Neutrality and some of its deviations

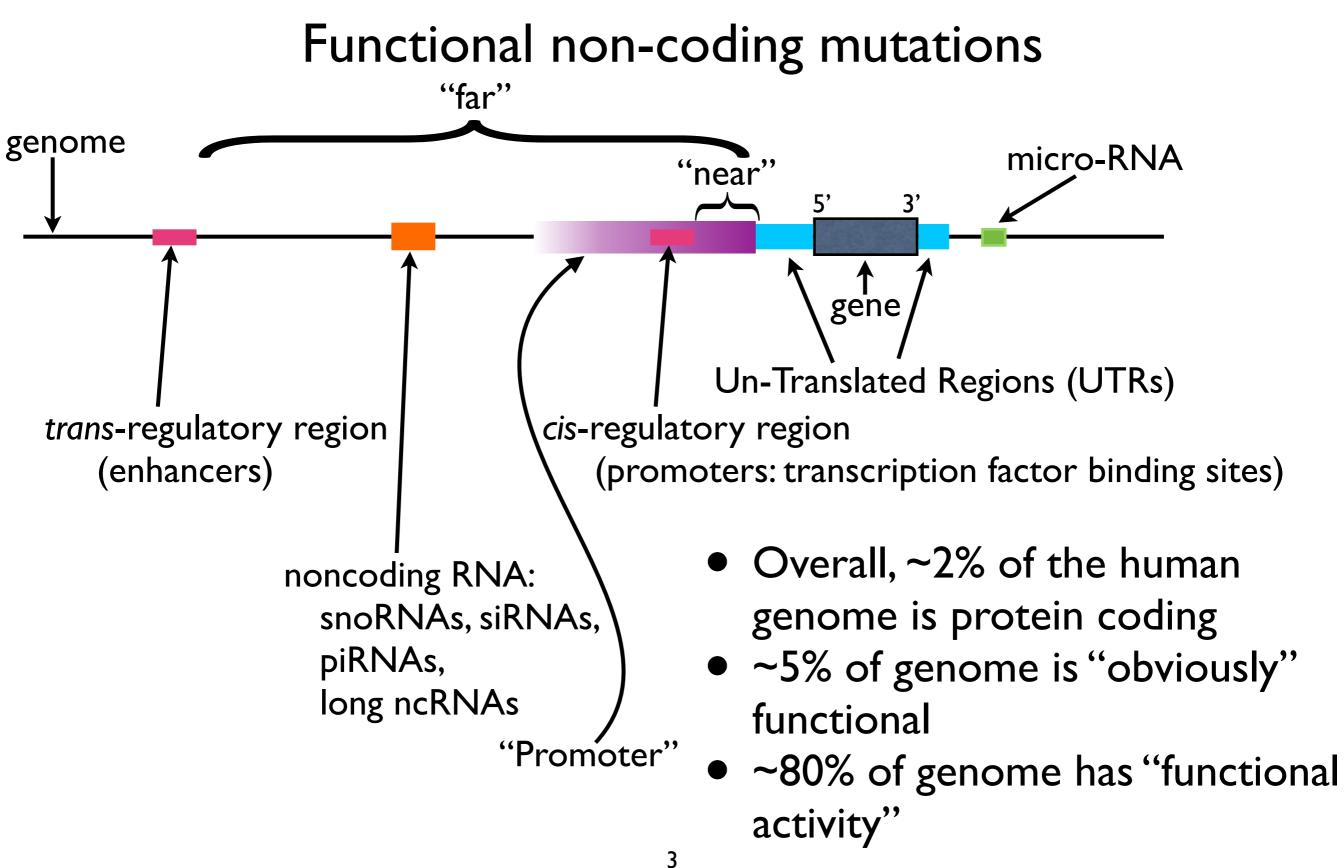
Ryan Hernandez

ryan.hernandez@UCSF.edu

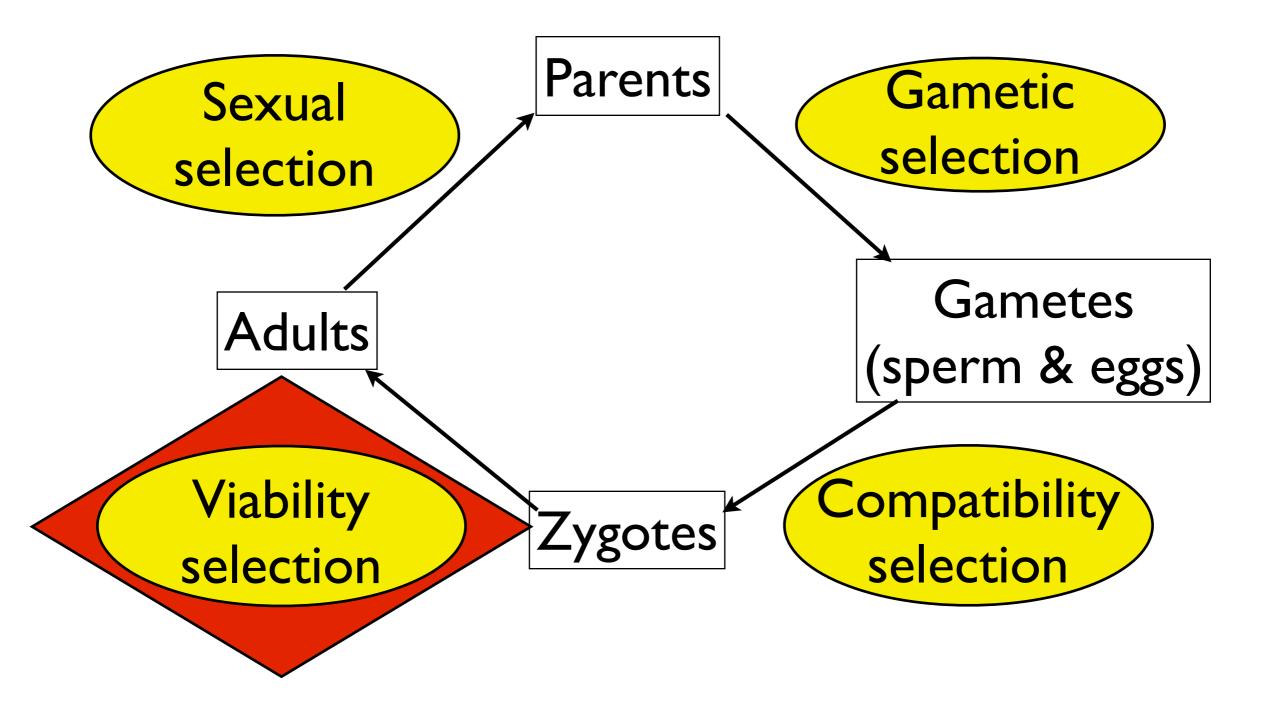
Goals

- Learn about the population genetics view of the life cycle
- A few Pop Gen summary statistics
- Revisit Hardy-Weinberg Equilibrium Assumptions
 & violations

