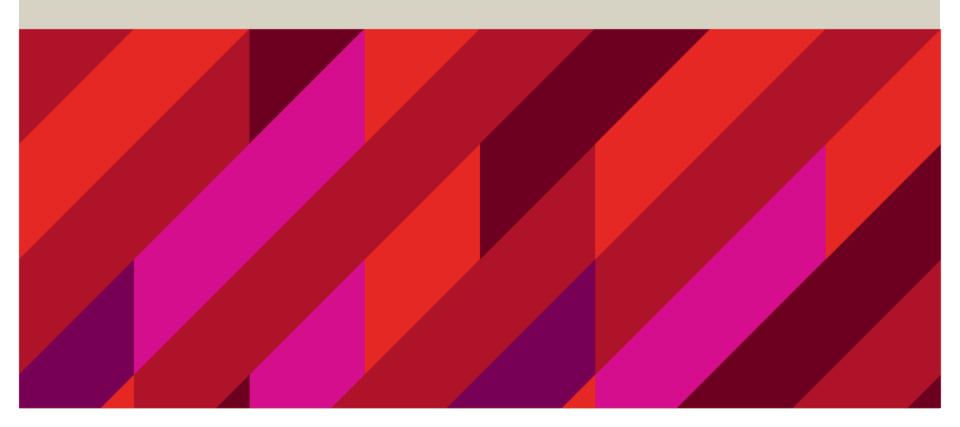


### **BIOL3110 Conservation & Ecological Genetics**

**LECTURE 19:** ADAPTIVE POTENTIAL 2



#### Case Study: Research by Jeffrey Good

**University of Montana** 

Climate change and the evolution of seasonal camouflage

#### Research to understand

- 1) the evolution of seasonal camouflage and
- 2) genomic responses to climate change in mammals

