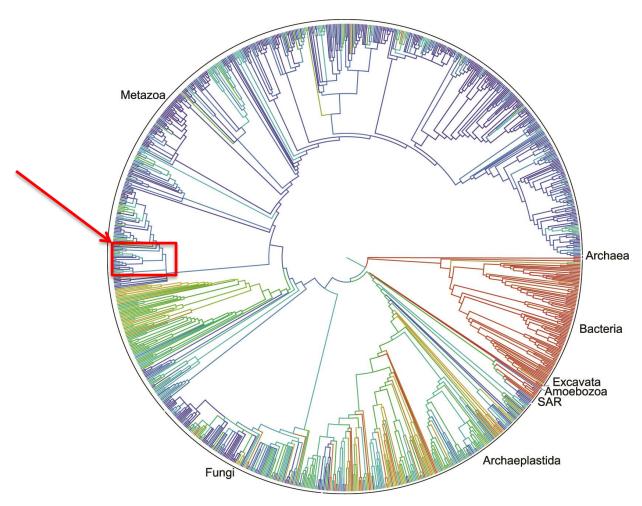
The Lab of Molecular Systematics & Ecology

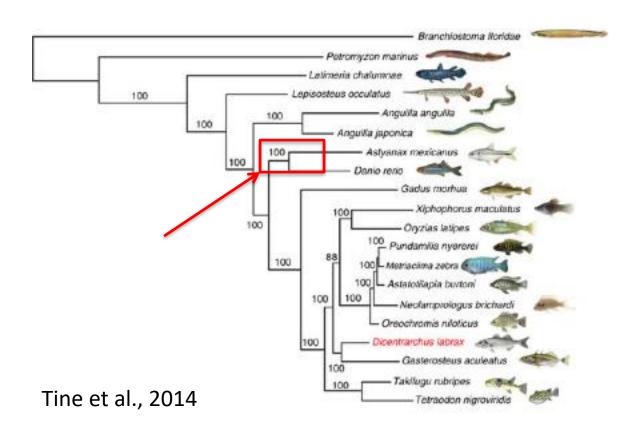
Workshop in Molecular Evolution

Jan 7 - 11, 2019 Shanghai

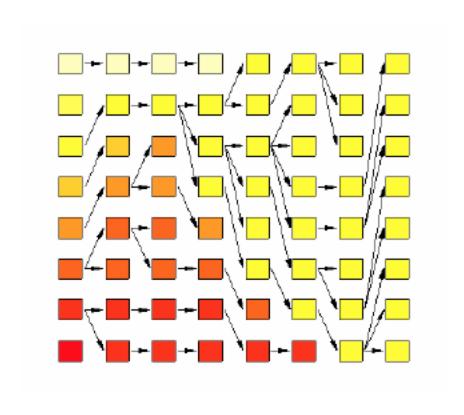
Two faces of one process: phylogenetics vs. population genetics



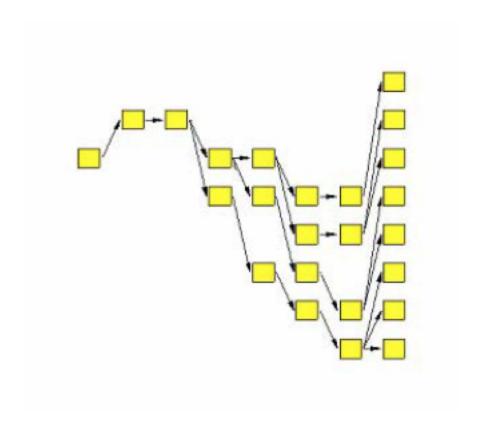
Phylogenetics – model of speciation



Population genetics – model of coalescence



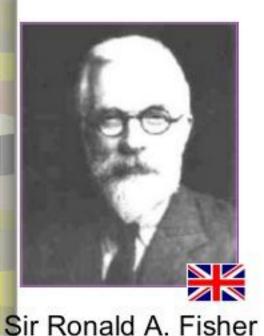
Population genetics



Idealised population (Fisher-Wright population)

