

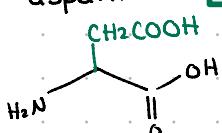
MENORIZE

BIO/BIOCHEM

→ AMINO ACIDS

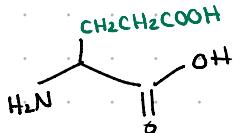
acidic - very charged

aspartic acid Asp/D



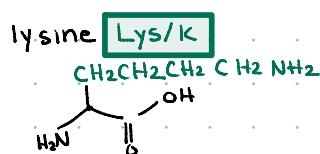
basic - very charged

glutamic acid Glu/E

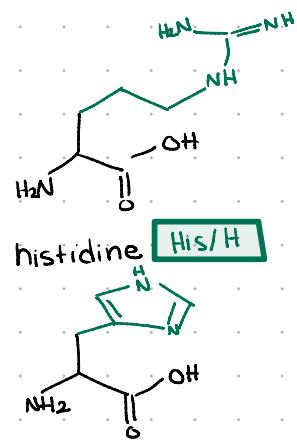


highest isoelectronic point
basic - very charged

lysine Lys/K

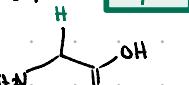


arginine Arg/R

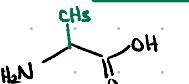


hydrophobic → aliphatic → aromatic

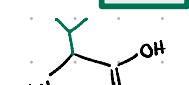
glycine Gly/G



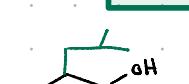
alanine Ala/A



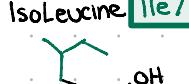
valine Val/V



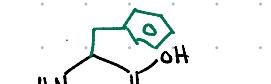
Leucine Leu/L



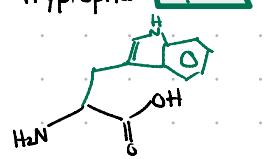
Isoleucine Ile/I



phenylalanine Phe/F

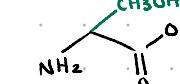


tryptophan Trp/W

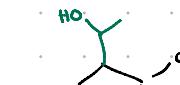


polar amino acids

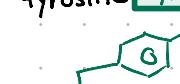
serine Ser/S



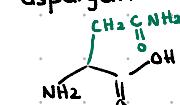
threonine Thr/T



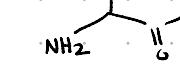
tyrosine Tyr/Y



asparagine Asn/N

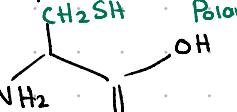


glutamine Gln/Q

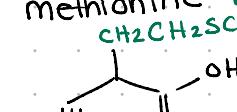


sulfur containing

cysteine Cys/C



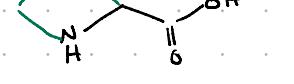
methionine Met/M



unique

proline Pro/P

Non polar

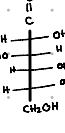
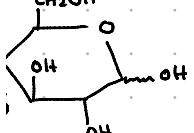