Basic Biology of Human Genome



Life Cycle



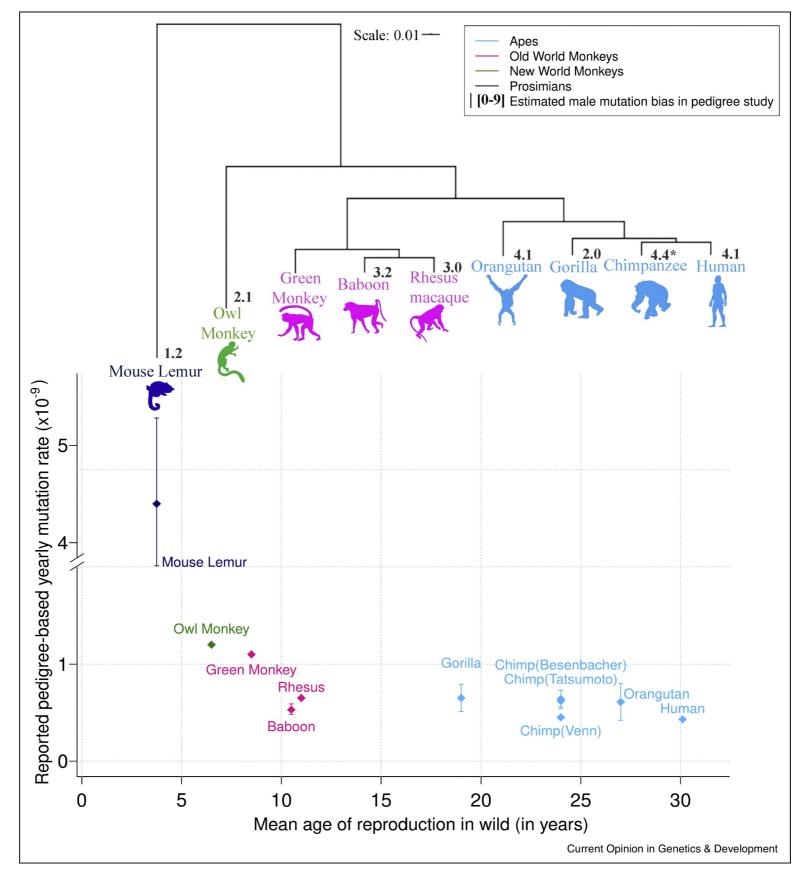
MODERN HUMAN GENOMICS: A CASE FOR RARE VARIANTS?

 $1.1 \times 10^{-8} \times 6 \times 10^{9} = 66$ [muts / person]

66 [muts/p] × 130M [p/y] ÷ 3B [bp]

2.86 muts/bp/yr

MUTATION RATE EVOLUTION IN PRIMATES



Chintalapati & Moorjani (Curr Op G&D, 2020)

<pre># Pairwise differences</pre>	3	4	3	3	3	3	
4	А	С	G	A	С	Т	
3	G	Т	G	А	Т	Т	5
2	А	Т	G	A	С	Т	
1	А	С	А	G	С	С	*
Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6]

 π = average pairwise diversity

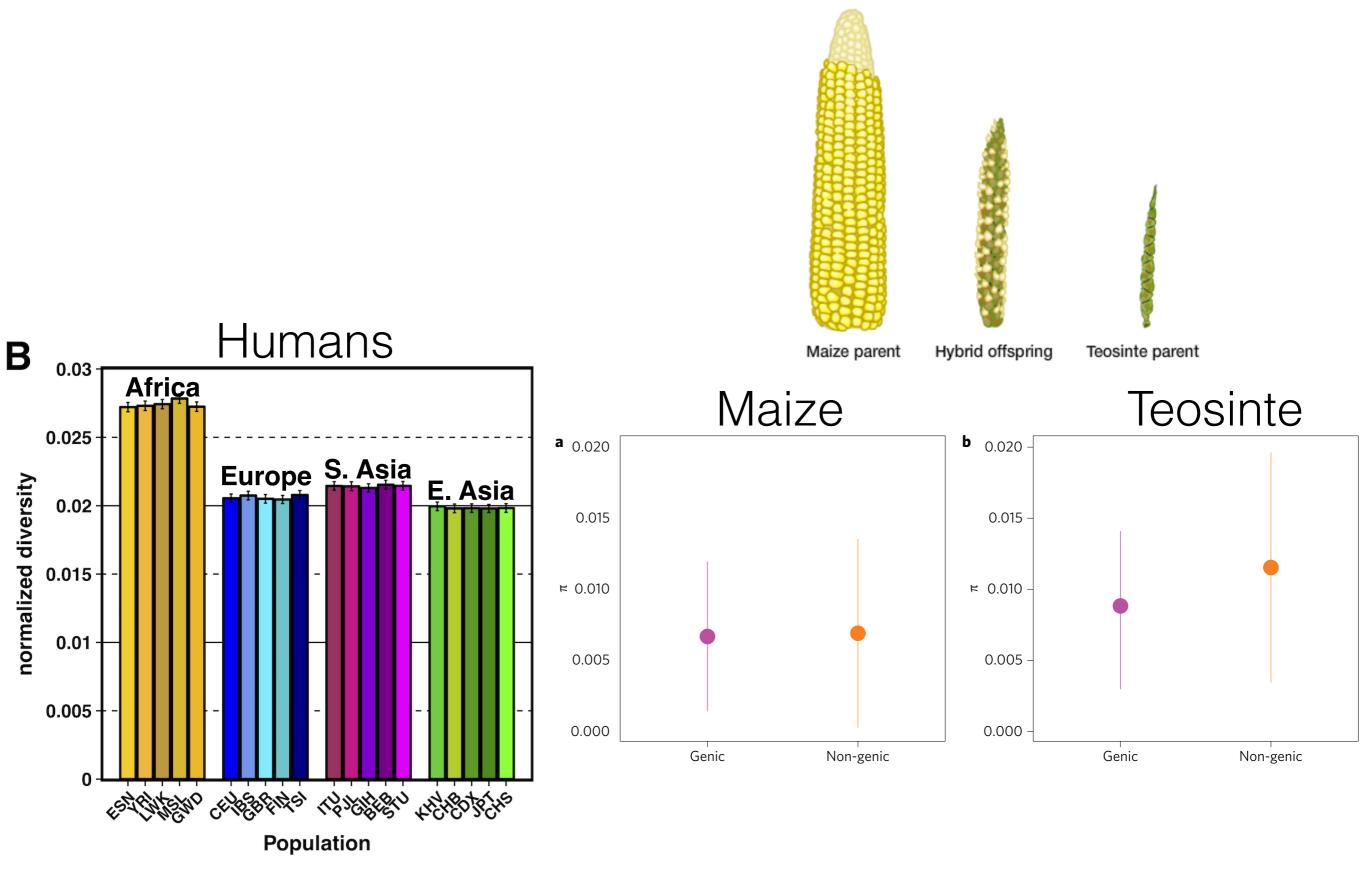
Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6	
1	А	С	А	G	С	С	
2	А	Т	G	А	С	Т	
3	G	Т	G	А	Т	Т	5
4	А	С	G	А	С	Т	
# Pairwise differences	3	4	3	3	3	3	
# Compared	6	6	6	6	6	6	

 π = average pairwise diversity

Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6	
1	А	С	А	G	С	С	*
2	А	Т	G	А	С	Т	
3	G	Т	G	А	Т	Т	
4	А	С	G	А	С	Т	
# Pairwise differences	3	4	3	3	3	3	
# Compared	6	6	6	6	6	6	
Avg. Pairwise Diff	0.5	0.67	0.5	0.5	0.5	0.5	

Number of variants: 6 SNPs Diversity (π): 3.1667/L 6

DIVERSITY ACROSS POPULATIONS



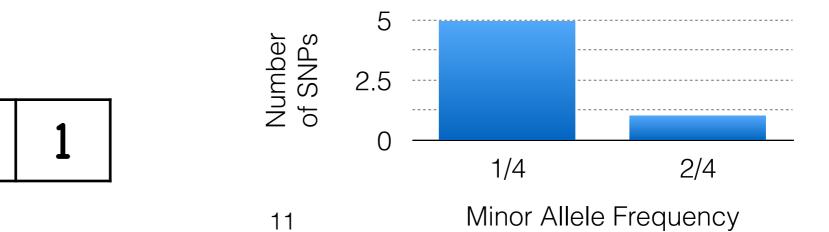
Torres, et al. (2018)

Beissinger, et al. (2016)

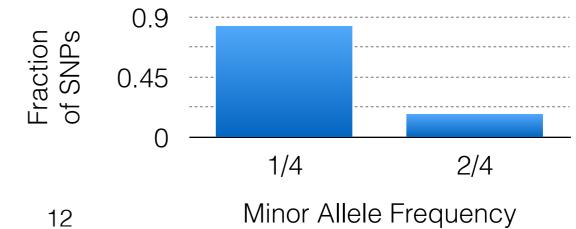
Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6
1	А	С	Α	G	С	С
2	А	Т	G	А	С	Т
3	G	Т	G	А	Т	Т
4	А	С	G	А	С	Т
Minor Allele	G	Т	А	G	Т	С
MAF	0.25	0.5	0.25	0.25	0.25	0.25