- Random mating
 - each copy of the gene found in the new generation is drawn independently at random from all copies of the gene in the old generation
- No selection
- No migration
- No mutation
- Large population size, no drifting

Genética De Poblaciones



(1890-1962)

Sewall Wright (1889-1988)



John B. S. Haldane (1892-1964)

British biologist and statistician Ronald Fisher

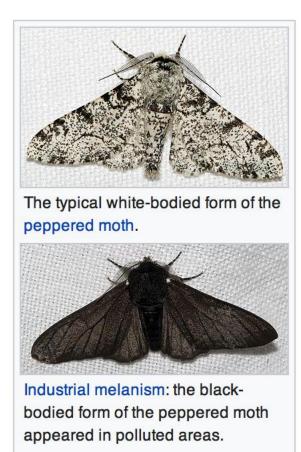


- In a series of papers starting in 1918 and culminating in his 1930 book The Genetical Theory of Natural Selection
- Fisher showed that the continuous variation measured by the biometricians could be produced by the combined action of many discrete genes, and that natural selection could change allele frequencies in a population, resulting in evolution.

British geneticist J.B.S. Haldane

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- worked out the mathematics of allele frequency change at a single gene locus under a broad range of conditions.
- Haldane also applied statistical analysis to real-world examples of natural selection, such as peppered moth evolution and industrial melanism



The American biologist Sewall Wright



- animal breeding experiments, focused on combinations of interacting genes, and the effects of inbreeding on small, relatively isolated populations that exhibited genetic drift.
- In 1932 Wright introduced the concept of an adaptive landscape and argued that genetic drift and inbreeding could drive a small, isolated subpopulation away from an adaptive peak, allowing natural selection to drive it towards different adaptive peaks.

Idealised population (Fisher-Wright population)

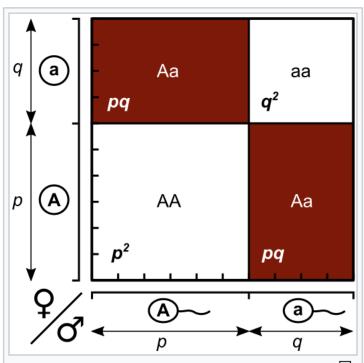
- Random mating
 - each copy of the gene found in the new generation is drawn independently at random from all copies of the gene in the old generation
- No selection
- No migration
- No mutation
- Large population size, no drifting

Idealised population (Fisher-Wright population)

- Hardy-Weinberg equilibrium
 - allele frequencies stay constant over time, genotype frequencies are related to allele frequencies
- Linkage equilibrium

Hardy-Weinberg equilibrium





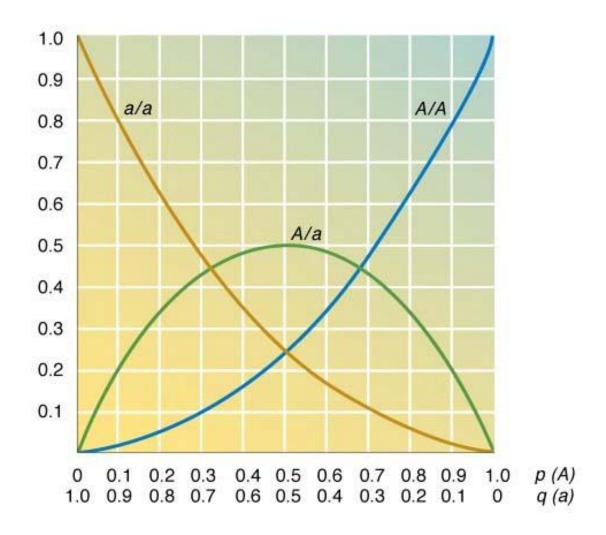
Length of $p,\,q$ corresponds to allele frequencies (here $p=0.6,\,q=0.4$). Then area of rectangle represents genotype frequencies (thus

AA : Aa : aa = 0.36 : 0.48 : 0.16).

