

INTRODUCTORY

Explaining Bio - how & why

- Time dimension
 - Proximate: immediate cause of an effect (how, mechanism)
 - Ultimate: higher level reason for an effect (why, historical or functional)
- * Biological organization dimension
molecules, proteins → genes, → organisms → communities → ecosystems, biomes
- * Spatial dimension
local to global

B. Organism - alive?

- Energy: acquire and use energy
- Cells: membrane bound units that regulate structure and function
- Genes/information: to pass on info to next gen, process and respond to stimuli.
- Replication: sexual and asexual

* Evolution: organisms are products of evolution and populations of organisms themselves continuously evolve

C. Diversity

- Genetic diversity: variation in genes → adapt and evolve
- Taxonomic diversity: number, abundance, distribution of species (taxa)
- Ecosystem diversity: variety of habitats and communities of different species that interact in interdependent relationships

D. Ecology vs Evolution

- Ecology: hows, whys organisms interact, and their physical environment
- Evolution: change in allele (genetic variant) frequency in a population over time

02 Evolution

change in allele (gene variant) frequency over time.

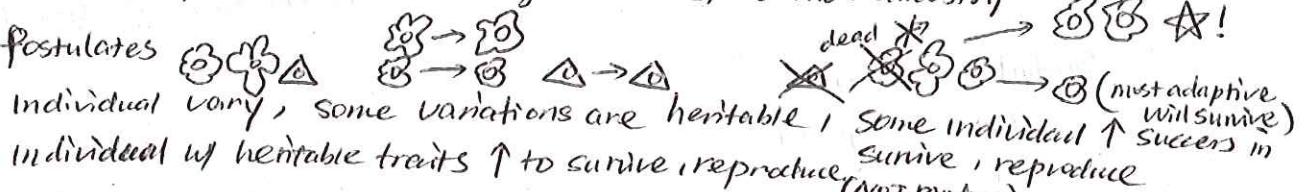
- A. Mechanisms of change:
 - Selection (fitness advantage to some alleles)
 - Randomness (drift)
 - Movement (gene flow)
 - Mutation (direct change to genes)

B. Darwin's Revolutionary Theory

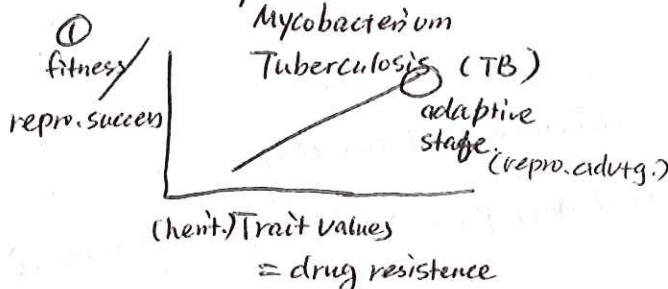
1. Natural Selection → slight variation of trait will be preserved, if useful

2. Descent w/ modification: change overtime, common ancestry

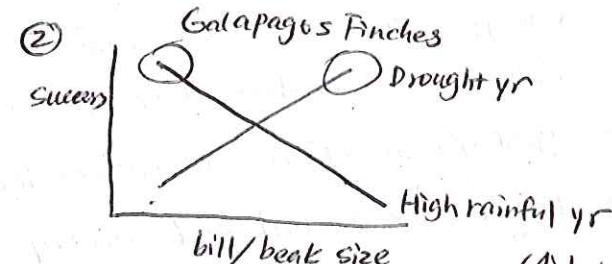
four postulates



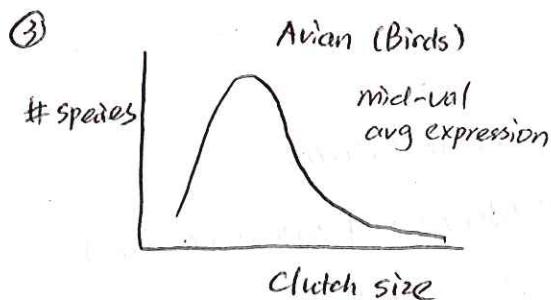
case study



Some TB mutated → drug → ↑ TB (mutated)
Individual cells did not evolve!



Drought yr → tough fruits → ↑ beaksize ↑ to eat large seed
High rainfall → ↓ beak size → ↑ small seeds
(↓ body size)
↑ pointed beak



- Natural Selection does NOT change individuals
- Acclimatization ≠ Adaptation

Theory: well-substantiated explanation of some aspect of natural world based on body of facts, repeated / confirmed thru exp. and observation. used to make predictions abt natural events not been observed

Hypothesis: a proposed explanation for a natural phenomenon (prediction)

03 Behavior Ecology

Behavioral → population → community → global

- Some complex behavior can occur without complex cognition or decision making
- = some behaviors involve cognition and are learned
- Behavior is a heritable trait that can respond to selection pressure

Tinbergen's 4 Questions

proximate (immediate) (1) Causation / mechanism: genes, physiological pathways to cause trait to occur (biological environmental)
 (2) development + how this trait/behavior develops over the lifetime of the individual (developmental mechanism or processes)

ultimate (long term) (3) Function: selected advantage related to fitness increase + function

(4) Evolution: Evolution history (behavior in a shared common ancestor), phylogenetic interpretation, with other species

CASE STUDY

1. Rovers vs Sitters in fruit fly

- ① A gene's associated w/ rover, sitter difference to cause difference in foraging traits.
- ② Innate behavior depending on food supplies
- ③ Rovers: ↑ population density, ↓ food supply, ↑ find unused energy cued to more forage
- ④ Sitters: ↓ inheritance from ancestors

2. Male Zebra Finch Song

- ① hormones promote sexual behavior, cued by day length exchange
- ② Air passing syrinx produces song
- ③ song learned from tutor (father), requires acoustic & visual information
- ④ ↑ attractiveness to females → individual fitness

3. Sexual Activity in Lizards

- ① hormones → male bobs up & down, ↑ sexual organs in female
- ② change in day length, bobbing develops (visual cue) → trigger female readiness
- ③ to synchronize the sexual readiness → avoid predators
- ④ show good male quality → sexual select. (mate w/ good traits)
- ⑤ Inherited from ancestor, the push-up ↑ during select.

4. Sexual Select. in Australia Redback Spider

- long copulation, ↑ eggs, ↑ nutrients for offspring.
- ↓ accept 2nd suitors to mate

{ Reciprocal Altruism (exchange of good act) Kin Selection. Apparent Altruism

Br > C
Benefit / cost coefficient

$$r = 0.5 \times 0.5 + 0.5 \times 0.5 = 0.5$$

Full-Siblings

Actor Recip.

0.5 → 0.5 → 0.5 → 0.5

Half-Siblings

Actor Recip.

0.5 → 0.5 → 0.5 → 0.5

$$r = \frac{1}{2} \times \frac{1}{2} = \frac{1}{4}$$

Cousins

Actor Recip.

0.5 → 0.5 → 0.5 → 0.5

$$r = \frac{1}{2} \times \frac{1}{2} \times \frac{1}{2} = \frac{1}{8}$$

direct fitness: care for own offspring → ↑ offspring → ↑ fitness of parents

indirect fitness: others (relatives) → ↑

(Haplo-Diploid male) (female)

Mom Dad D
0.5 → 0.5 → 1
Sister Sister (D)
0.5 → 0.5 → 1
Sis-Sis.
R = 0.5

Mom D
0.5 → 1
Sist. Sist.
R = 0.5

Mom-Daughter
0.5 → 1
D R = 0.25

Father Son Daughter
R = 0.5

Mom D
0.5 → 1
Sist. Sist.
R = 0.5

D R = 0.25

3

04 ANOVA

we test H_0 instead of H_1

H_0 - Null hypothesis = no effect of treatments

H_1 - Alt. hypothesis = not explained by chance, notable difference between groups

Variance. $s^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \bar{x})^2$ sum of squares

(N) degree of freedom = SAMPLE SIZE - 1

Sum of squares
(SS)

$$\text{TOTAL} = \sum (x_i - \text{total mean})^2$$

$$\text{WITHIN GROUP} = \sum (x_i - \text{treatment mean})^2$$

$$\text{BETWEEN GROUP} = \text{TOTAL} - \text{WITHIN}$$

df

sample (total) - 1

Mean Square = $\frac{s^2}{MS}$

sample - # of treatment
1 (if 2 groups).
number of treatment - 1

$$F = \frac{MS_{\text{between}}}{MS_{\text{within}}}$$

$F > F_{\text{critical}}$ (1) \Rightarrow reject H_0
 $F < F_{\text{critical}}$ \Rightarrow accept H_0

P = possibility to fit H_0
(reject or accept H_0)

$F \uparrow P \downarrow$
 $F \downarrow P \uparrow$

(chance to incorrectly reject H_0)

example.