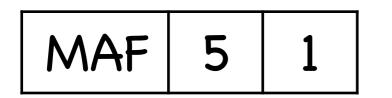
MAF

5



Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6
1	А	С	Α	G	С	С
2	А	Т	G	А	С	Т
3	G	Т	G	А	Т	Т
4	А	С	G	А	С	Т
Minor Allele	G	Т	А	G	Т	С
MAF	0.25	0.5	0.25	0.25	0.25	0.25





Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6
1	А	С	А	G	С	С
2	А	Т	G	А	С	Т
3	G	Т	G	А	Т	Т
4	А	С	G	А	С	Т
Chimp	А	С	А	G	С	Т

Chromosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6
1	А	С	А	G	С	С
2	А	Т	G	Α	С	Т
3	G	Т	G	Α	Т	Т
4	А	С	G	Α	С	Т
Chimp	А	С	А	G	С	Т

Chr	comosome	SNP 1	SNP 2	SNP 3	SNP 4	SNP 5	SNP 6
	1	А	С	А	G	С	С
	2	А	Т	G	Α	С	Т
	3	G	Т	G	Α	Т	Т
	4	А	С	G	Α	С	Т
(Chimp	А	С	А	G	С	Т
	erived count	1	2	3	3	1	1
Proportion of SNPs	0.5 0.333 0.167 0					-Frequ ctrum	

3

Site-Frequency Spectrum (SFS)

Derived frequency in sample

2

1

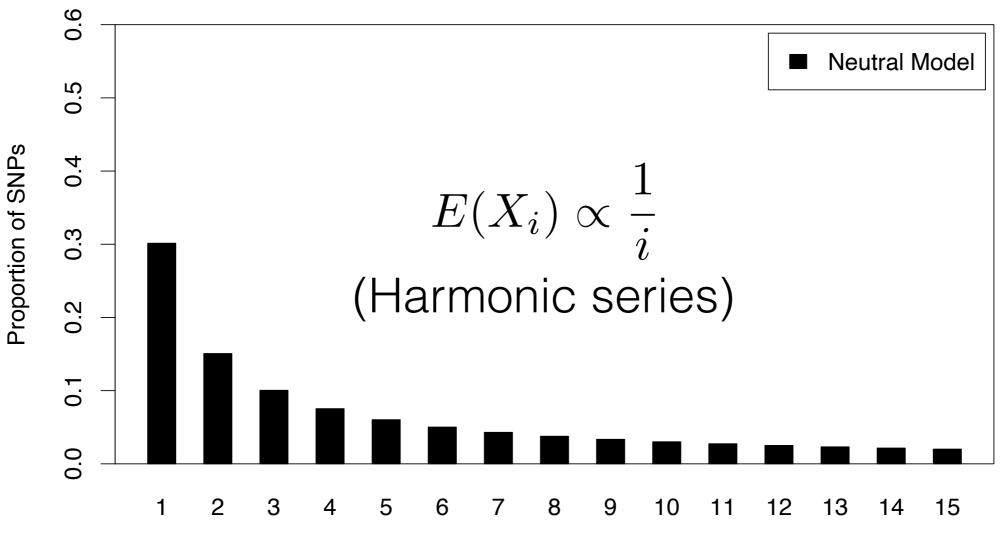
Site-Frequency Spectrum

	*		*	*		*	*			*	*	*	*		*	*	*	*		*	*	*
1	С	A	Т	Н	С	G	Α	Α	G	С	G	A	Т	С	Α	G	G	С	Т	Α	Т	Α
2	С	A	Т	Т	т	G	A	G	Α	С	G	A	Т	С	А	G	G	С	Т	Α	Т	A
3	С	G	Т	Т	Т	G	Α	G	A	С	G	A	Т	Т	A	G	G	С	С	Α	Т	A
4	С	A	Т	Т	С	G	Α	G	A	С	G	Α	Т	С	A	G	G	С	Т	A	Т	Α
outgroup	т	A	С	С	С	Α	G	G	A	G	Α	т	Α	С	G	С	A	Т	Т	Т	A	
					U							Ľ						Ľ				

- = non-coding
- = synonymous
 - = nonsynonymous
- * Substitution between species

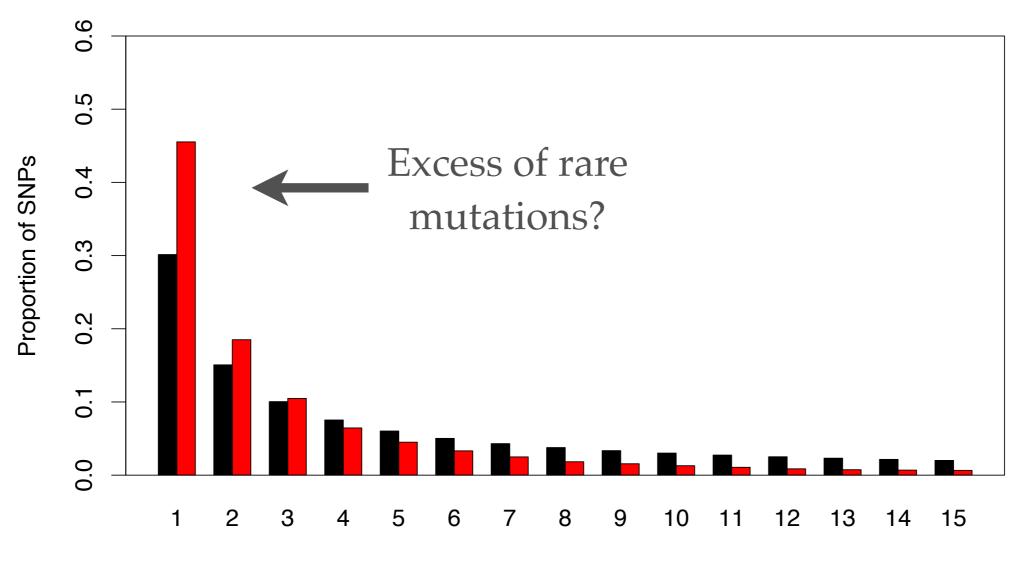
Site-Frequency Spectrum

The proportion of derived mutations at each frequency in a sample of chromosomes



Derived count in sample of 16 chromosomes

SITE-FREQUENCY SPECTRUM



Derived alleles in sample of 16 chromosomes

What evolutionary forces could cause an excess of rare variants? (one-word-answers-please)

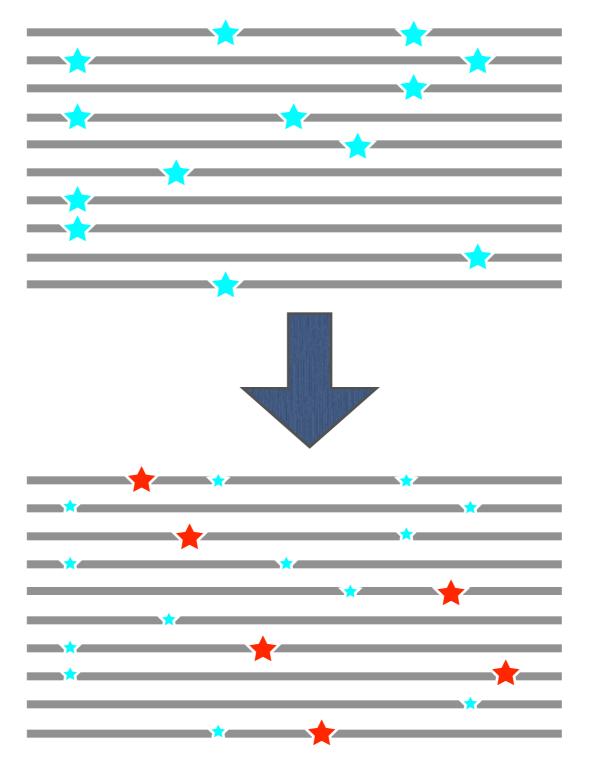
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The Effect of Negative Selection