using population and functional genomic approaches

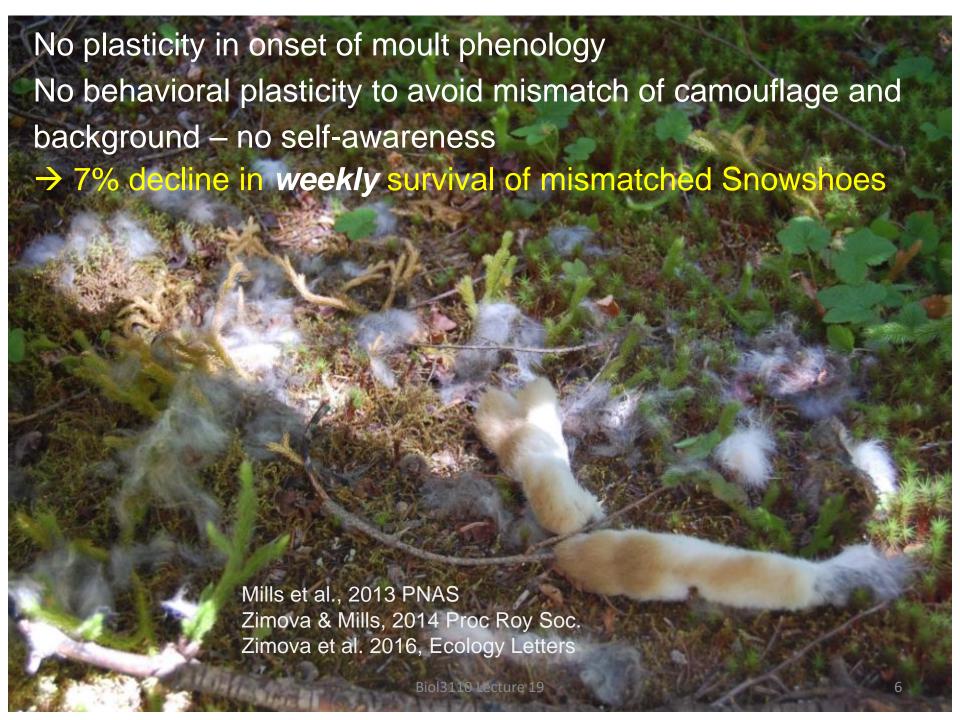


### Snowshoe hare seasonal camouflage









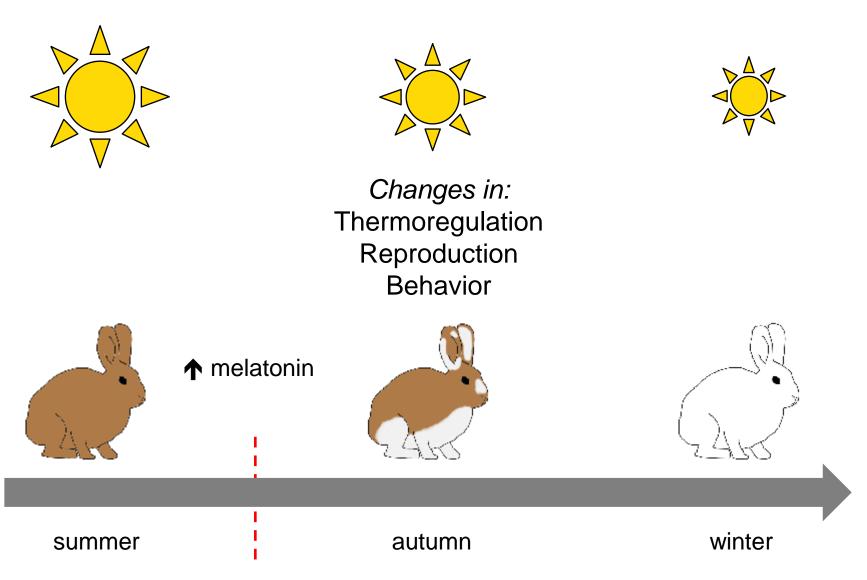
How does seasonal change in phenotype (phenology) happen?

Animals need a cue that warns them of the upcoming season, eg when there will be snow on the ground

Often use photoperiod (daylength), detected by a molecular receptor system (e.g. melatonin) that communicates with the systems that cause phenotypic seasonal change

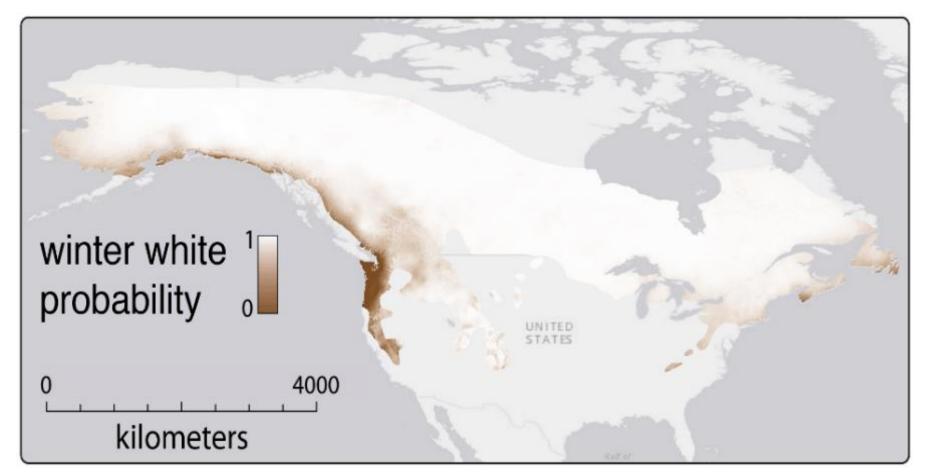
e.g. photoperiod connects to systems of circadian rhythm ('clock') genes, controlling expression of many genes, driving changes in phenotype

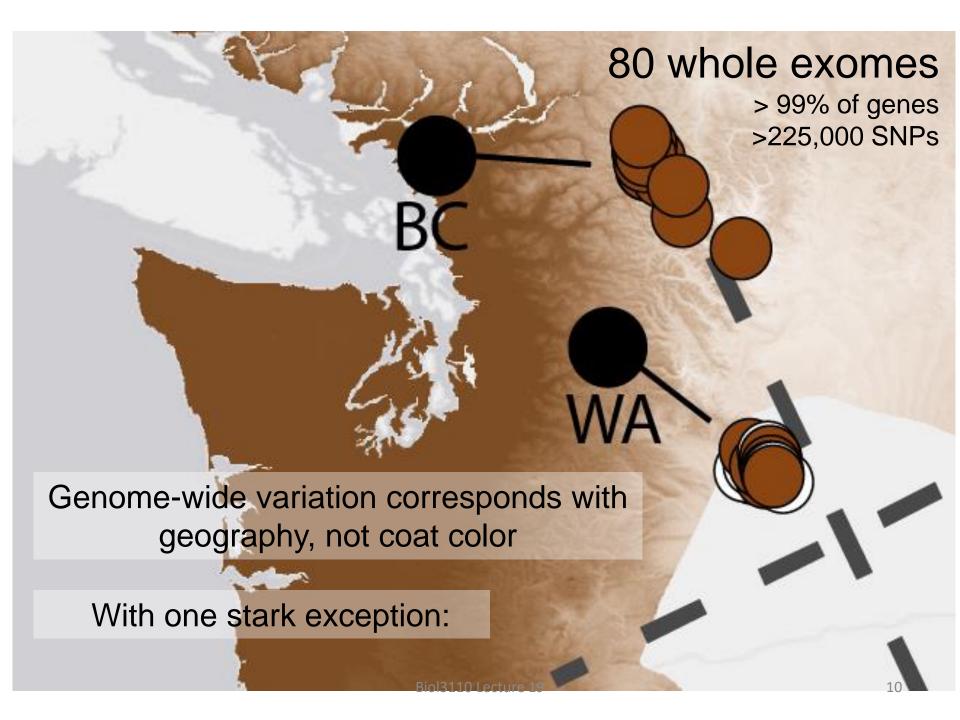
#### Coat color change is tied to seasonal signaling molecules