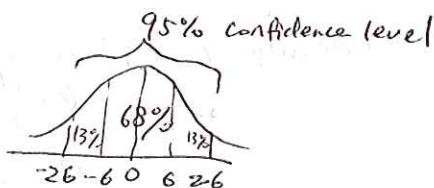
SS

$$\text{TOTAL} = \sum (x_i - \text{total mean})^2$$

$$\text{Within} = \sum_{(\text{choice})} (x - \text{mean walking})^2 + \sum (x - \text{mean running})^2$$

$$\text{between} = \text{Total} - \text{Within}$$

$$\text{Standard deviation } (s) = \sqrt{\frac{1}{N} \sum_{i=1}^N (x_i - \bar{x})^2}$$



$$\text{Standard Error of Mean (SEM)} = \frac{s}{\sqrt{N}}$$

$2 \text{SEM} = 95\% \text{ confidence level}$

population: a group of individuals of the same species that live in the same area at the same time

05. POPULATION ECOLOGY

Behavioral → Population → Community → Global

3 properties of populations

1. Size
2. Distribution
3. Density

Life Table in Conservation.

Biology $\Delta N = N_2 - N_1 = \text{Birth} - \text{Death} + \text{Immigration} - \text{Emigration}$

sea turtle ↓ adult mortality + fec.

Protect eggs → protect adult female

1. Population Size

- Assessing Birth and Death

Life Tables

TABLE 5.2 Life Table for Lacerta vivipara Females in the Netherlands

Age Class	Age	# Offspring	Survivorship	Average Age	Average Fecundity
0	0	500	1.00	0.65	0.65
1	1	474	0.92	0.64	0.62
2	2	308	0.80	0.63	0.45
3	3	158	0.68	0.62	0.38
4	4	67	0.57	0.61	0.24
5	5	12	0.29	0.60	0.21
6	6	2	0.00	0.59	0.02
					R ₀ = 1.00

$$\text{Survivorship} = \frac{\# \text{ of offspring survived}}{\# \text{ of initial population}}$$

Fecundity = # of offspring produced by a female

$$\text{Net reproductive rate } R_0 = \sum (\text{survivorship} \times \text{fecundity})$$

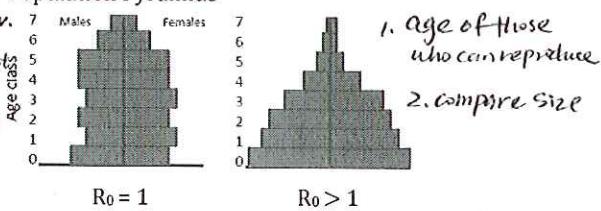
$R_0 > 1$, Population increase; $R_0 < 1$, population decrease

Application: increase survivorship of adult female

Life-history continuum

low resist. low fec.
High fec.
High surv. long life
High resist.
Fitness trade-offs
(e.g. limited resources)

Population Pyramids



Factors limiting population size

- Density-independent abiotic environs.
- Density-dependent rate competition for resources disease & parasitism predation toxic wastes social behavior

- Prediction of Population Size - Per Capita Growth Rate

$$r = \ln(R_0)/g$$

g = generation time, r = B rate - D

$$\text{Population size} = \text{Initial Population} \times (e^r)^n, n = \text{year (exponential growth)}$$

$$\frac{dN}{dt} = rN \left(\frac{K - N}{K} \right) \quad (\text{logistic growth})$$

(Density-dependent population growth)

K = carrying capacity determined by the environment (food, resources, etc.)

N = current population size,

N > K, population decrease; N < K, population increase.

Example: fecundity of sparrows declines at high population density

- Measuring Population Size

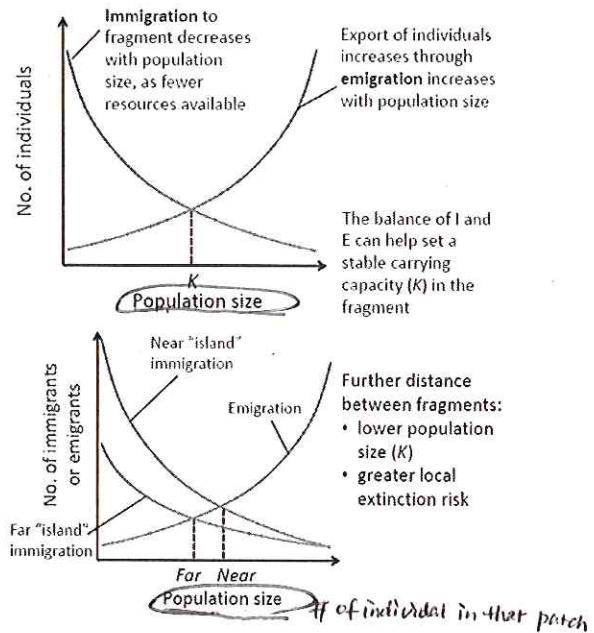
1. Transects: count everything along a predetermined survey line (within a range)

2. Quadrats: count at random locations, inside rectangular plots

$$3. \text{Capture-mark-release-recapture: } \frac{n_{\text{mrk}}}{N_{\text{total}}} = \frac{n_{\text{mrk in recap}}}{N_{\text{recap}}}$$

$$- \text{Metapopulations (E and I)} \quad \left(\frac{M}{N} = \frac{m}{n} \right)$$

Individuals from a species occupy many small patches of habitat ⇒ independent populations
⇒ meta-pop. (beyond populations)



2. Population Distribution

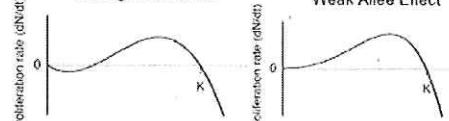
- Random: individuals don't exclude each other
- Clumped: patchy, clustered resources and enhance fitness/survival by presence of others
- Uniform/overdispersed: competitive exclusion by others (resource competition)

3. Population Density

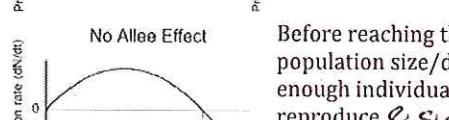
Density = population size/range

- Allee effects

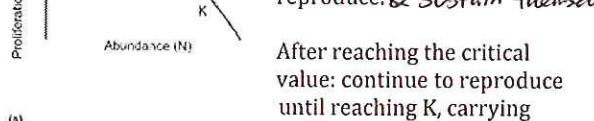
Strong Allee Effect



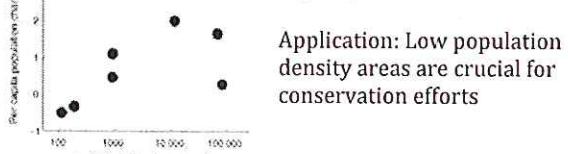
Weak Allee Effect



No Allee Effect



Abundance (N)



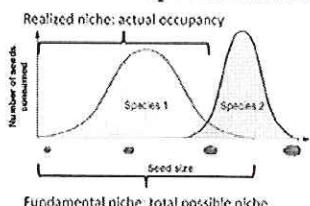
After reaching the critical value: continue to reproduce until reaching K, carrying capacity.

Application: Low population density areas are crucial for conservation efforts

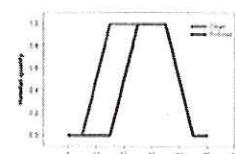
06. COMMUNITY ECOLOGY

Behavioral → Population → Community → Global
Interactions between populations/species. in a certain area

1. Niche and Species Interaction



Niche: the range of resources (shelter from predator, weather, environment conditions like temp, etc.) that a species uses and the habitat it can tolerate.