Chromosomes in	
a population	

The Effect of Negative Selection

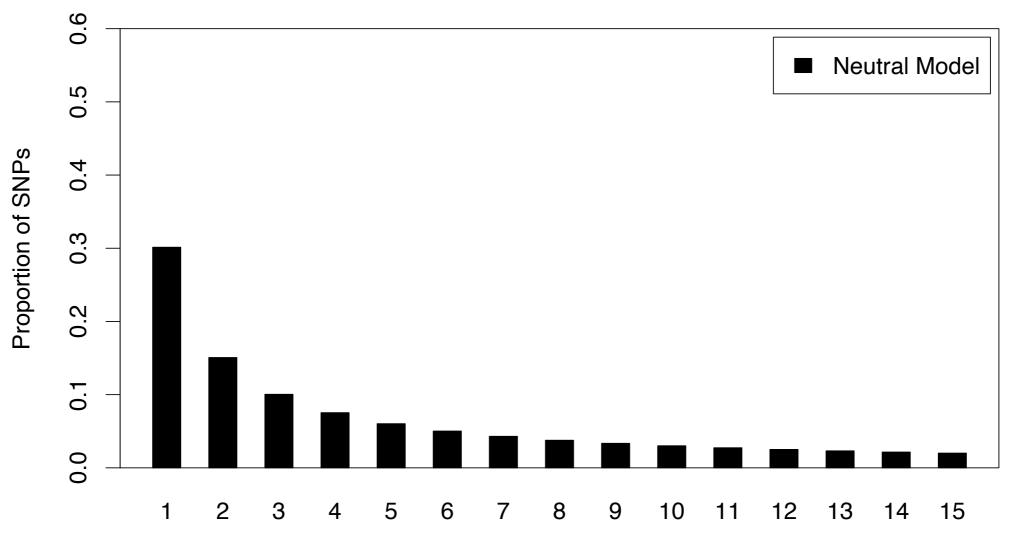
Chromosomes in a population with standing variation



Negative selection: the action of natural selection purging deleterious mutations.

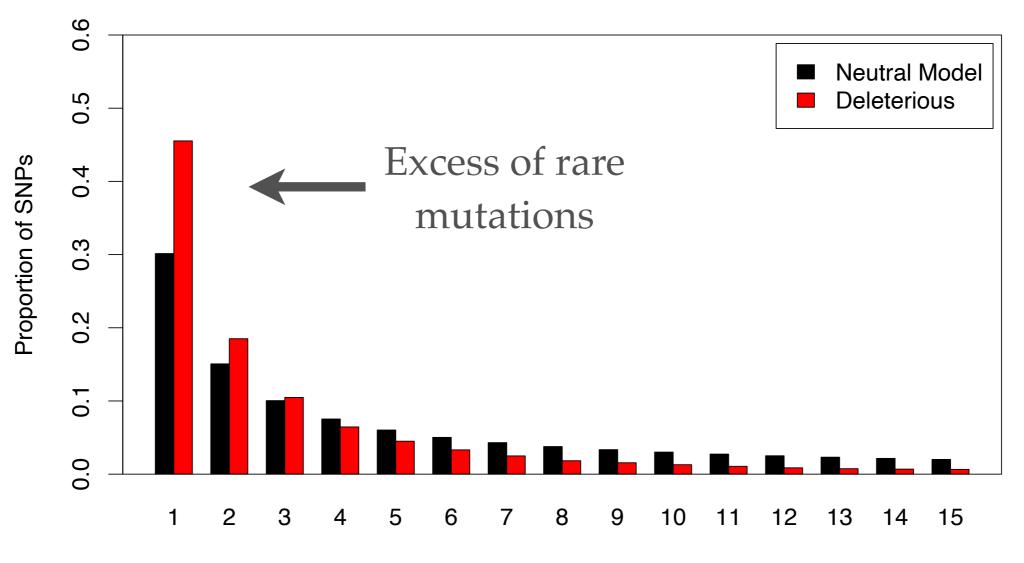
Deleterious mutations will arise in the next generation