One locus with 2 alleles at

HWE: $p^2 + 2pq + q^2 = 1$

Implications of HWE (two-allele locus)



- Rare alleles mostly in heterozygotes
- Common ones mostly in homozygotes
- Maximum He is 0.5 at 2 allele locus, rising to 1.0 with more alleles

Linkage equilibrium

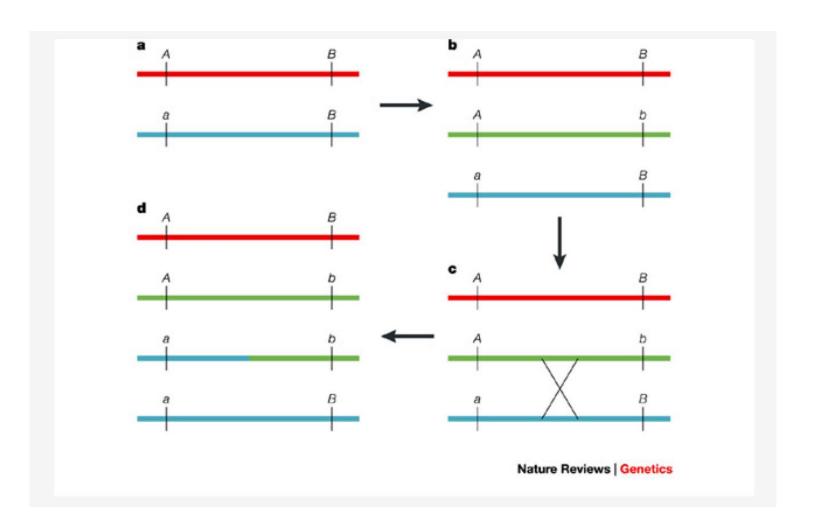
•Alleles at separate loci are expected to segregate independently during meiosis. They show **linkage equilibrium**.

Example:

- 2 loci with alleles A_1 and A_2 ; B_1 and B_2 their frequencies will be p_1 and p_2 and q_3 .
- Possible gametes A₁B₁; A₁B₂; A₂B₁; A₂B₂
- Genotype frequencies will be the product of constituent allele frequencies

	A_1	A_2	Total
B_1	$x_{11} = p_1 q_1 + D$	$x_{21} = p_2 q_1 - D$	q_1
B2	$x_{12} = p_1 q_2 - D$	$x_{22} = p_2 q_2 + D$	q_2
Total	p_1	p_2	

Linkage disequilibrium

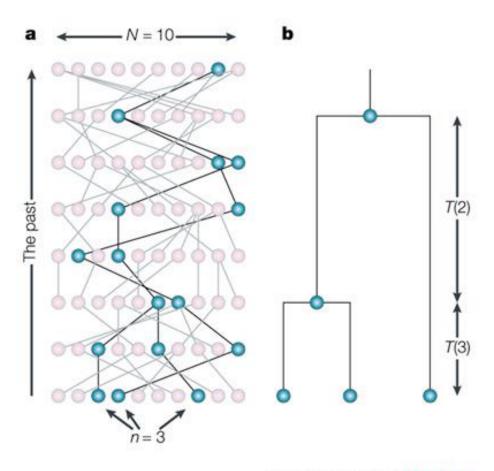


Linkage disequilibrium

Linkage disequilibrium = a deviation from random associations of alleles at different loci

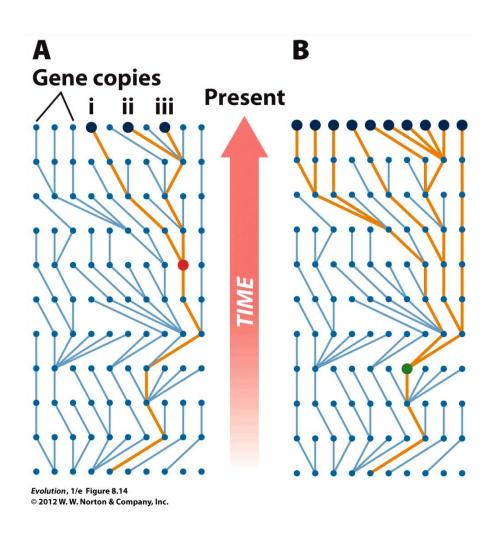
- Linkage disequilibrium can be caused by :
 - chance events
 - population bottlenecks
 - recent mixing of different populations
 - selection
- Linkage disequilibrium is important because:
 - It is common in threatened species with small populations
 - evolutionary processes are altered
 - functionally important genes may exhibit linkage disequilibrium
 - can be a signal of recent admixture of populations