Niche overlap determines interspecies competition.
We can only measure realized niche ← caused by competition and human cases like pollution

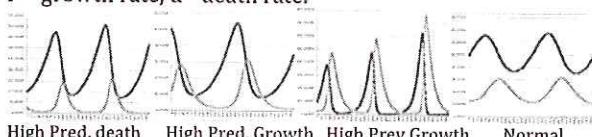
Species Interaction (Uniform pop up, distribution)
Competitive exclusion (evolved to avoid each other) leading to narrower realized niches. (-, -) e.g. fox & coyotes
2. Commensalism (+, 0) e.g. birds follow cattle
3. Mutualism (+, +) e.g. Plants & pollinators, digestive bacteria (service / produce)
Parasitism (+, -) has a cost, but not necessarily killing
4. Consumer Predation and Herbivory (+, -)

Predator (N_1) & Prey (N_2) (Predator depends on Prey)

$$\frac{dN_1}{dt} = N_1 + r_1 N_1 N_2 - d_1 N_1$$

$$\frac{dN_2}{dt} = N_2 + r_2 N_2 - r_1 N_1 N_2$$

r = growth rate, d = death rate.



2. Food Webs/Trophic Levels

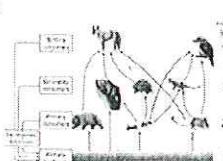
Species Classification

Keystone: low biomass (proportional mass, not individual mass → population), high impact

Dominant: high biomass, high impact

Rare: low biomass, low impact

Common: high biomass, low impact



Food Web

3. Determinants of Biodiversity

Top-down: predators → other species

Bottom-up: producers → other species

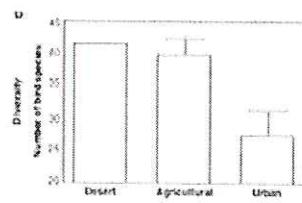
Case Study: (Dorrestein et al. 2015) The top-down effects of humans (on apex predators, mesopredators, and herbivores) is stronger than most "natural" relationships.

Island Biogeography Theory

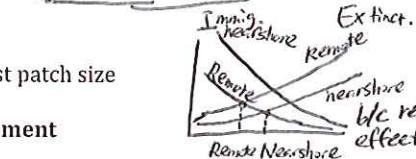
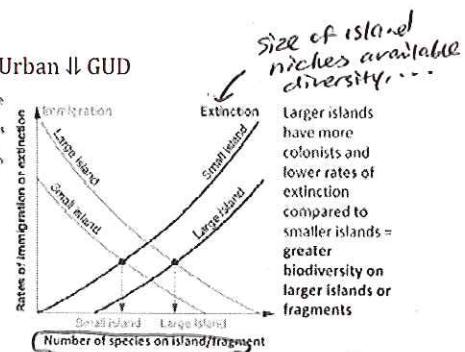
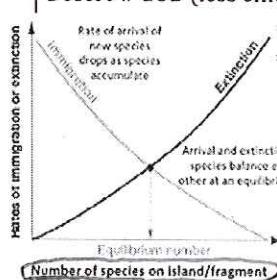
Desert, Agriculture, Urban

Urban dominant species ↑ foraging efficiency → exclude other species

Giving Up Density (GUD)



Desert ↑ GUD (less efficient); Urban ↓ GUD



Island Biogeography Theory: Extinction rate relates to forest patch size

4. Biodiversity and Measurement

Biodiversity

- Genetic Diversity: gene for evolution (↑ Variation in a heritable trait → ↑ fitness); Helps populations cope with environmental change

- Taxonomic Diversity

Richness (# of total different species in an area)

Evenness (evenness of different # of species exist in that area. Non-existent species not in consideration) "drop out"

Species composition (who the species are)
Be sure to look for changes in y-axis when examining different graphs.

- Ecosystem diversity

$$\text{Evenness} = \sum_{i=1}^S p_i \times \ln(p_i)$$

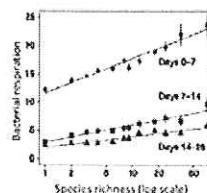
Measurement: Hotspots: species are condensed in biodiversity hotspots, but protecting the hot spots may not save endangered species b/c many mismatches between biodiversity and threatened species hotspots

5. Value of Biodiversity

A. Ecosystem Functions: water, Nutrient storage, Soil, Pollution, Climate stability

B. Biological Resources: Food, medical, wood, Breeding stocks, gene reservoirs

C. Social Benefits: Research, tourism, culture, natural disasters



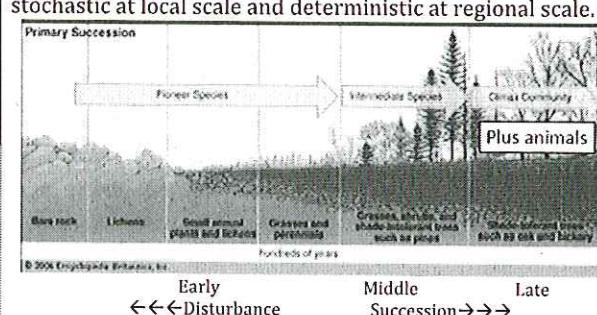
If diverse, If ecosys function,
If days, If food, If respiration.

6. Ecological Succession

Biodiversity (species composition), trophic interactions, and ecosystem structure are not static—change over time.

Can be fast (e.g. after major storm, fire, landslide,

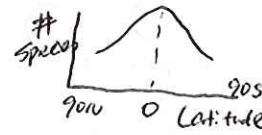
volcanic eruption, human influence) or slow (e.g. after mass extinction) Vary by scale and tend to be more stochastic at local scale and deterministic at regional scale.



07 Global Ecology

A. Seasonality

Seasonality affects, sunlight, temp. drive primary productivity affect other trophic level
 cause of seasonality: 23.5° tilt, annual rotation
 productivity of primary producers

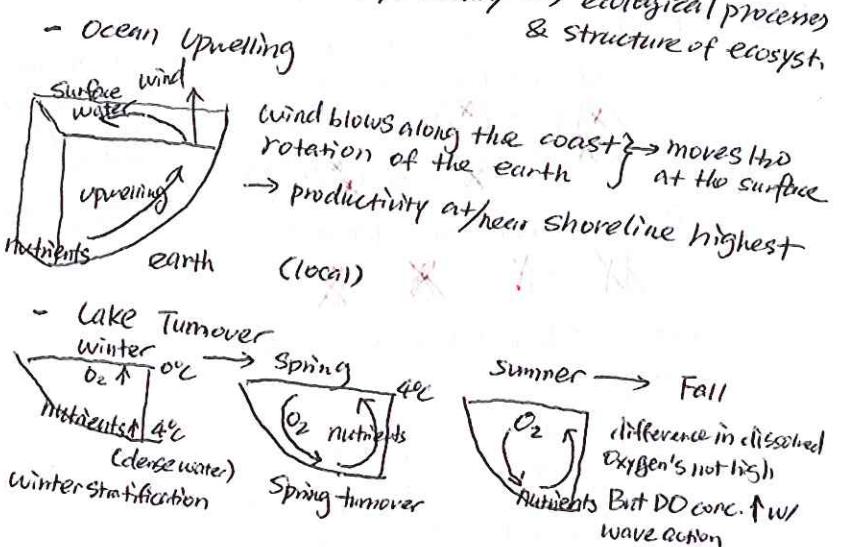
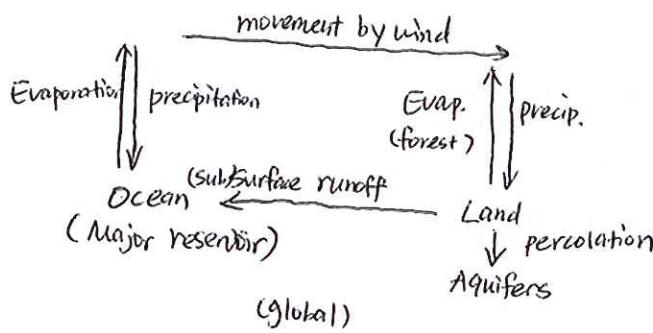


Latitudinal gradients:

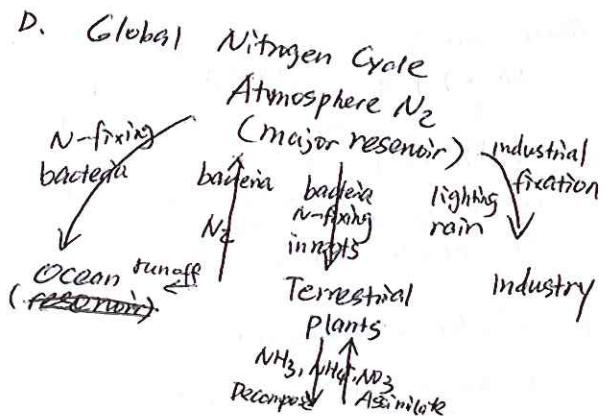
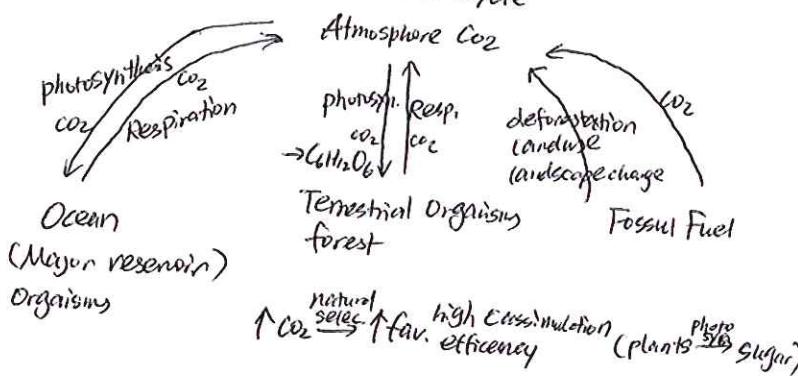
Poles are cooler than tropics
 Productivity tropics > poles
 Equator 90° - more sunlight - \uparrow Energy
 \uparrow Immune sys - \uparrow fecundity

B. Water Cycle:

Organisms need H₂O to grow & survive. H₂O nitrates more globally, locally drive ecological processes & structure of ecosystems.



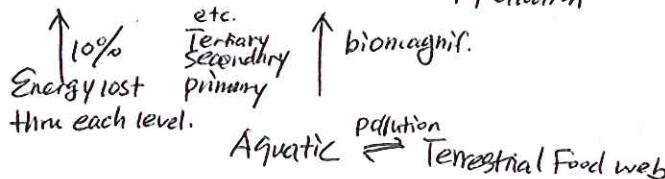
C. Nutrient Cycles = Global Carbon Cycle



E. Tropic Level & Pollution

bioaccumulation/biomagnification

Fossil fuel, metal production (excluding gold)
 Gold production, etc. \rightarrow mercury pollution

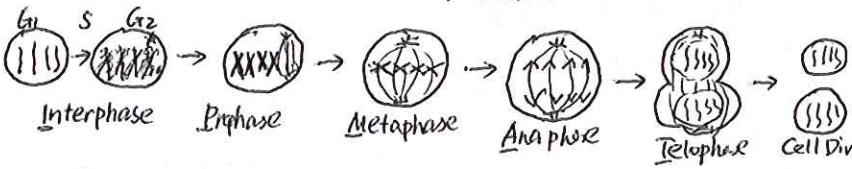


Overuse of fertilizer \rightarrow \uparrow Nitrogen \rightarrow \uparrow algae runoff \rightarrow \uparrow aerobic bacteria \rightarrow \downarrow oxygen \Rightarrow dead zone (river), fish kill, shrimp in Amer. Catch

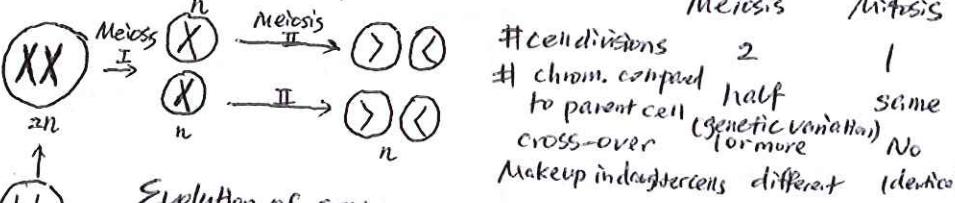
08 Microevolution

A. Mitosis, Meiosis, Phenotype genetics

1. Mitosis: repair, asexual reproduction

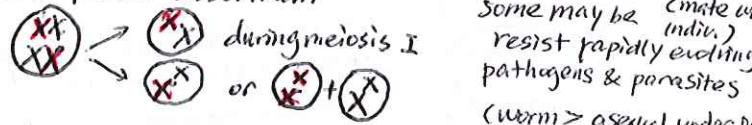


2. Meiosis: sexual reproduction, reduction division



Evolution of sex

- Benefits: greater genetic diversity
 - (a) Independent assortment



- (b) Cross-over in prophase I.

$$XX \rightarrow X \rightarrow X \rightarrow XY$$

~ Costs: "two-fold cost of males"

asex.	sex.
♀ n	♂ ♀ 2n
♂ n	♂ ♀ ♂ 4n
various indus 16n	♀ ♀ ♀ ♂ 4n

* inherit all deleterious alleles harmful
BAD!

Mendelian genetics and Punnett Square

RRYYxrryy	↓	RY Ry rY ry
RrYy x RrYy	↓	RY Ry rY ry
Dominant/Recessive genes		Homo/Heterozygous

Homo/Heterozygous

B. Evolutionary genetics

1. H-W = Hardy-Weinberg Equilibrium (without evolution)

Genotype	Frequencies	Allele Freq.	What will happen?
MM	MM	M	
NN	NN	N	

Observed: 0.179 0.502 0.319 → $\text{Allele Freq.} = P + Q = 1$ (Allele Freq.)

$$\text{Expected: } p^2 \quad 2pq \quad q^2$$

$$= P(M) + \frac{1}{2} MN = P(M) + \frac{1}{2} (P + Q)N = P(M) + \frac{1}{2} N = P(M) + \frac{1}{2} (1 - P) = P(M) + \frac{1}{2} - \frac{1}{2} P = \frac{1}{2} + \frac{1}{2} P = \frac{1}{2} + \frac{1}{2} (0.179) = 0.319$$

All should be portion instead of #s.
i.e. 0.179 = $\frac{\# \text{ of MM indv.}}{\# \text{ of total indv.}}$

$$\text{Chi-Square: } \chi^2 = \sum \frac{(obs. - exp.)^2}{exp}$$

chi-square val. < critical val.
→ fail to rej. H₀ ⇒ fall into H-W, no evolution.

Deviation from Ho

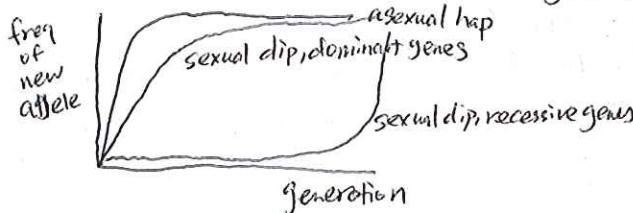
2. Selection

Modeling Selection

most common: asexual haploid
i.e. bacteria, cyanobact. prokaryotes

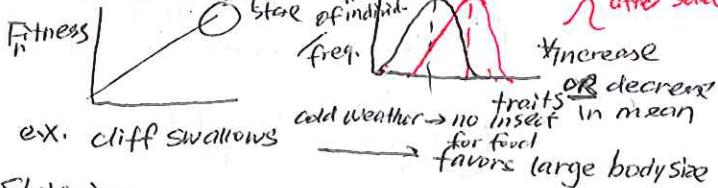
2nd most common: sexual diploid

i.e. multicellular plants, animals, fungi

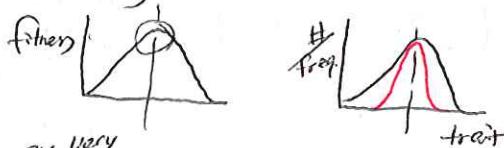


Directions in Selection

① Directional



② Stabilizing



③ Disruptive



Sexual Selection (under natural selection)

selection different reproduce. success \rightarrow nonrandom mating

- male use sexually selected traits to compete for mates

- female choose mates based on

physical character. Signals male genetic behav. charact. \rightarrow ability to provide parental care

quality

explaining/measuring selection

(1) heritability - heritable traits

(2) fitness - related to fitness differences

(3) Change overtime in population under selec. pressure (environment)