→ CARBOHYDRATES

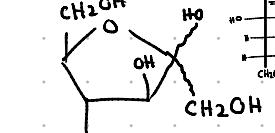
monosaccharides

D GC

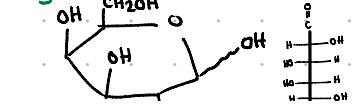
glucose



fructose

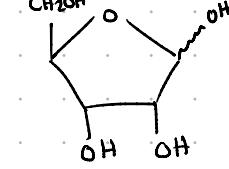


galactose

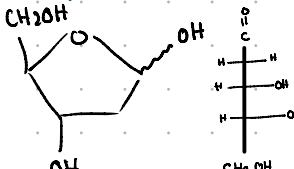


② 5C

ribose



deoxyribose



disaccharides

① maltose = glucose + glucose

reducing α 1-4

② sucrose = glucose + fructose

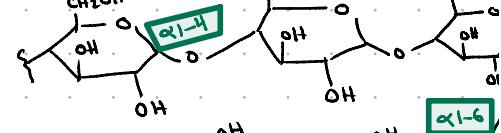
non reducing 1-1

③ lactose = glucose + galactose

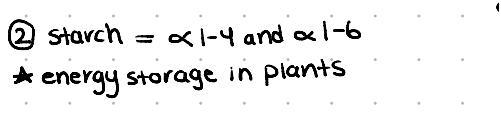
reducing β 1-4

polysaccharides

① glycogen = α 1-4 and α 1-6
branching
* energy storage in animals



② starch = α 1-4 and α 1-6
* energy storage in plants



③ cellulose = β bonds
* cell walls in plants

★ better at storing energy than carbs

① **triglyceride** = 3 fatty acid + glycerol
ester bond $R-O-C-OH$ backbone (3C sugar)

★ energy storage in adipose tissue

★ fat transport in bloodstream

② **phospholipid** = 2 fatty acids + phosphate head + glycerol backbone (3C sugar)

★ amphipathic

★ form cell membranes

③ **terpene** = multiple isoprenes $(C_5H_{10})_n$

★ 2 isoprene \rightarrow terpene

★ 3 terpene \rightarrow squalene

★ precursor for cholesterol

★ can form Vitamin A

VITAMINS

p395 - come back to this

TERMODYNAMICS

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

-ve \rightarrow spont exergonic

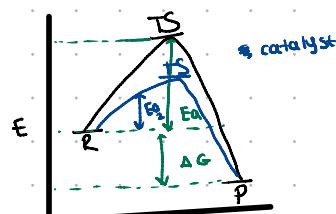
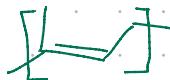
+ve \rightarrow non spont endothermic

0 \rightarrow eqm

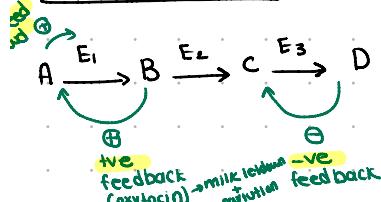
NOT changed by enzymes

④ entropy \uparrow in universe

coupling = spont rxn + unspont rxn \rightarrow drive unspont rxn



TYPES OF LOOPS



INHIBITION

$$V_{max} \rightarrow [E]$$

$$K_m \rightarrow [S]$$

comp

ACTIVE

$$V_{max} -$$

$$K_{max} \uparrow$$

non

ALLOSTERIC

$$V_{max} \downarrow$$

$$K_{max} -$$

un

E-S

$$V_{max} \downarrow$$

$$K_{max} \downarrow$$

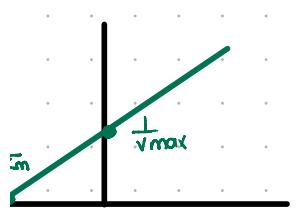
mixed

E-S OR E unoccupied

$$V_{max} \downarrow$$

$$K_{max} \text{ depends}$$

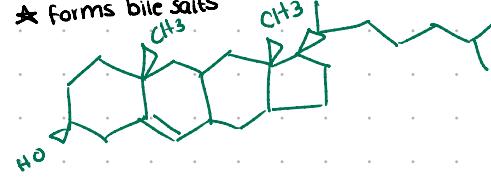
Lineweaver-Burk



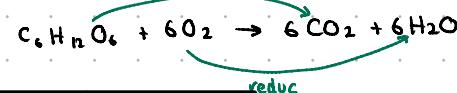
closer to origin
 \downarrow
denominator \uparrow

3-6C rings + 1-5C ring
precursor to estrogen and testosterone
regulates membrane fluidity + forms lipid rafts

forms bile salts

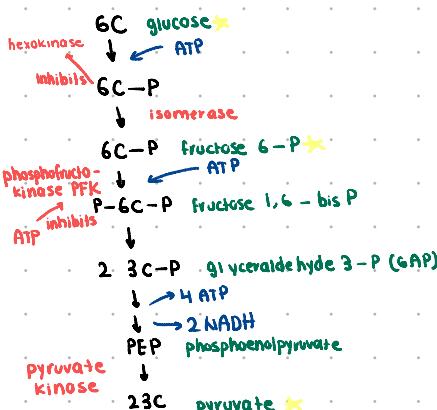


CELLULAR RESPIRATION

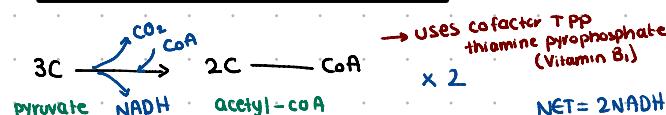


process	location	O ₂
glycolysis	cytosol	No
PDC/Krebs	matrix of mitochondria	Yes, indirectly
ETC/oxidative phosphorylation	inner membrane of mitochondria	Yes, directly

