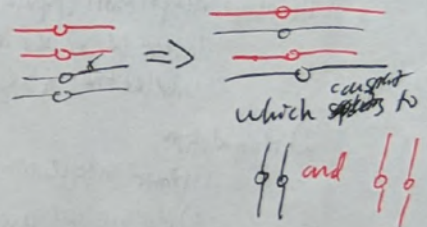
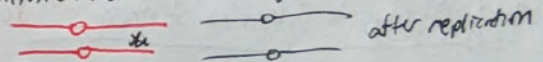
(1) De novo



(2) Chromosome Loss



(3) Mitotic recombination during repair



Mutations needed for cancer

- Multiple hit hypothesis (1 + gene mutation)

Normal \rightarrow hyperplasia \rightarrow dysplasia \rightarrow tumor \rightarrow metastasis

- old age
- radiation

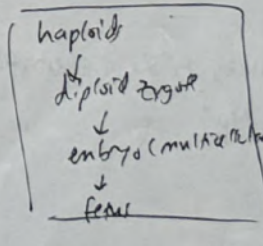
DEVELOPMENT

- Model organisms
- genes can be ubiquitous, restricted, or cell-type specific
- FACTORS impact cell type

Effector proteins differentiate cells

Regulatory genes help cells commit

- \rightarrow cell-cell signalling
- \rightarrow determinants
- \rightarrow inherited factors



ectoderm
epidermal
mesodermal

Lengthening an Epithelial sheet

- Epiboly (lengthening)

- intercalation

Organogenesis - organ - many cell types in 3D org
 - MIGRATION - SEQUENTIAL INDUCTION
 - COINDUCTION

Tissue engineering

- platform
- cell types
- signaling molecules
- 3D information

Morphogenesis - acquire shape

stick together through junctions

- tight (closed)
- gap
- adhesion junction

sort

cadherins (some cells attach to each other)

mut

- Fact'n poly

Mesenchyme - single units/cells that are loosely connected, no junctions

Epithelium

- cell sheets, yes junctions

E → M (metastasis)
MET

Stem Cells

→ measure lifetime, migration, differentiation by tracking shape/size of a certain cell

Pulse/chase assay - measures cell turnover ($t_{1/2}$) (implies stem cells by cell replacement)

- 1) pulse w/ BrdU or other label, chase w/ regular thymidine/base
- 2) Measure how many cells are still there.

Assays to determine "stemness" → measure SC potency

- They all solve "Is this a stem cell?"
- 1) Embryo incorporation (popular)
 - Inject labeled SCs into blastocyst + pseudopregnant mouse
 - labeled fates of SC
 - 2) repopulation
 - 1) remove endogenous cells (like w/ radiation)
 - 2) replace w/ possible SC, see survival
 - 3) In vitro induction
 - 1) Pump in diff inducers to see differentiated cell types
 - 2) see the diff cell types

FACS

- use labels and antibodies to ISOLATE stem cells
- use labeled antibodies

Making Stem Cells

- Embryonic SC
- iPS

Potency

Totipotent - everything
Multipotent/Pluripotent

Bipotent

↓
3 germ layers potential
Medium Potential

Cloning - SCNT

Organoids - scaffolds,
- hydrogels
- matrices

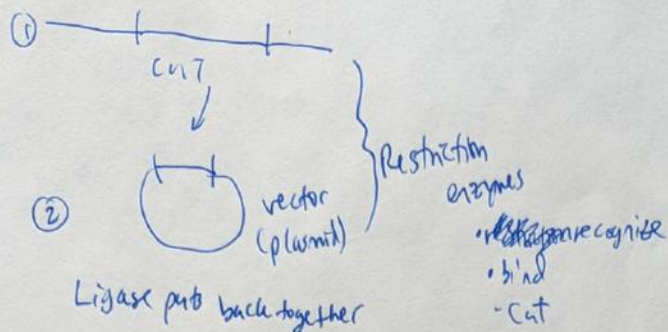
3D!

Stem Cells

- divide asym
- low acetylation Lfxn (over diff, increases acetylation)
- low methylation Pfxn

Stem cell niche - micro environment "non stem cells" + SCs

Recombinant DNA - Cloning



Ex $5' \text{G/AATTC} 3'$
 $3' \text{CTTAA/G} 5'$

↓
5' overhang,
staggered cut

- ③ Vector can replicate in host
- * Selection - some vector enters bacteria
 - * Screening

SNPs → substitution of one base pair

- noncoding SNPs change gene exp
- certain SNPs are related to certain alleles

Testing how well you pasted: restriction mapping



CRISPR

- 1) Target gene → delete/edit it
- 2) guide RNA complementary
- 3) Cas-9 makes cuts

guide RNA design

- Find AUG
- Find N6G → on target
- Box 20 bp before N6G

- gRNA is complementary to template (so identical to target)

Fusion proteins

* Don't mess up protein 2's reading frame

* include protein 1's AUG

* include stop codon

* to connect/cut → RE + ligase

* usually protein 2 is a GFP or label for localization or something

DNA Sequencing

- ① put some dNTPs in solution with replicated DNA strand

↳ probability of terminating

↳ separate DNA pieces synthesized by GE

↳ read 5' → 3'

PCR - amplifies sample of DNA - dsDNA + primers

- ① Denature DNA @ High temp
- ② Anneal DNA primers @ Low temp
- ③ Tag DNA pol @ med temp

Libraries - Genomic
↓
all DNA

cDNA
reverse transcribed mRNA
ie all coding sequences (identify protein exp)

ENZYME activity impacted by:

- active site regulation

activators increase enzyme activity
- reversible

inhibitors (dec enzyme activity)

competitive

mostly
rev.