Site-Frequency Spectrum



Derived count in sample of 16 chromosomes

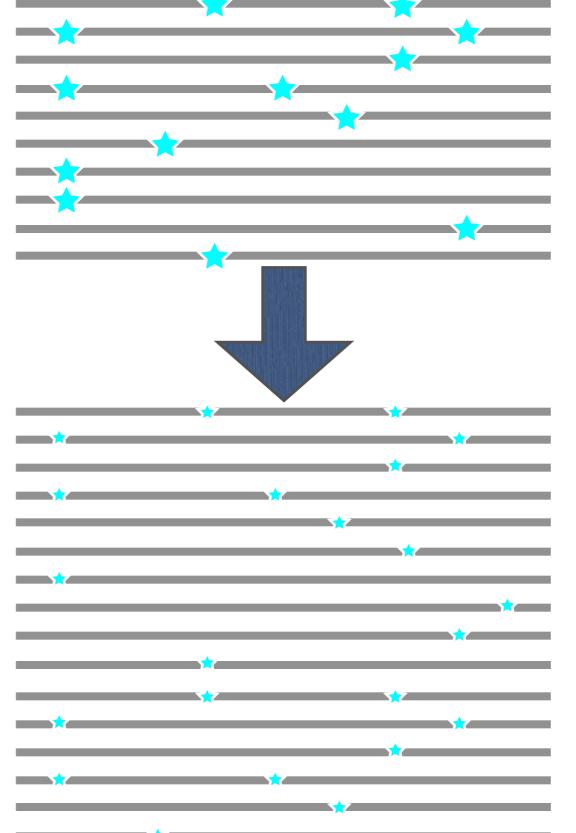
SITE-FREQUENCY SPECTRUM



Derived alleles in sample of 16 chromosomes

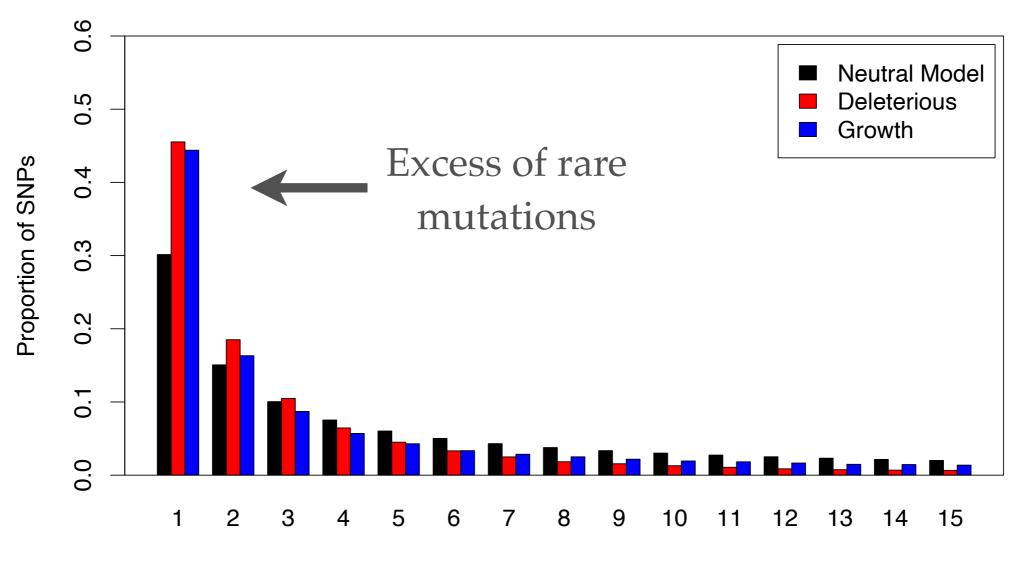
The Effect of Population Growth

Chromosomes in a population with standing variation



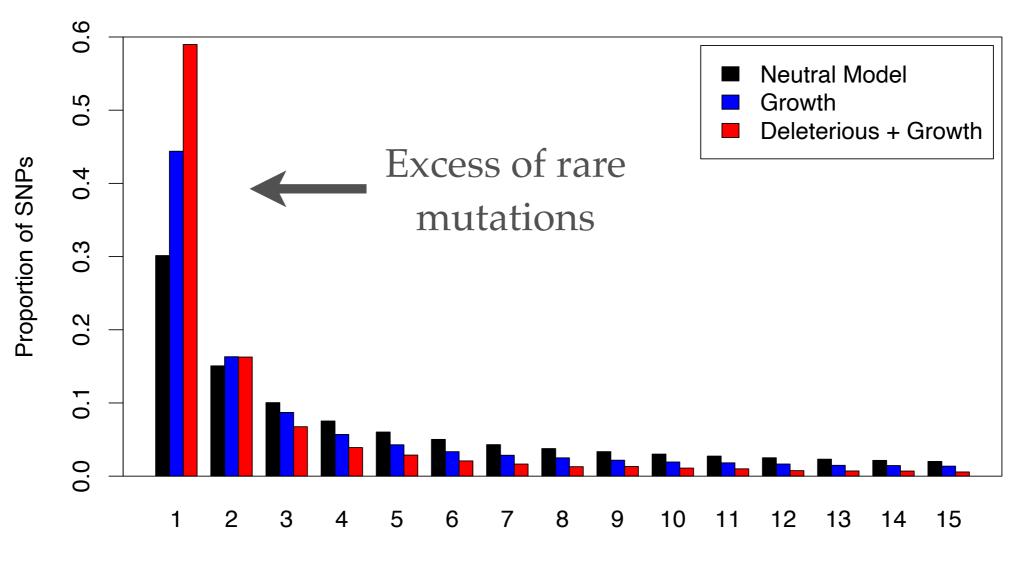
Population Growth: introduction of new (rare) variants.

SITE-FREQUENCY SPECTRUM



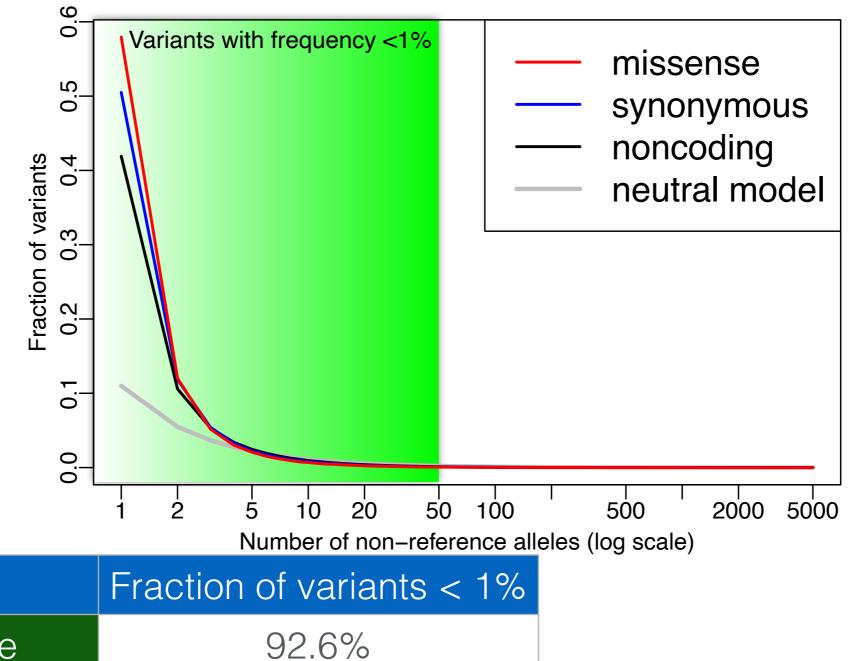
Derived alleles in sample of 16 chromosomes

SITE-FREQUENCY SPECTRUM



Derived count in sample of 16 chromosomes

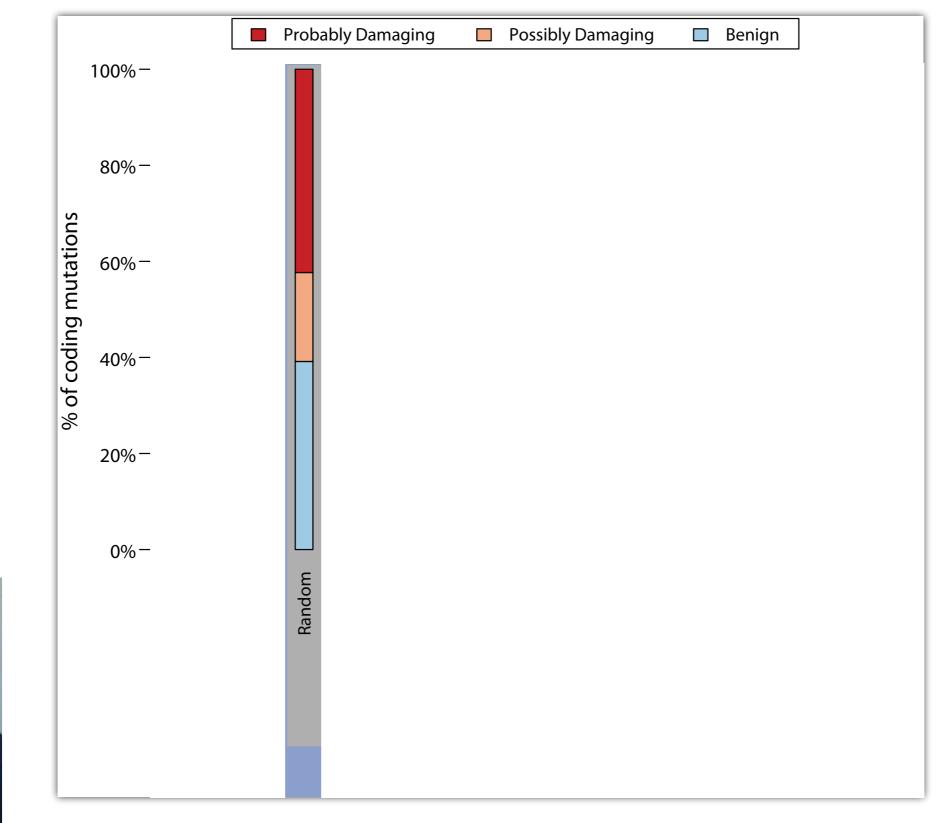
Majority of human genetic variation is rare



Class	Fraction of variants $< 1\%$
Missense	92.6%
Synonymous	88.5%
Non-coding	82.3%