3. Genetic Drift

a change in allele freq in a population due to chance

Mendelian Lottery

$\left\{ \begin{array}{l} \text{randomness in meiosis: crossover \& indep. assortment} \\ \text{randomness in gamete fusion.} \end{array} \right.$

$\left\{ \begin{array}{l} \text{Random (non-fitnessrelated) Events: volcanic eruption} \\ \text{conseq. of sampling error} \end{array} \right.$

$\text{zygote } \frac{\text{allele A}}{\text{allele B}}$ due to chance

allele A due to chance

Population reduction: founder effect

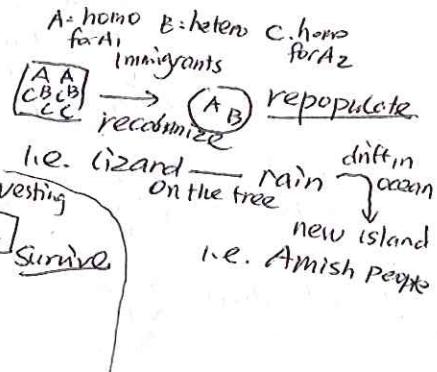
change in allele freq - when a new population's established

Bottleneck effect - sudden red. in allele freq.

Δ random events can be significant in small population (drift most important) over long time periods

Outcomes

- (1) loss of genetic diversity: random fixation & loss of alleles
- (2) random changes to traits may/may not influence fitness



(migration)
4. Gene Flow = immigration of new alleles
or emigration of old alleles → change in allele freq.



Human effect gene flow: (biased gene flow)
hatchery fish - lower fitness
highway - human bandits



Outcomes

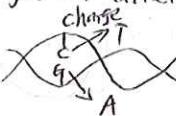
- (1) Homogenization of populations
- (2) Potential breakdown of local adapt., (hatchery fish)
- (3) Potential intro of new alleles

5. Mutation = physical changes in alleles

- Point Mutation

change in nucleotide

seq. → diff. protein



i.e. sickle cell anemia

less O₂ uptake

- DNA Duplication (chromosome-level)

change in # or composition of chromosome

XX → XXX change in # or gene deletion

- Inversion: single chromosome rearrange after breakage

XX → XXX change in composition

- Lateral gene transfer

Gene transfer from one species to another
or inclusion of DNA from environment

i.e. Finches get carotenoid pigments from food

genome from fungus → pea aphid → synthesize own carotenoid

Outcomes:

- (1) Intro of new alleles

Primary mech for generation & genetic variation

- (2) Slowest/weakest evol. force

(3) works w/ other 3 forces
w/out mutation, evolution will stop

Natural Selection



Variation

~↑, or ↓

avg. Fitness

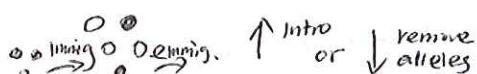


Genetic drift



↑ or ↓ (most)

Gene Flow



↑ Intro or ↓ remove alleles

→ ↑ or ↓

Mutation



/ ↑ or ↓ (most)

example p47)

deletion of red genes

→ green aphid genome

→ avoid predation

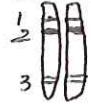
09 Coevolution

Coevolutionary interactions

Within-species

A. Genetic correlations and pleiotropy

chromosome A



chrom. B.

genes located close to each other on the SAME chromosome tend to be related

Gene 1, 2, 3 correlated

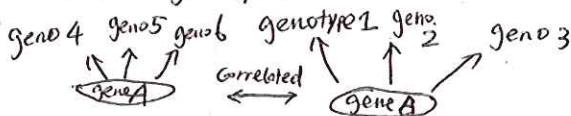
Gene 1, 2 more correlated & ↑ likely to exchange together

Gene 3 could be uncoupled by crossover

Gene 4 not correlated w/ any gene here.

Pleiotropy: 1 gene has multiple trait/phenotypic effect

Selection on gene A can cause correlated responses in other genotypes



① gene B highly beneficial (selected for)

Gene A Slightly deleterious

⇒ gene A maintain in population

② Genotype 3 highly beneficial

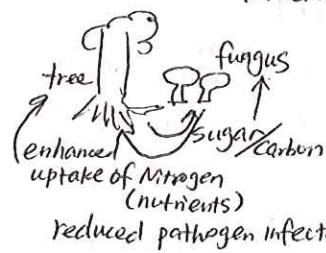
Genotype 1 slightly deleterious

⇒ genotype 1 maintain in population

i.e. P-53 in human { Prevent rep of mistakenly broken cells in mitosis (highly beneficial)
- avoid cancer aging (deleterious)

Among species

B. Mutualistic interactions



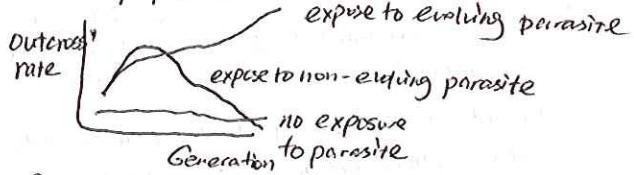
processes/functions
nitrogen uptake
communication
disease detection
nutrient cycling

Honeyguides (birds) & humans

C. Predator-prey

(1) Red Queen hypothesis

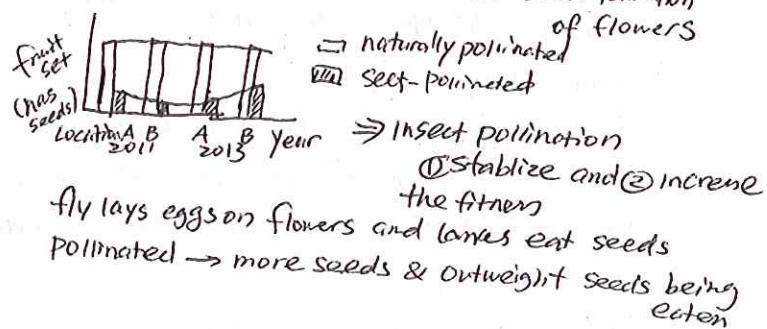
Worm both produces sexual & asexually. (more cost) + deadly parasite see 8-1



- const: ① host evolve to fight off parasite ② parasite evolve to fight off defense mech. co-evolu.

(2) predator-prey Song et al 2014

H0: the fly has no effect on the seed formation



① stabilize and ② increase the fitness
fly lays eggs on flowers and larvae eat seeds
Pollinated → more seeds & outweigh seeds being eaten

10. Macroevolution

A. Paleontology & History of Life

trace of an extinct organism

Fossil: morphology, behavior; rarely genetic info

- phenotypic record of the history of life

Formation of Fossil

- Compression = organism cemented into rocks by sediment LEFT: external
e.g. leaf imprint, dinosaur footprints phenotype
- Casts/Molds = buried in sediment, then decomposes to leave external cast or internal molds LEFT: external or internal phenotype (shell)
- Permineralization: mineral infiltrate cells & replace organic material w/ rock LEFT: ex or internal phenotype, 3-D shape, fine structures (dinosaur)
- Intact remains: protected (e.g. in amber), decap. step.
LEFT: entire organ, possibly genetic info

Limitations / Bias

- Taxonomic/Tissue: hard shells > soft parts
- Abundance = large population size
- Habitat = fast sedimentation (forest, deserts & marine syst.)
Anaerobic (less decap.)
- Temporal: Younger > older rocks

Bias in Biology

- (day)
- Ecology: terrestrial (land), diurnal organisms
 - Evolution: short gen times, simple genetic syst.
 - Genetic: model organisms, human medical correlates

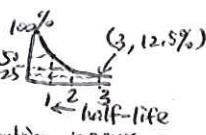
Interpretation of Fossil



- Adaptive Radiation: origination of large number of lineages from a common ancestor in a short time
- Driven by:
 - new niches available (ecological opportunity, environ.)
 - evolution of new morphology → useful niches (change)
 - e.g. feathers originally in display → gliding & fly → dinosaurs → birds
 - e.g. Cambrian Explosion (Radiation): animals w/ different body plans occurred
 - ↑ O₂ levels → ↑ body size (adaptation)
 - Ca²⁺ in oceans from volcanoes → shells, skeletons
 - Arm race bet. predator-prey → coevolution

History of Life

Radiometric dating: isotope decay



Earth formation	First record of life	Orig of Eukaryotes	first animals	Cambrian explosion	K-pmbs extinction	human
4.6 b	3.7 b (prokaryotes)	1.8 b orig. sexual reprodu.	1.2 b	600m 540m	65m 200,000	

B. Speciation

Biological Species Concept (BSC)

- reproductive isolation (no hybrid, no gene flow)

(1) prezygotic isolation

(time) Temporal (breed at different times)

(space) Habitat (breed in diff. habitats)

(behavioral) Behavior (courtship displays differ) song type of birds

(aftermating) Gametic barrier (egg/sperm incompat.)