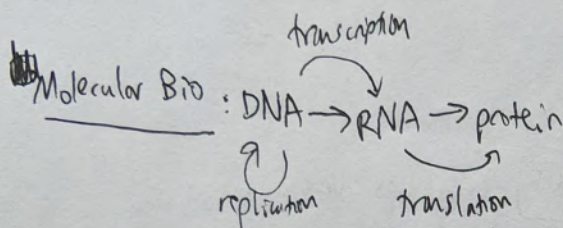
binds to
active site

allosteric/
noncompetitive

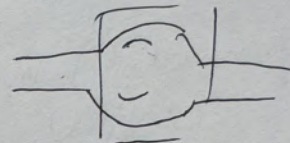
binds elsewhere

sometimes
(irreversible)
covalently binds
to enzyme

Uncompetitive inhibitors bind to ES, prevent $\rightarrow P$



Replication:

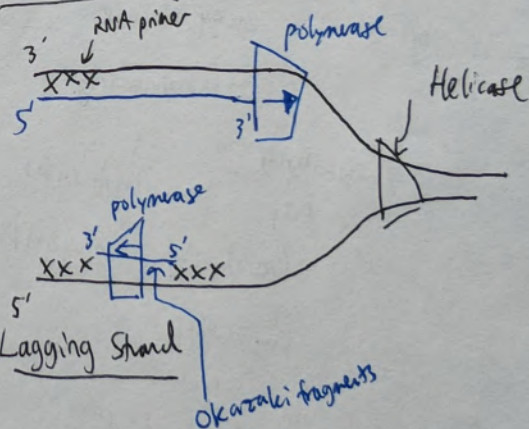


ori, tends to T-A rich

4 Rules of DNA:

- 1) Complementary base pairing
- 2) Antiparallel strands
- 3) Nucleotides added to OH 3'
- 4) Double stranded DNA can denature & reanneal

Leading strand



DNA repair

- proofreading 3' \rightarrow 5' exonuclease DNA pol (during rxn)
- Mismatch repair - single nucleotide error - identifies by methylation 5' Cytosine (diff in methylation)
- Nucleotide excision repair - thymine covalent bond errors

telomerase
repairs shortening
of chromosomal ends

INGREDIENTS FOR REPLICATION:

- nucleotides
- polynase - elongate from primers, seals gaps from primers
- RNA primase - attaches primers
- RNase - removes primers
- clamp - keeps polynase on template
- SSBP - keep DNA single stranded
- Helicase - breaks H bonds, unwinds DNA
- ligase - binds on lagging strand

BIOCHEM

1. Chemical bonds/structures

CHNOPS

Bonds: (by strength)

- 1) covalent (within molecule)
 - 2) ionic (diff charges)
 - 3) H bonds (IMF) (partial δ^- and partial δ^+ on H)
 - 4) hydrophobic - avoids water (nonpolar)
 - 5) VDW - doesn't happen as much
- } functional groups regulate 3D struct of molecule, interactions with other molecules

2. Macromolecules

Carbohydrates (monosaccharides) - hydrophilic
 $C:H:O$

LIPIDS

- energy, recognition markers
- hydrophobic
- energy storage
- cell signaling

Steroids - rings

phospholipids

PO_4

Glycerol

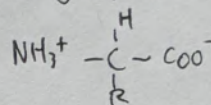
Fatty acids

Triglycerides

glycerol + 3 fatty acids

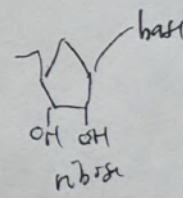
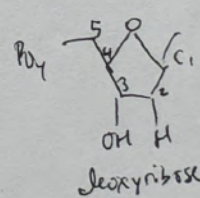
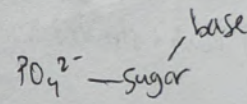
PROTEINS

- R groups determine polarity



NUCLEIC ACIDS

- nucleotides
- hereditary info



3. Energy & Enzymes

Exergonic: $\Delta G < 0$

Endergonic: $\Delta G > 0$

$$\Delta G = \Delta H - T\Delta S$$

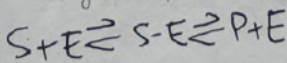
generally endergonic
Anabolic: build new molecules

Catabolic: breakdown of more complex organic molecules

Sometimes gives off energy

generally exergonic

ENZYMES - rxn specific



* reversible

* lowers E_a

* makes rxns faster

* behavior impacted by

- pH, temp

- coenzymes

Impacts substrate by:

- promoting transition state
- changing structure
- changing orientation
- creating conformational change