

LIFE on Earth

- chemical basis (carbon-based life on water-planet) proteins, sugars, fats, nucleotides
- genetic code (DNA)
- replication, transcription, translation
 four bases (2 bit (binary digit) code)
- cellular organization (membranes, organelles, nuclei)
- basic units of life
- evolution (natural selection, survival of fittest)
- mutation, recombination
- symbioses (living together)

 - organelles (SET) organisms (life styles)
 - → collective co-existence (ecology)



Ecology (hierarchy) • biosphere (all environments on Earth inhabited by life) ecosystems (all living and non-living things within given area) (matter recycles while energy flows through) communities (all species within given area) interactions between species, e.g. food chains, competition, predation, herbivory, disease) populations (all individuals of single species) (single species distribution and abundance)

Population ecology

- distribution
 - temporal (any time frame, but esp. seasonal)
 - spatial (any space, but esp. regional)
- abundance
 - size (number)
 - density (number/area)
 - concentration (number/volume)
 - intensity (e.g. number parasites/host)
 - prevalence (e.g. proportion infected)
 - incidence (change in prevalence over time)

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Population growth

Dynamics (numbers) + kinetics (time)

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- reproductive population (births)
- senescent population (deaths)
- open population (immigration/emigration)
- closed population (no migration)
- discrete change (step-wise)
- continuous change (constant)
- unconstrained (infinite resources) (exponential)
- constrained (finite resources) (logistic)

Population characters

- · demography (categorize!)
 - biology (age, sex, breed, size ...)
 - other (income, postcode, religion, ...)

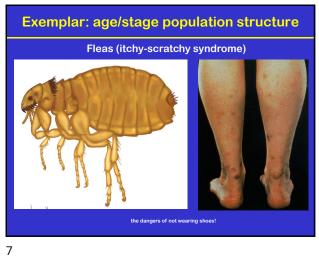


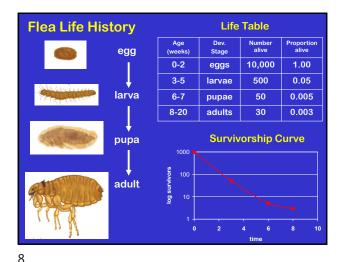
- · life tables
 - age/stage-structure, numbers
 - births, deaths, migrations



- survivorship curves
 - life expectancy

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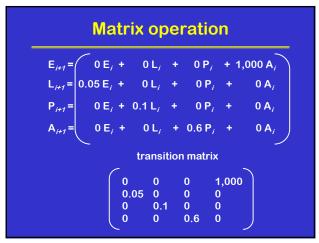


Stage	Stage-structured diagrams		
used to depict life-cyclesarrows show proportion in transition			
Time period	Flea developmental stage	Number alive	
1	Eggs	10,000	
2	Larvae	500	
3	Pupae	50	
4	Adults (50% female) (females lay 2,000 eggs)	30	
	1,000		
(E) 0.0	$ \begin{array}{c c} \hline E & 0.05 \\ \hline L & 0.1 \\ \hline P & 0.6 \\ \hline A \end{array} $		

Simultaneous equations 1,000 0.05 $= 1,000 A_i$ 0 E, + $0 P_i + 1,000 A_i$ $L_{i+1} = 0.05 E_i +$ $0A_i$ $= 0.05 E_i$ $0A_i$ $= 0.1 L_i$ 0 E, + $= 0.6 P_{i}$ 0 L, + 0.6 P; + 0 A,

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Matrix multiplication assume population N_0 = 100 E, 50 L, 10 P, 2 A calculate N₁ 100 50 10 2,000 5 5 6 1,000 100 0.05 0 50 0.1 0.6

Population growth

- · vital statistics of populations (changes over time)
- individuals join (births and immigration)
- · individuals leave (deaths and emigration)

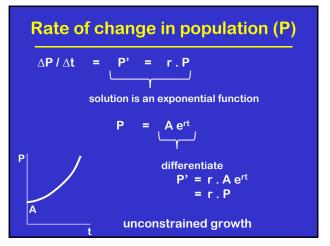
 $P_2 = P_1 + (births + immigration) - (deaths + emigration)$

assume closed population (no migration)
 P₂ = P₁ + (births - deaths)