Coalescent theory

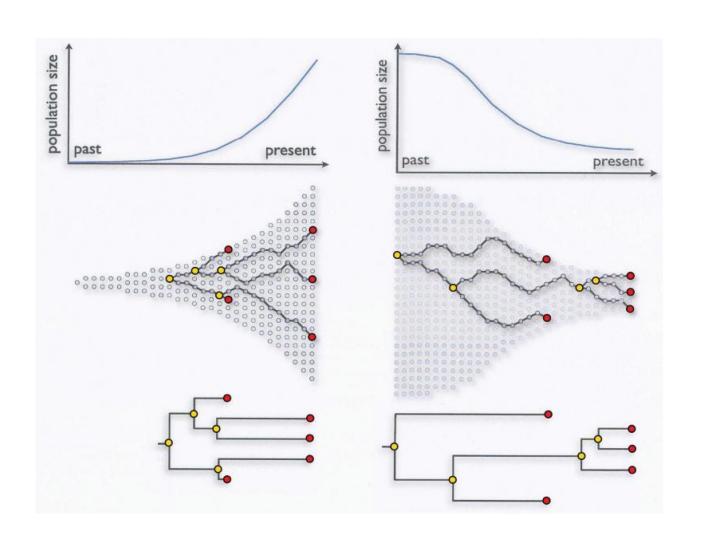


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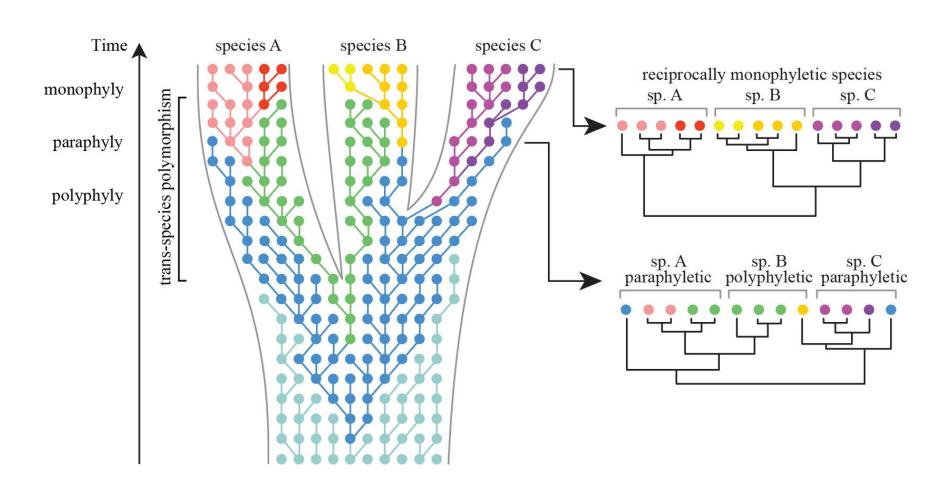
Sampling for coalescent analysis



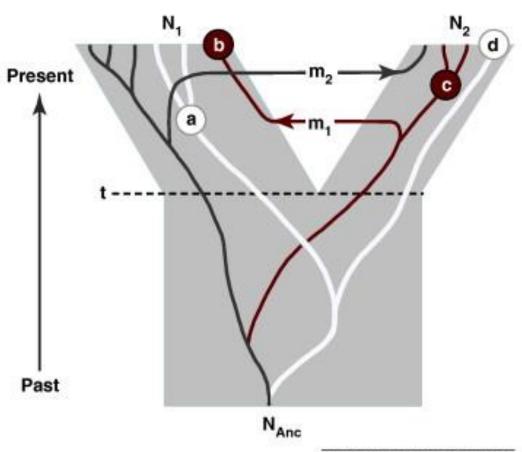
Population growth



Species delimitation



Migration



TRENDS in Ecology & Evolution



Thank you!