Observed Effect of Selection

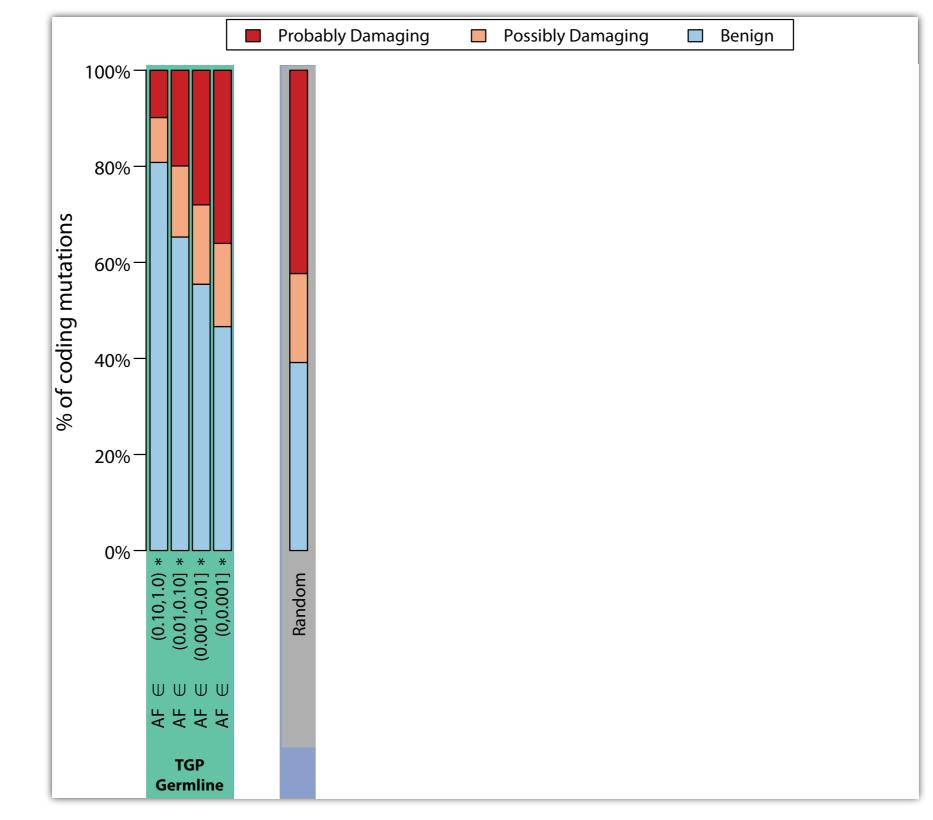


Zach

Szpiech

PolyPhen2

Observed Effect of Selection

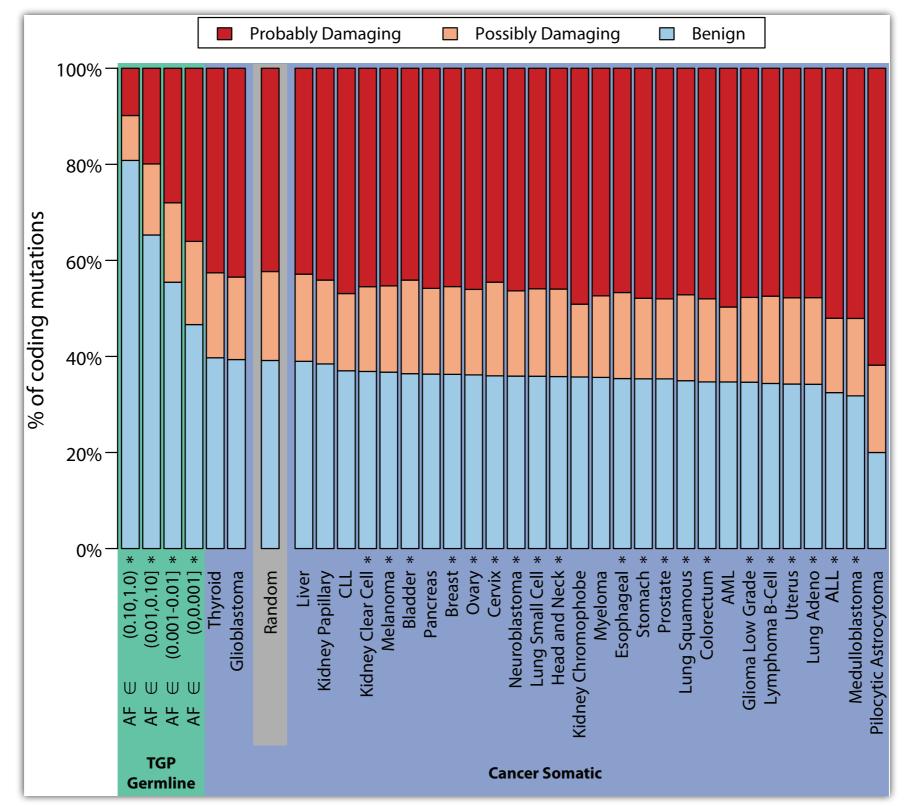


Zach

Szpiech

PolyPhen2

Observed Effect of Selection



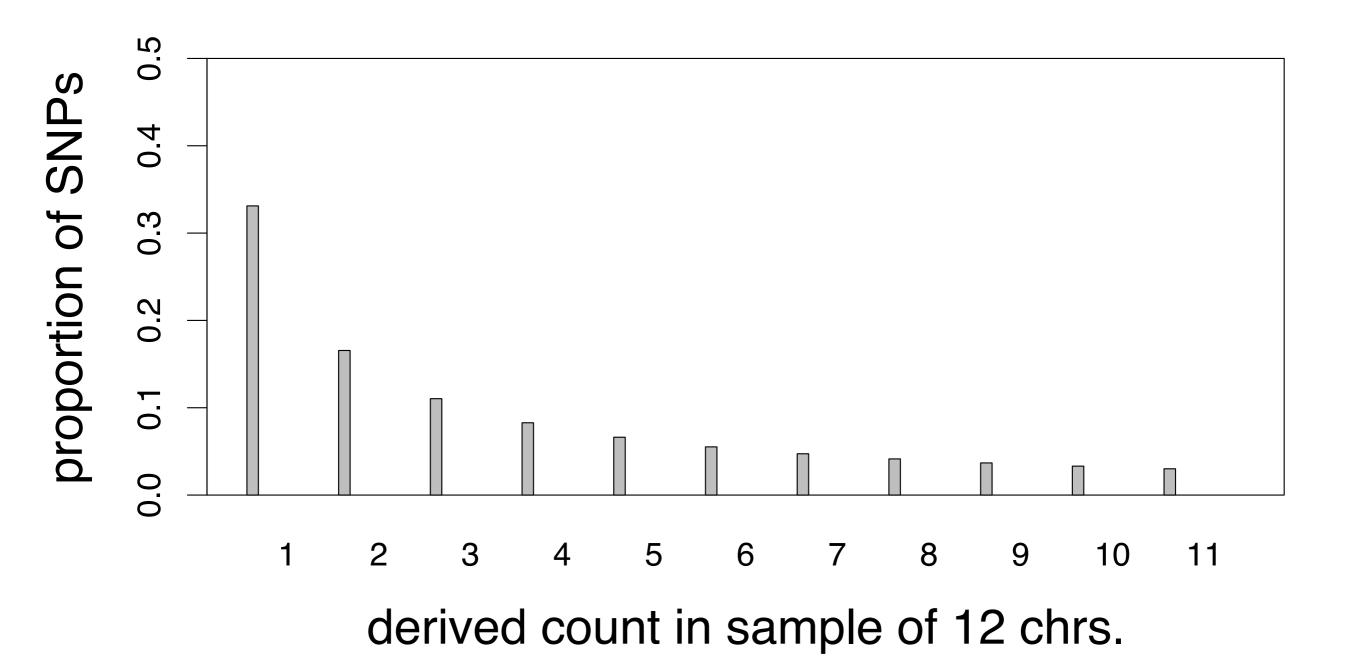
Zach

Szpiech

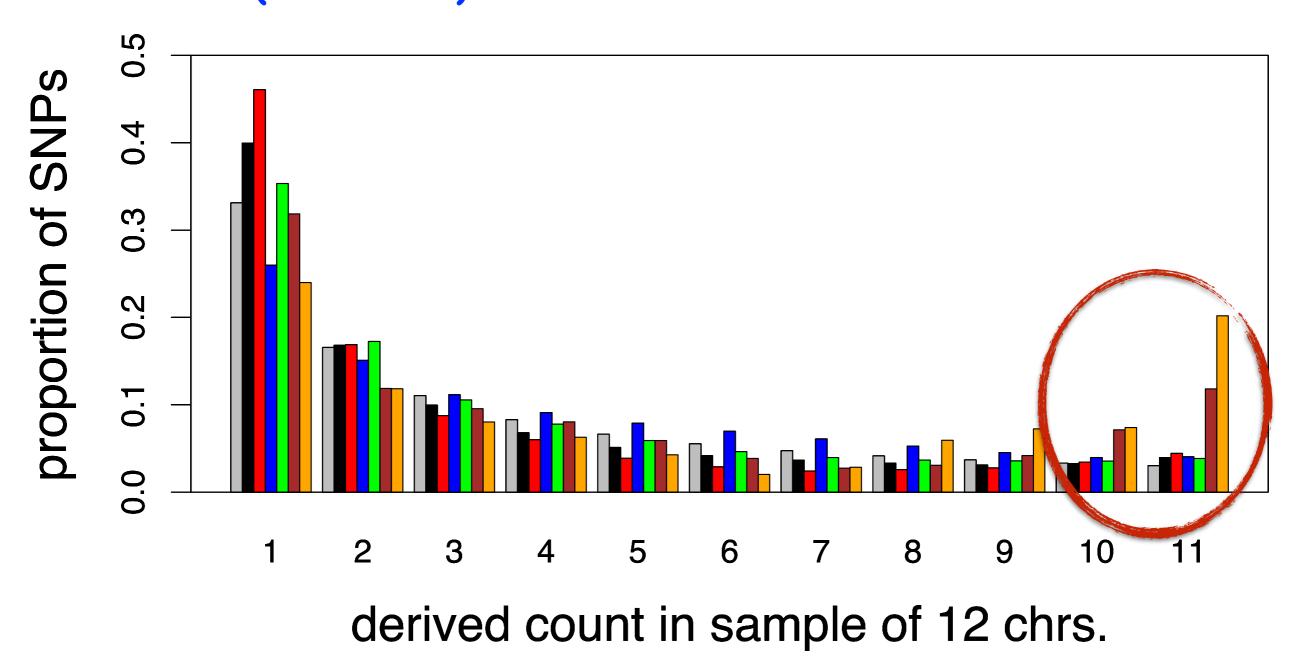
PolyPhen2

Site-Frequency Spectrum

The proportion of SNPs at each frequency in a sample of chromosomes.