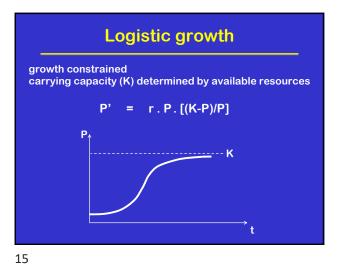
that is, population is proportionate to:

- current size (P)
- growth rate (r) (= birth rate death rate)

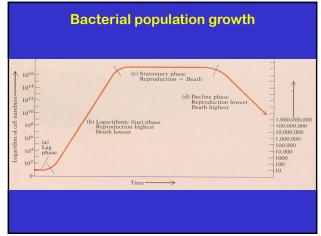
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Epidemiology?

Like ecologists, epidemiologists seek to understand:

- species richness (biodiversity)

- species abundance (populations/communities)

- species distribution (temporal, spatial)

Epidemiology

Study of occurrence, spread and control of diseases

Prevalence (number infected)
Incidence (change in prevalence over time)
Distribution (density, intensity, concentration,..)
exhibit longitudinal fluctuations (esp. seasonal)
influenced by many factors:
demographic, socioeconomic, behavioural
geographic, climatic

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Epidemiological studies Four main types: **Maths** not quantitative Case series (descriptive) index, incidental, miscellaneous statistics Case control studies (retrospective) cases + controls interviewed **Odds Ratio** statistics **Cohort** studies (prospective) cohort followed forward in time **Relative Risk** calculus Outbreak studies (predictive) **Differential Equations** - rate of change in population

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OR and RR Odds Ratio (OR) = AD BC **CASES CONTROLS** Relative Risk (RR) = **EXPOSED** В A / (A+B) C / (C+D) UNEXPOSED С D causative? no association protective?

BUT, nothing is IDEAL

proportion lose resistance (susceptible)

Population varies in susceptibility

 proportion just infected (latent) proportion patent (infectious)

proportion immune (resistant)

proportion die proportion migrate

⇒ all influence Ro

OUTBREAKS Pathogens must reproduce themselves to survive (reproduction rate = R) R>1 epidemic

R=1 perpetuation R<1 extinction time

Basic (case) reproduction rate (Ro) = average number of secondary infections resulting from primary case in susceptible population

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• etc

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Micro-parasites

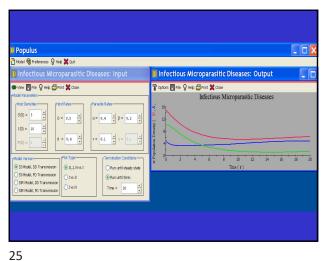
viral, bacterial, protozoal pathogens

- · reproduce quickly to reach high intensities
- thus causing acute transient infections
- often limited by host immune responses
- recovered individuals develop protective immunity

population dynamics driven largely by transmission rate (contact between susceptible and infected individuals)

· famous SIR model

SIR model (N = Susceptible + Infected + Recovered/immune) β = infection rate v = recovery rate γ = immunity loss b = births d = deaths α = infection mortality bN + γ R - dS - β SI - dl - α l - ν l **BSI** R' = ν l - dR - γ R



Macro-parasites helminths and arthropods · cause chronic and persistent infections disease depends on number present (which in turn depends on exposure to free-living infective stages) But many infections over-dispersed (a few hosts have most of the parasites), so must track intensity of infection Anderson & May model: infective stages short-lived Dobson & Hudson model: hypobiosis (larval arrest)

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Anderson & May HPW model
Host population, Parasite population (adults), free-living stages (W) λ = parasite birth rate γ = infective stage mortalit d = host deaths in out H' = - dH - αP bΗ P' = βWH - $\mu_p P$ - αP - dP - $\alpha P^2/H[(\kappa+1)/\kappa]$ - γ**W** - β**W**H $\lambda \mathbf{P}$ W' = 27

Model 📵 Preferences 🖓 Help 💥 Quit 🗱 View 🔚 File 💡 Help 😅 Print 💥 C 👕 Options 🔚 File 💡 Help 🖨 Print 💢 Close Macroparasitic Infections

Disease prediction models SIR model of rubella S(t) 8000 6000 4000 2000 time (weeks)