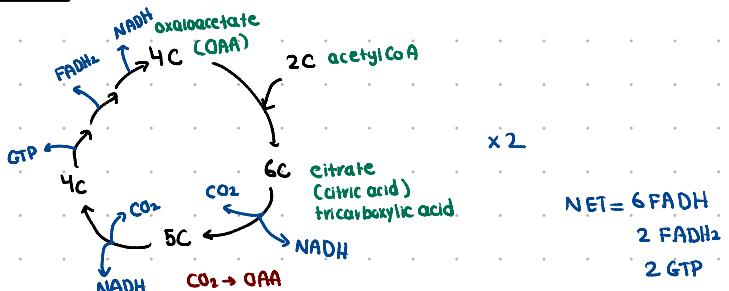
GLYCOLYSIS



Pyruvate Dehydrogenase Complex (PDC)



KREBS



ETC/Oxidative Phosphorylation

① make ATP

② oxidize e-carriers

matrix $NADH \rightarrow NAD^+$

inner $FADH_2 \rightarrow FAD$

inner e^- space $CoQ \rightarrow CoQ$

outer $NADH \rightarrow NAD^+$

outer $FADH_2 \rightarrow FAD$

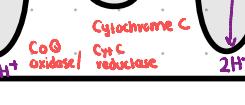
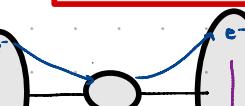
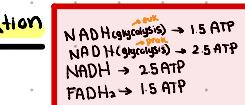
outer e^- $CoQ \rightarrow CoQ$