(morpho) Mechanical (incomp. rep. structures, sizes)

(2) postzygotic isolation

(embryos) Hybrid Viability (hybrid offsp. die as embryos)

(adults) Hybrid Sterility (mature but sterile as adults)
horse \times donkey
male \downarrow
mule

Probs

(1) difficult to apply in field studies (need to be close geographically)
Species need to be co-localized to demonstrate barriers

(2) cannot test in fossil records/specimens (sample)

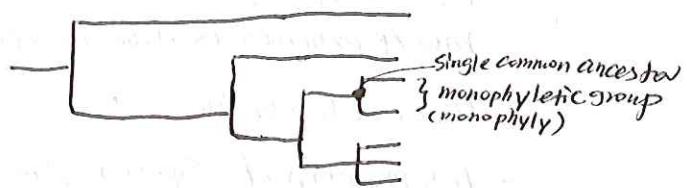
BSC = observ. of habitat, behavior difference

(3) does not apply to asexual, parthenogenetic, selfing poput
~~茱萸子~~ ~~茱萸子~~

(4) plant species - many can hybridize

Phylogenetic Species Concept (PSC)

(evolutionary)



- need to generate hypotheses

- species defined by stas. differences in diagnostic traits (DNA) among populations \rightarrow testable!

Probs

(1) not too many good phylogenies

(2) controversy of trivial traits
may result from mutation & drift

(3) # of extant species would \uparrow w/ PSC

process of Speciation

1. Isolation of populations

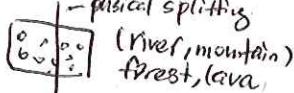
physically/Aeupatric (different location) isol.

= dispersal, colonization



founder effect

= vicariance



physical splitting

(River, mountain)
forest, lava

Genetic/Sympatric (same-hab) isolation.

= change in chromosomes

3. Reproductive Isolation of poput

population diverged

if hybrid, offspring \rightarrow ↓ fitness

\rightarrow selection for assortative mating

\rightarrow Reinforcement \rightarrow speciation

2. Divergence between populations

= Genetic drift

= Selection = speciation via adaption to novel environments

i.e. apple tree - novel food source
apple fly vs. hawthorn fly

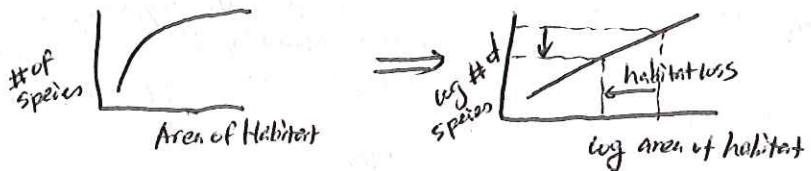
C. Extinction

flip-side of speciation

rate of extinction is the concerning element of current biodiversity crisis

- Direct Observations

- Interpretation of Species-Area Curves



Evidence/observations from Pimm et al. 2014.

1. rate of human-induced mass extinction is 1000 times stronger than background rate of extinction
2. Species w/ small geographic ranges were more at risk of extinction

Solutions

1. protected terrestrial areas are slowing extinction. Marine areas less effective
2. Crowd-sourced citizen science provide crucial data about real-time extinction

Mass Extinction

global, broad range, rapid effect

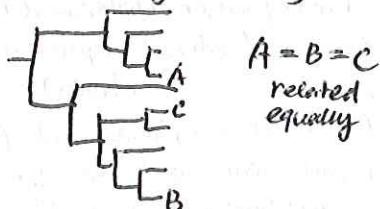
K-P mass extinction

meteonte impact, (high concentration of Ir in impacted layer, shocked quartz, microtekites, impact crater)

effects of impact: SO₂, dust, fires, earthquake, volcanic
cooling, darkening ↑CO₂, ↑Temp
loss of lineages = dinosaurs, etc. (long term)

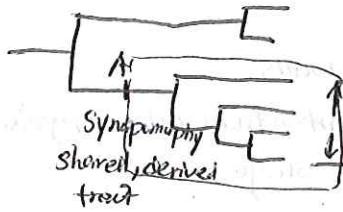
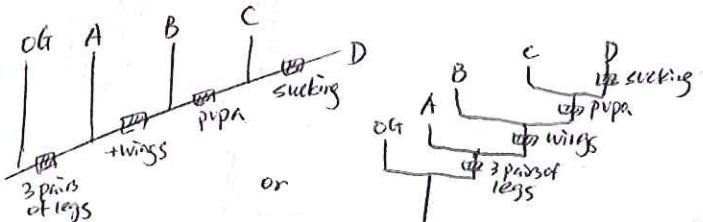
11. Phylogenetics

- nodes can be rotated
- find common ancestors
Instead of counting nodes when finding relations



- Collecting data to construct phylogeny
morphological, molecular (occasionally behavioral)
develop., physiological)
must be homologous across all taxa studied
data must vary, have different character states
- From data matrix to a tree
parsimony rule

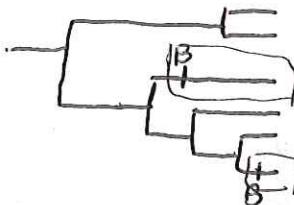
Synapomorphy		Outgroup	A	B	C	D
# of pairs of legs	3+	3	3	3	3	3
wings	absent	abs.	pres.	pres.	pres.	pres.
develp mode	Direct	Dir	Dir	Pupa	Pupa	
multiparts	Grinding	Gri	Gri	Gri	Sucking	



homology: similarity thru common evolutionary descent (shared ancestry)

monophyletic group

Ancestral population + all its descendants, no others

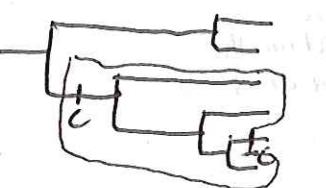


Homoplasy = shared traits not thru shared ancestry

Analogy: similarity due to convergent evolution

polyphyletic group (NOT shared ancestry)
does not include the most recent common ancestor

Analogy: similarity not due to shared ancestry

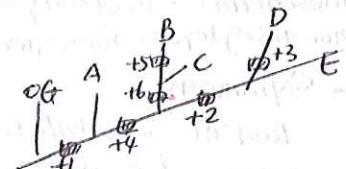


paraphyletic group

does not include all descendants from the common ancestor

Site/trait	OG	A	B	C	D	E
1	0	1	1	1	1	1
2	0	0	0	0	1	1
3	0	0	0	0	1	1
4	0	0	0	1	1	0
5	0	0	0	1	1	1
6	0	0	1	0	0	0

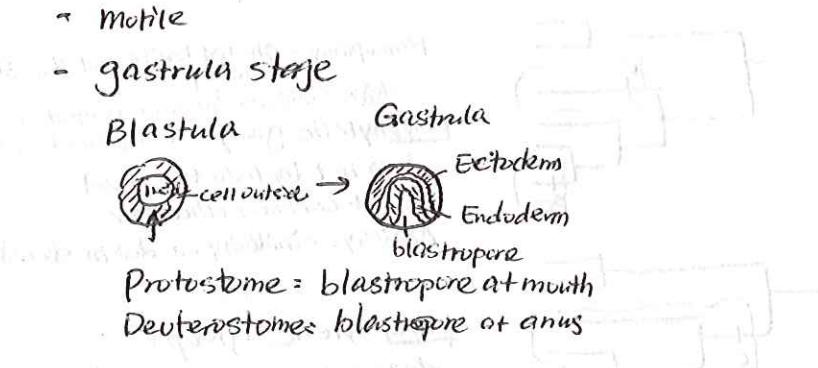
$A = 1$
 $B = 1, 4, 5, 6$
 $C = 1, 4, 6$
 $D = 1, 2, 3, 4$
 $E = 1, 2, 4$



traits can add or loss

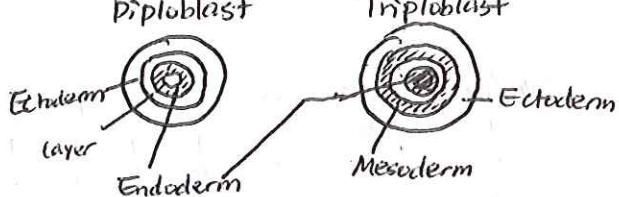
12 Animals

- multicellular, lack cell walls
 - Heterotrophs : obtain carbon from other organisms
 - embryos show gastrula stage
 - Collagen : fiberlike protein
 - Mobile
 - gastrula stage