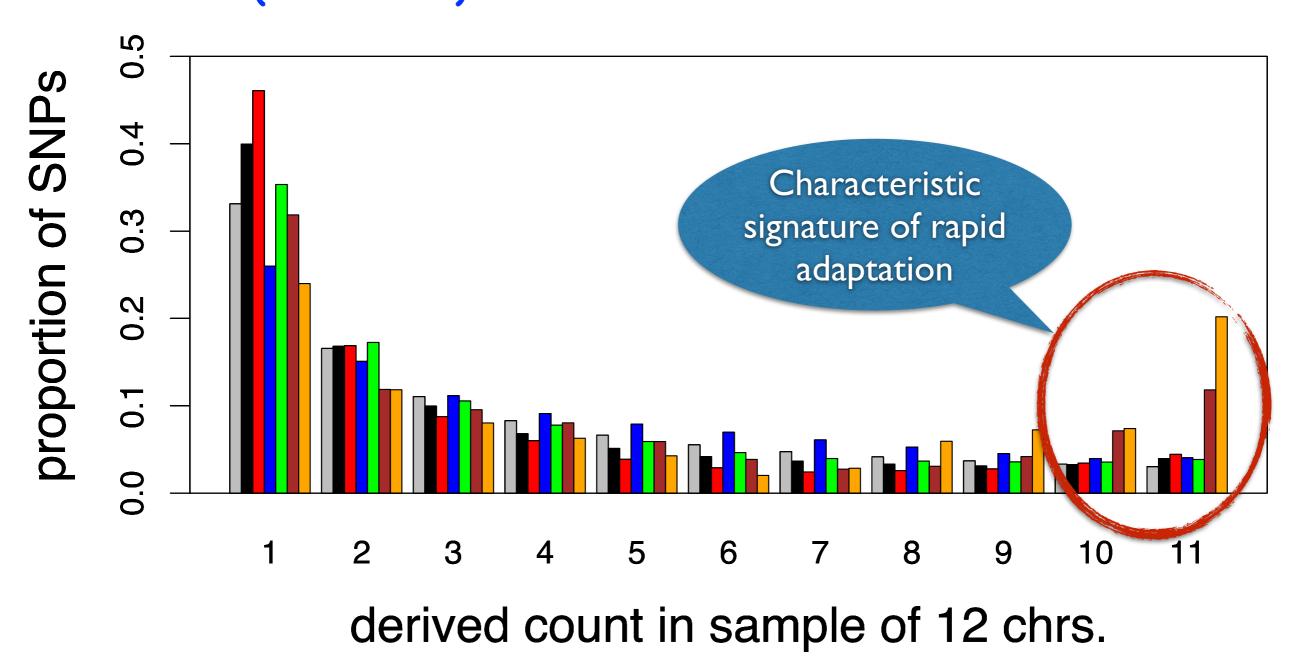


SNM AfAm (Human) Ch (RheMac) In (RheMac)



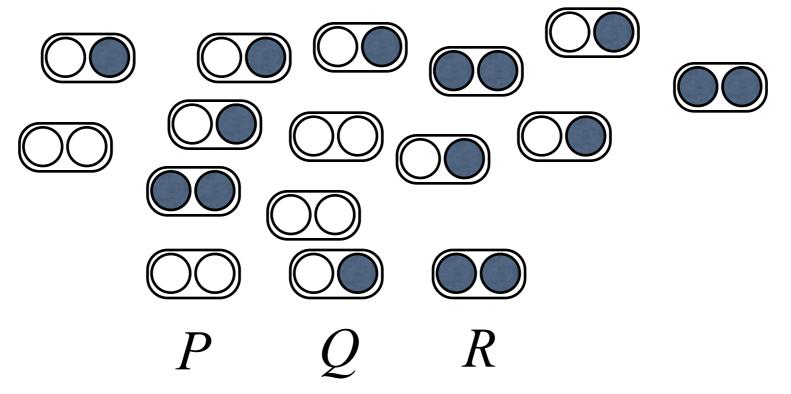
What evolutionary forces could cause an excess of high frequency derived variants? (one-word-answers-please)

SNM AfAm (Human) Ch (RheMac) In (RheMac)



Population Genetics

• Imagine a population of diploid individuals



- Principles of **random mating**:
 - Any two individuals are equally likely to mate and reproduce to populate the next generation.
 - Either chromosome is equally likely to be passed on.

Hardy-Weinberg Principle



Godfrey H. Hardy: 1877-1947



Wilhelm Weinberg: 1862-1937

What are the assumptions of the Hardy-Weinberg Principle?

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Hardy-Weinberg Principle

• Assumptions:

- Diploid organism
- Sexual reproduction
- Non-overlapping generations
- Only two alleles
- Random mating

Conclusion I: Both allele AND genotype frequencies will remain constant at HWE generation after generation... forever! 38

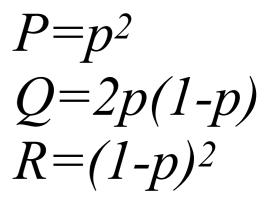




Godfrey H. Hardy: 1877-1947

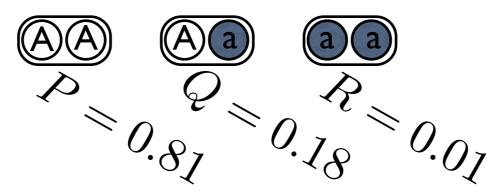
Wilhelm Weinberg: 1862-1937

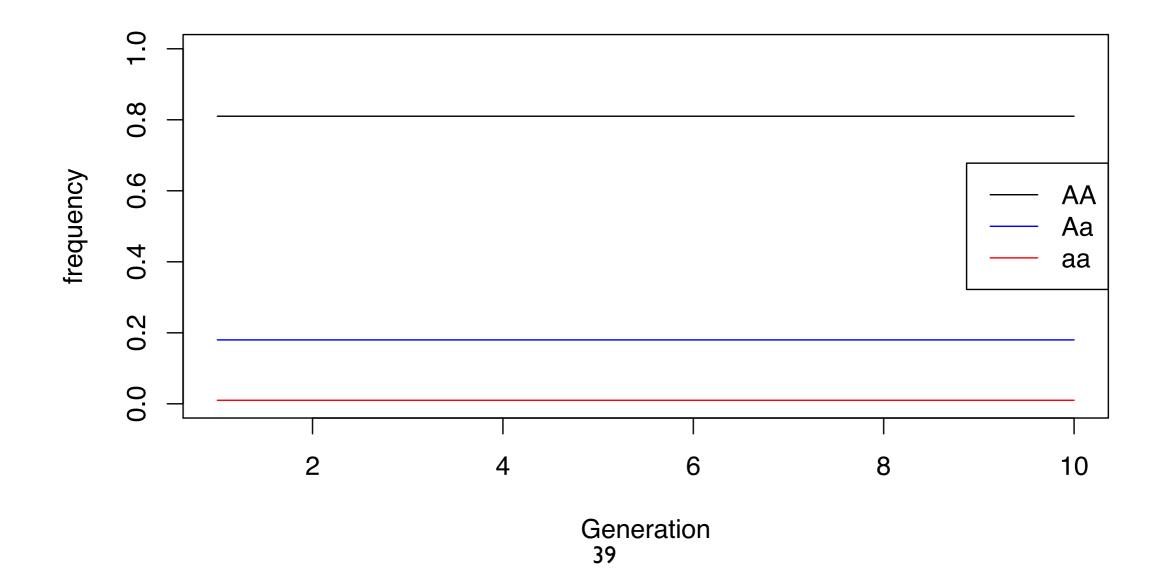
- Identical frequencies in males/females
- Infinite population size
- No migration
- No mutation
- No natural selection