outer $Cyt C \rightarrow Cyt C$

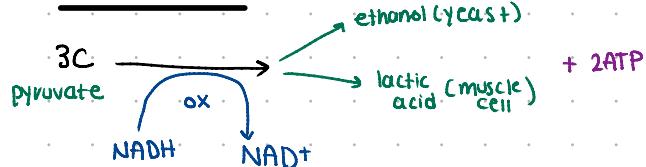
outer $2H^+ \rightarrow 2H^+$

outer $Cyt C \rightarrow Cyt C$

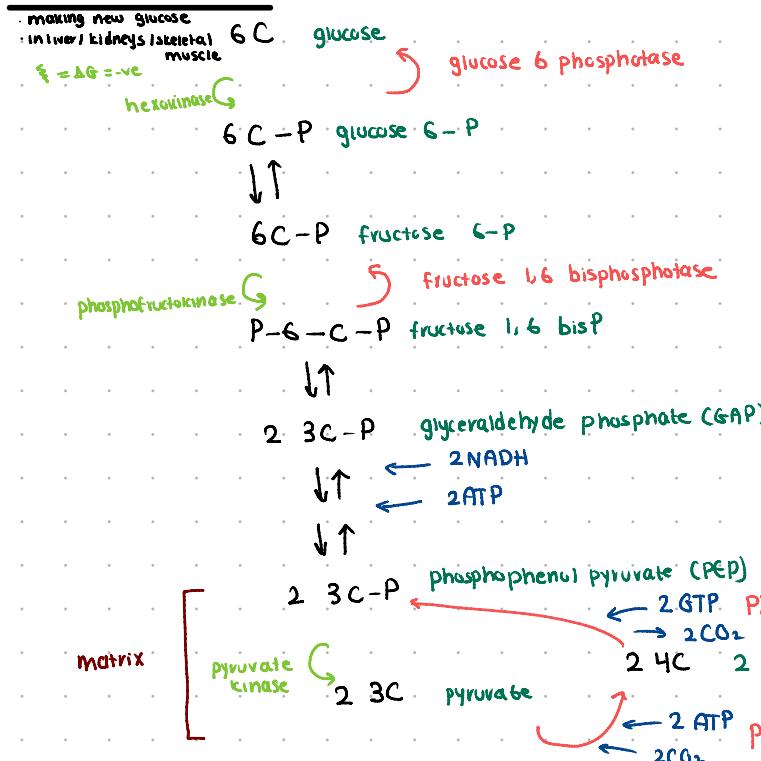
outer $ATP \text{ Synth}$



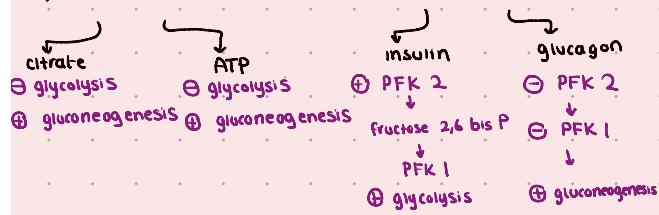
→ FERMENTATION



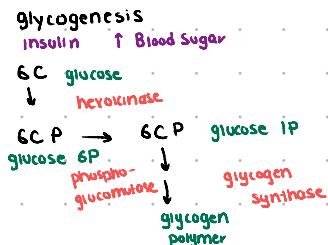
→ GLUCONEOGENESIS



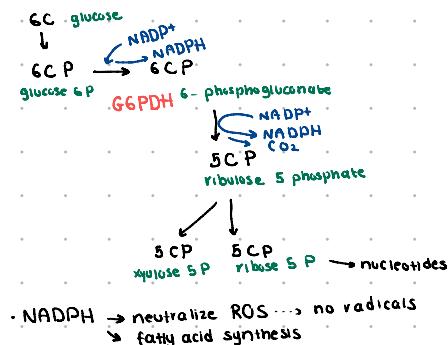
reciprocal regulation



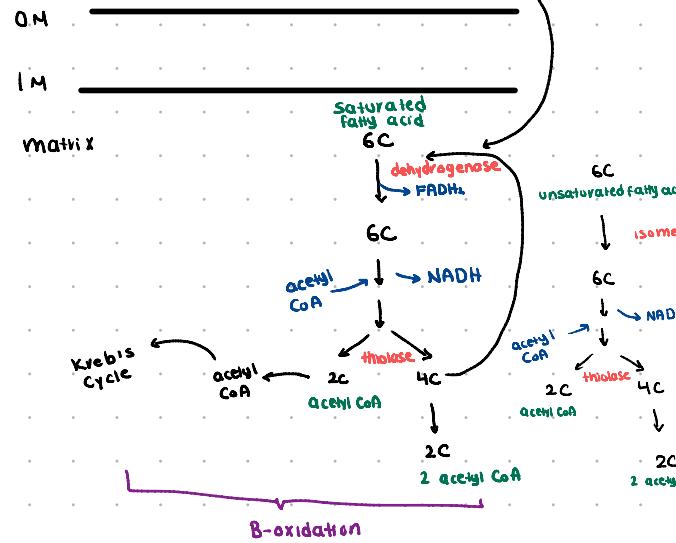
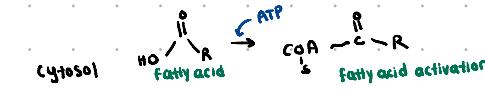
→ GLYCOGEN METABOLISM