Key synapomorphies / traits

- Tissue layers: Diploblast vs Triploblast



Ectoderm: skin, nervous syst

Endoderm: lining of digestive tract

Mesoderm: circulatory sys., muscle, organs, bones

non-living layer = mesoglea 中膠層

- ## - Symmetry

Radial vs bilateral

bilateral { distinct head, tail

direct ways to find prey, seek shelter

Sensation organs

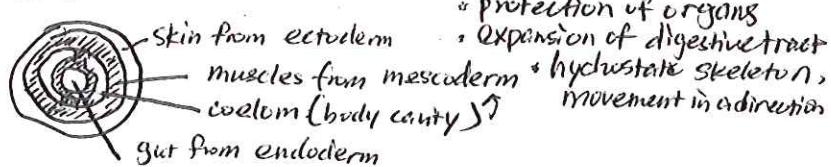
Structure on sides for movement/sensing

- ## - Central Nervous Syst. & Cephalization

Radial = diffuse nervous net

Bilateral = CNS, cluster nervous sys, directional sensation
sensation in head (Cephalization)

- custom



Coelemates = ~~coelom~~^{body cavity} surround internal organs
(digestive tract)

Acoelomates: no coelom survives

Pseudo coelomates = body cavity does not completely surround internal organs

- ## - Hox genes

group of related genes that control the body plan of an embryo

Small variation in location and expression
Timing gives variation in body plans

- ## - other features

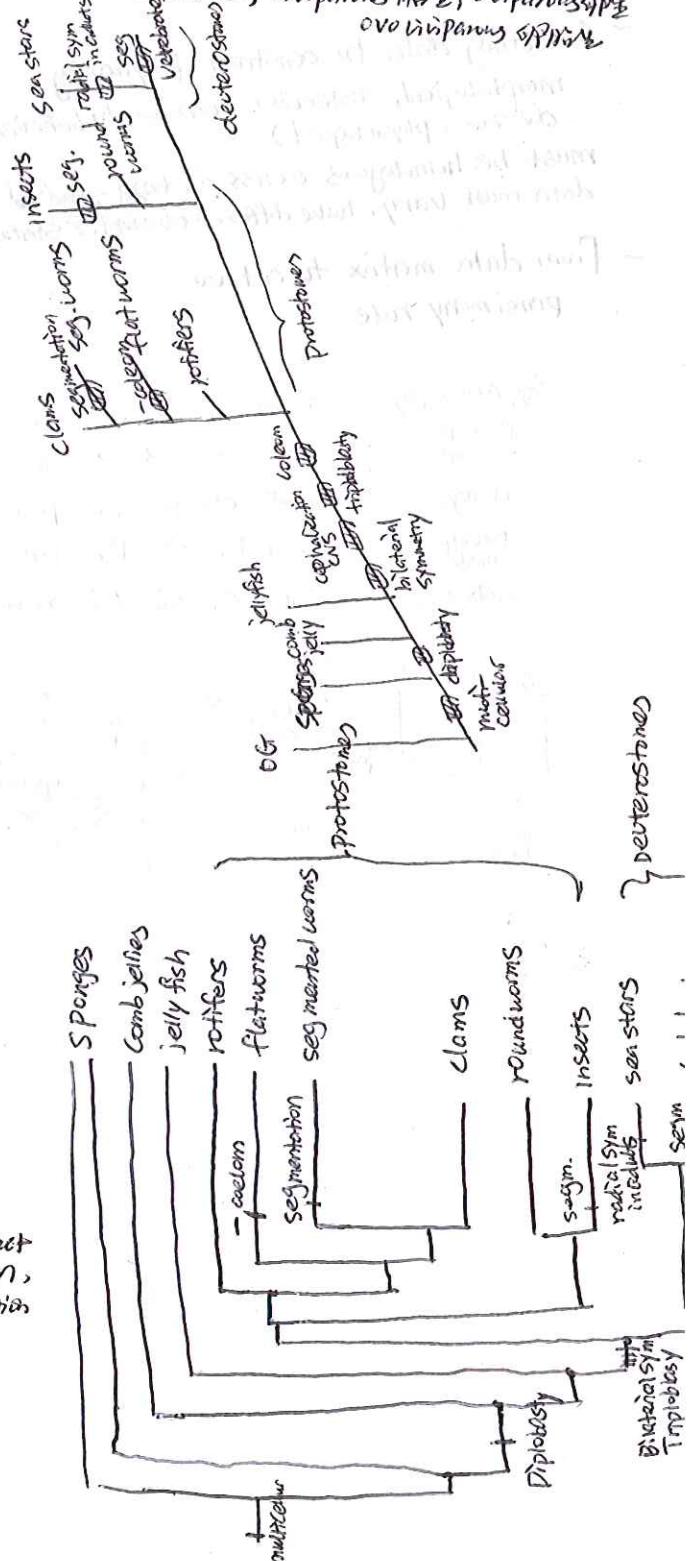
Sensory organs (light, sound, touch, electromagnetic field)
types of feeders (suspension, fluid, deposit, chunks)

limbs (lobe-like, jointed, tube feet, tentacles)

reproduction (sexual, asexual, internal fertilization)

in shark, viviparous type, oviparous type
ovoviviparous common

Ovulipanus spp.



13 Plants

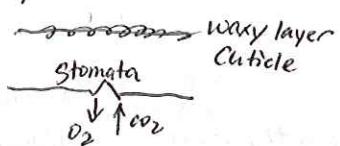
A. Societal importance, Biological importance

- Fuel & energy, food, Secondary compounds (drugs, rubber, glue, oil etc.), quality of life
- photosynth, carbon sink, slow down run-off, Primary producers

B. Key Evolutionary Events

1 - Origin of Chloroplast from cyanobacteria within a eukaryotic cell (endosymbiosis)

2 - Ability to live on land



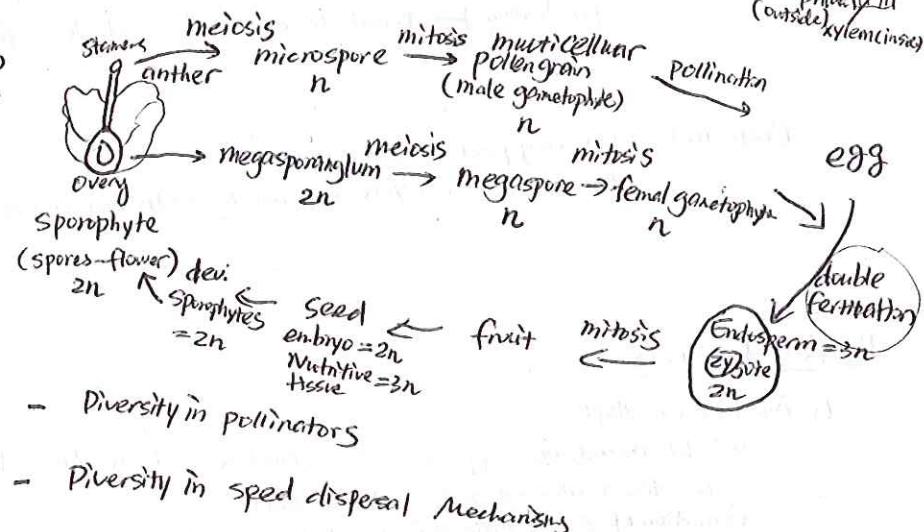
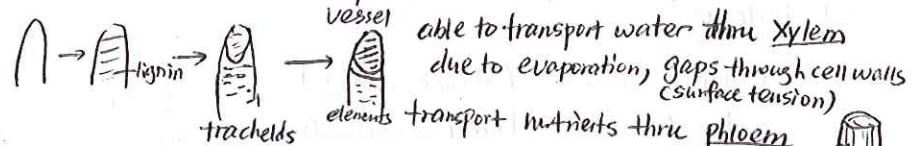
4 - Seeds protected in an Ovary
Gymnosperm vs Angiosperm



key features

3 - Vascular Tissue

water and nutrient transportation

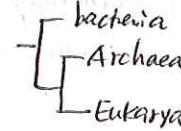


- Diversity in pollinators

- Diversity in seed dispersal mechanisms

14 Protists

- paraphyletic: all eukaryotes except for fungi, animals, and plants
- multicellular
- endosymbiosis
- cell wall outside or inside plasma membrane
- motile: cilia, flagella, amoeboid motion
- sexual/asexual reproduction