Hardy-Weinberg Principle

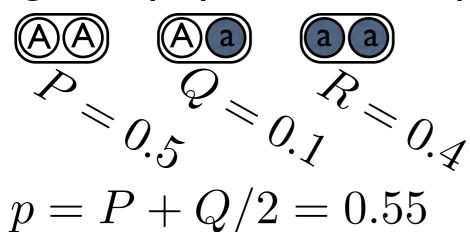
Imagine a population of diploid individuals





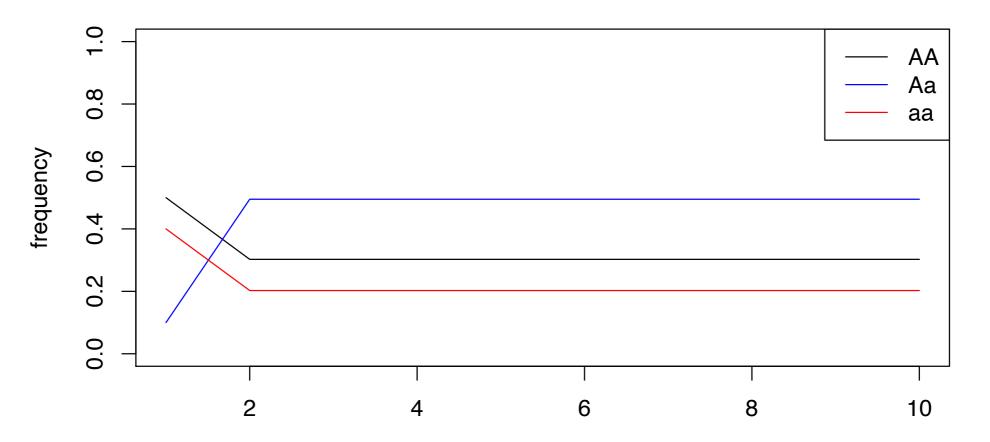
Hardy-Weinberg Principle

• Imagine a population of diploid individuals



$$p^2 = 0.3025$$

 $2p(1-p) = 0.495$
 $(1-p)^2 = 0.2025$



Conclusion 2: A single round of random mating will return the population to HWE frequencies!

Hardy-Weinberg Principle

• Assumptions:

- Diploid organism
- Sexual reproduction
- Non-overlapping generations
- Only two alleles
- Random mating



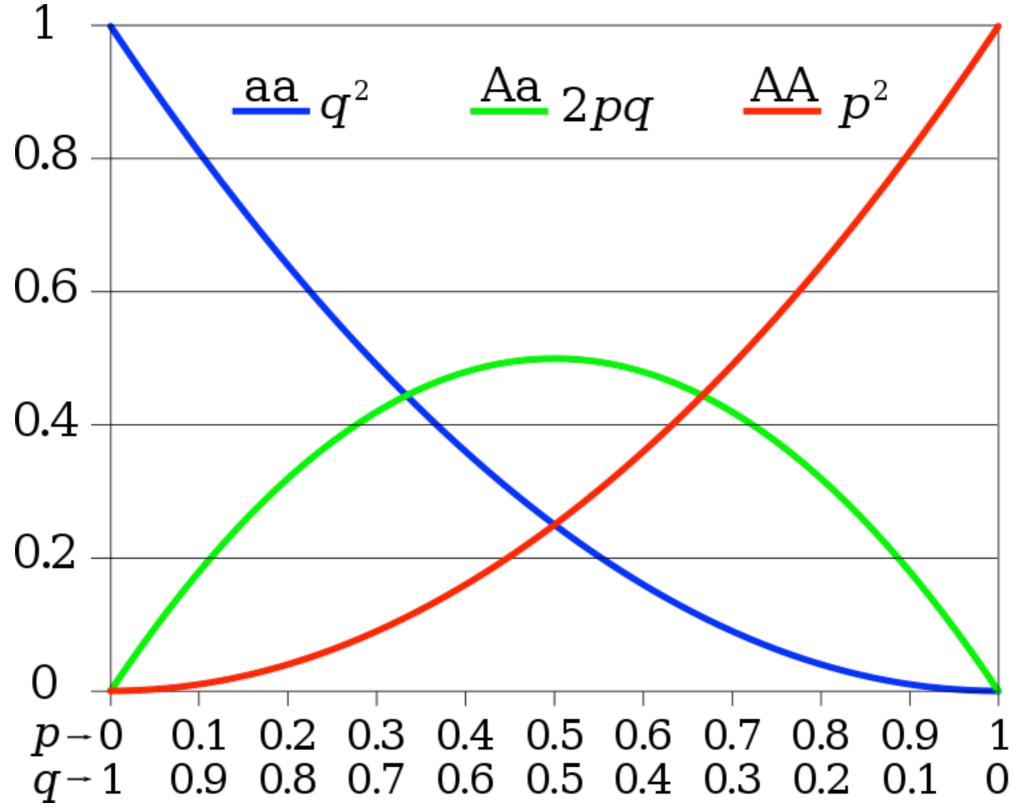


Godfrey H. Hardy: 1877-1947

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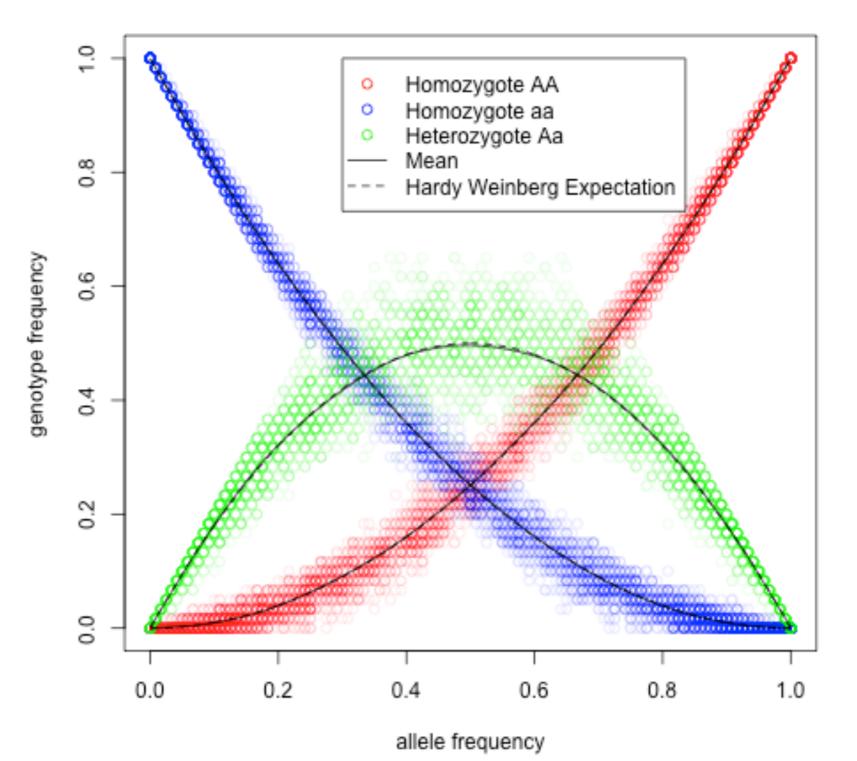
- Identical frequencies in males/females
- Infinite population size
- No migration
- No mutation
- No natural selection

Hardy-Weinberg Equilibrium



Hardy-Weinberg Equilibrium

HapMap YRI (Africans)



Graham Coop

Summary

- Hardy-Weinberg Equilibrium requires many assumptions, all of which are routinely violated in natural populations.
- Nevertheless, the vast majority of variants are in HWE.
 - Deviations almost always due to technical artifacts!
- Natural selection changes the expected allele frequency in the next generation.
 - But drift still acts in finite populations!