→ Pentose Phosphate Pathway



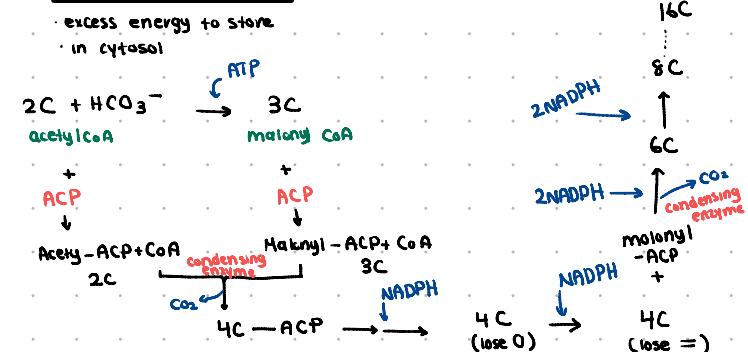
→ FATTY ACID OXIDATION



$$\# \text{ of cycles} = \frac{\# \text{ carbons}}{2} - 1$$

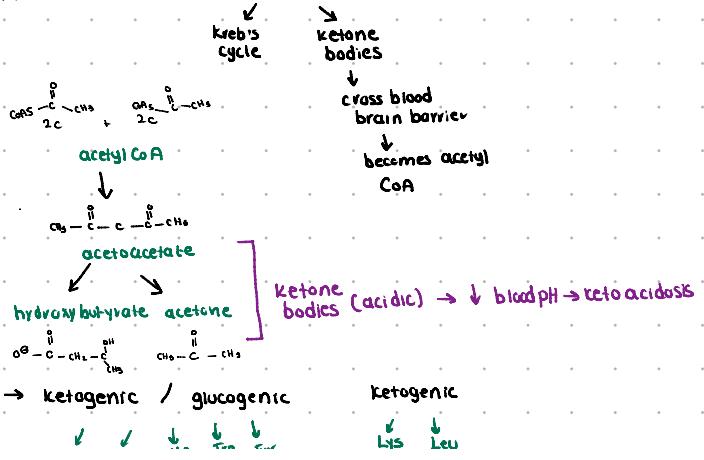
odd # = 2 acetyl CoA + 2 NADH CoA

→ FATTY ACID SYNTHESIS

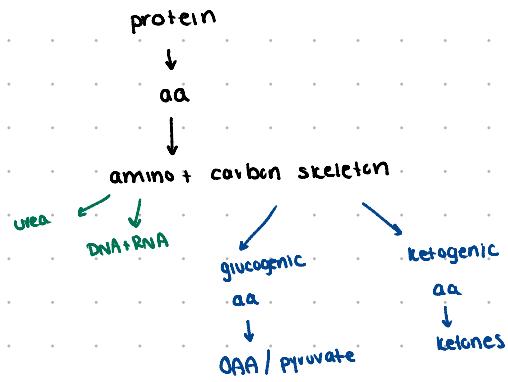


→ KETOGENESIS

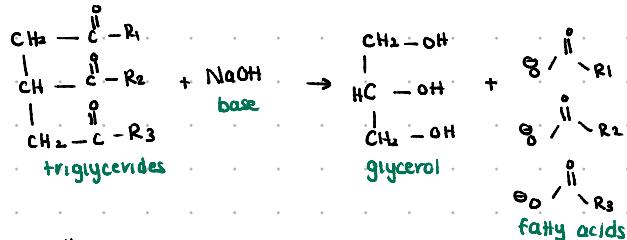
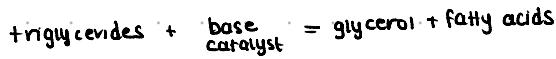
long term starvation
 blood glucose → ↓
 fatty acid oxidation occurs → ↑ acetyl CoA



→ PROTEIN METABOLISM



→ SPONIFICATION



→ "OTHER" LIPIDS

i) sphingolipids = sphingosine + 2 fatty acids + phosphate.

* sphingomyelin = myelin sheath of neurons

j) waxes = long chain fats esterified to hydrophobic long chain alcohols

k) fat soluble vitamins =

A → vision
D → bone
E → antioxidants
K → clotting

) prostaglandins = 20c fatty acids w/ different roles in tissues depending on receptor they are bound to (5 membered ring)

→ CLASSES OF ENZYMES

hydrolase	hydrolyzes chemical bonds
isomerase	rearranges bonds to form an isomer
ligase	forms a chemical bond
lyase	breaks chemical bonds w/out hydrolysis
kinase	transfers P group from a high energy carrier
oxidoreductase	runs redox rxns
polymerase	polymerization
phosphatase	removes P group
phosphotransferase	transfers P from inorganic P
protease	hydrolyzes peptide bonds