A. Importance

Chagas disease (zoonotic disease)

虫媒传播 (传播途径)
bug takes a blood meal → epimastigotes in midgut

传播媒介 (传播途径)

→ multiply in midgut → trypomastigotes in blood → blood meal
(in human body) → penetrate cells → multiply in tissues → enter bloodstream

Ecological = - primary producers in oceans

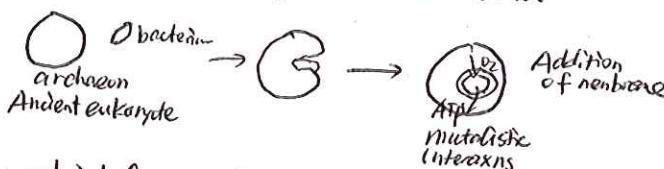
- marine carbon cycle (C into hard structures on ocean floor)

B. Key Features

1. Nuclear envelope

Infold membrane separated from plasma membrane (mutation, selection)
Separates transcription from translation
Formation of endoplasmic reticulum (ER): translation of RNA to protein

2. Endosymbiosis - origin of mitochondria



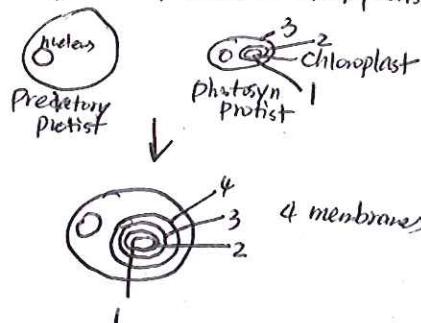
3. Amoeboid form, Supercell

Pseudopodia moved by actin and myosin against a cytoskeleton
Supercell = amoebas fuses to form a single cell w/ multiple nuclei

4. Plant Chloroplast

Primary endosymbiosis in plant

Secondary endosymbiosis in other protist clades



C. More Lineages in Detail

1. Red Algae

can fight off viruses and break down cell wall of virus → health

2. Dinoflagellates

Ecological: cause red tide

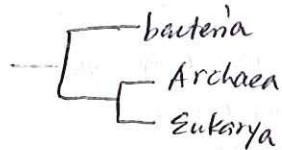
Toxic algae bloom
Kill animals, hypoxia
Health = toxin into human food chain
through oysters, clams, etc.

3. Foraminiferans

morphology: CaCO_3 shells

bio indicators of pollution
indicators of paleoecology and time periods

15. Bacteria and Archaea (Prokaryote)



Bacteria, Archaea differences

Bacteria

- peptidoglycan in cell wall
- transcription differs from archaea

Prokaryotes

- single-celled
- no membrane around DNA, single circular chromosome
- no energy-producing organelles
- very few cell compartments
- no sex
- flagella present

Archaea

- phospholipids from isoprene in cell membrane
- transcription similar to Eukarya

Values of Studying them

- old
- a lot!
- can live in extreme conditions

- medical

- if virulence is heritable } most bacteria not pathogenic
- a large genome for protein toxin } \rightarrow pathogenic bacterium
- bioremediation

key features

- morphological diversity

Size, shape, motility

Gram-positive bacteria = cell wall made of peptidoglycan

- negative : polysaccharides in outer membrane

- horizontal Gene transfer

conjugation = cell to cell contact

transformation = take up DNA from the environment

- Metabolic Diversity

photoautotrophs

energy

carbon

example

analogous

sun

CO₂

Cyanobacteria

plants, algae

chemoautotrophs

chemical rxn

CO₂

unique to prokaryotes

photoheterotrophs

sun

environment

heliobacteri

N/A

Chemoheterotrophs

environment

many eukaryotes, animals
fungi

16. Viral Diversity

Definition of the virus

- obligate intracellular parasite, dependent on host cells
- parasite of biological information systems

Can propagate

- through an extracellular stage
- Within the host genome

1. Are viruses Alive?

- No:
 - Not made of cells
 - No metabolic systems, capable of synthesizing biological molecules
 - No enzymatic system to generate ATP
 - No ribosomes

- Yes:
 - life is self-propagating
 - can be inactivated
 - can be reactivated, like seeds, be active under right conditions
 - "borrowed life"

* Importance

- Subject to evolution
- important role in the evolution of other forms of life

2. Key features

- Morphology



capsid or one or more membranous envelop



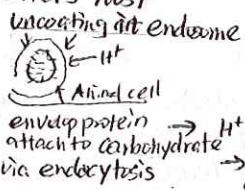
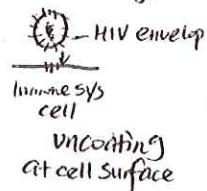
- Reproduction

Lytic cycle

animal: host specific receptor

plants: insects and damage cell wall

① Viral genome enters host



Env protein

attach to carbohydrate

H⁺ pump
via endocytosis
charge and fuse

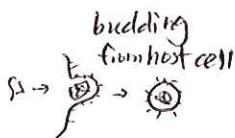
viral genome release

② Viral genome is transcribed, production of viral protein

③ Replication of viral genome

④ Assemble particles inside the host

⑤ exit and release to environment or new host



Burst from host cell (non-enveloped)



3. Why study viruses?

- Disease outbreak

- Global ecology

Algae w/ Caco₃ shells → viral infection die, sink → carbon sink
→ producers

- Diversity of life

- Red-queen hypo: viruses will continually adapt to host's defenses via natural selection, mutation
- high mutation rate in virus
high RNA polymerase error rate
lack repair enzymes

- Diversity of viruses

genomic reassortment generates new flu strains

2 different strains → infect → replication, → reassortment of genomic seg. → new strain!

phylogenetic & biogeographical info to understand emerging strains

continuous immunization, flu shots

geographically-predicted

17. Lyticco-Bodig Disease

Lyticco: progressive paralysis

Bodig: parkinsonism, occasionally w/ dementia

Hypothesis by Oliver Sacks

1. Human Genetics

colonized by Spanish, Huge bottleneck ~1% survived (heritable mutation)
family pattern

BUT, other ppl immigrated also developed this disease

2. Cycad - derived toxins

seeds made into flour

3. Mineral Deficiency or metal poisoning

Ca^{2+} , Mg^{2+} deficiency in water

Dinoflagellate toxins in local fish

4. Viral (pison or bird)

brown tree snake killed all birds

Hypothesis (other)

Consumption of flying foxes ("Conservation Implications of Chemono consumption of flying foxes")
Clark Morrison et al.

difference in body parts male & female eat

fruit bats of cycad toxin bioaccumulation

flying fox eradicated - overlapping the end of disease

18. Research

1. Establish Questions, hypothesis, predictions
critical analysis of others work, literature searching, discussing ideas
2. Design a methodology and completing it
methodology actually test your hypothesis, analyze data, techniques, skills practicing.
3. Drawing Conclusions from findings
4. Communicate findings and conclusions

Required Hw

prof Allison

In cell proliferation, in response to changing environment, requires extensive communication between cells. Understanding the transcription mechanism by which cells enter and exist cell nucleus in response to extracellular signals.

prof Saha

how cell acquire and maintain their unique identity during early vertebrate embryonic development, by focusing on nervous system

prof Bradley

single proton emission detectors to image gene expression