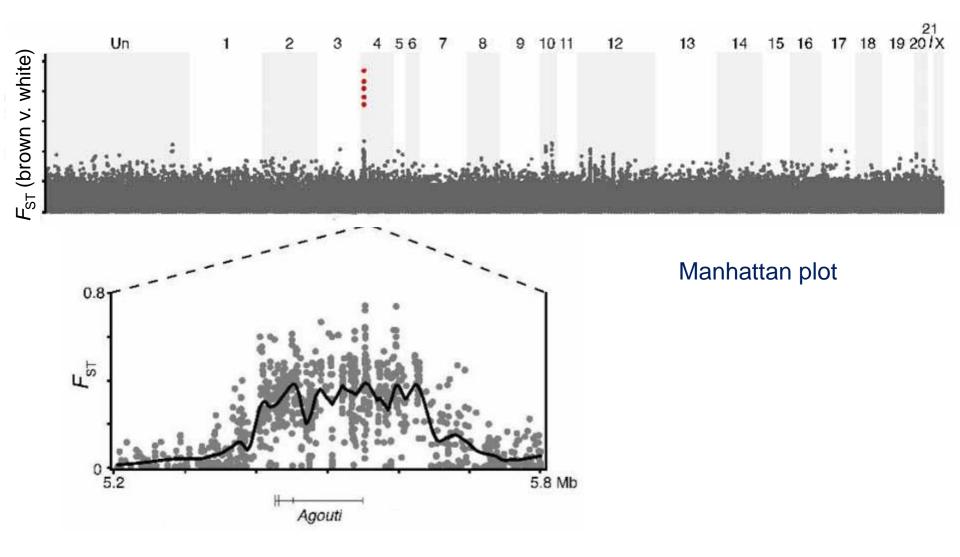
### Two components of variation:

- If change colour
- When to change colour





### Agouti is perfectly associated with winter colour



# The Agouti gene determines winter coat colour

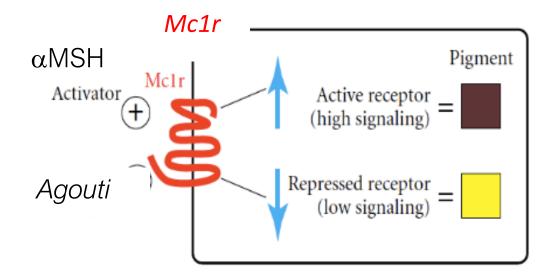


aa (a recessive)

AA Aa (A dominant)

# How does *Agouti* determine winter colour?

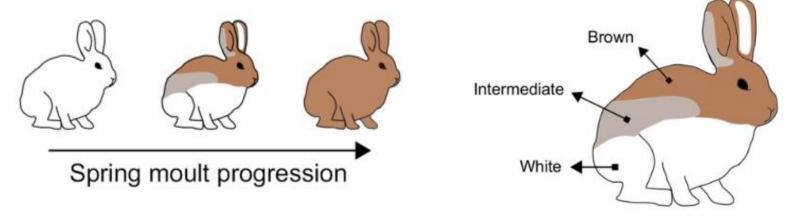
Agouti influences coat colour by reducing the binding of the melanocortin-1 receptor (MC1R), impeding the action of  $\alpha$ -melanocytestimulating hormone ( $\alpha$ MSH) that would cause production of dark eumelanin



From which you would predict that () is associated with seasonal upregulation of *Agouti – was that true?* 

# Finding genes that have different seasonal expression to do with coat colour

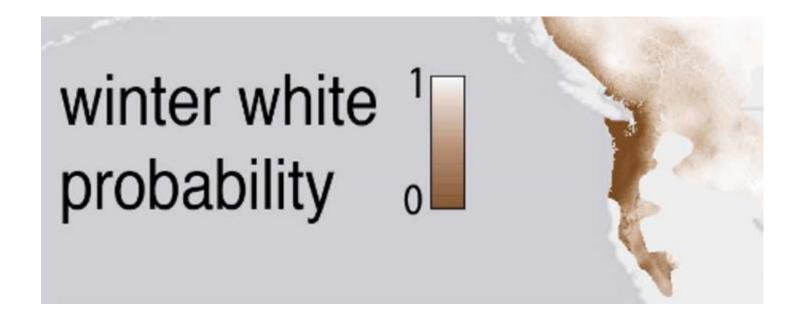
Progressive moult over the body allowed sampling of cells containing expressed genes, so that the changes in gene expression leading to differences in coat colour could be detected



Gene expression tested using RNAseq (RNA sequencing, i.e. transcriptomics) to estimate the abundance of gene transcripts in different-coloured areas of Snowshoes – result: The Agouti Signalling Peptide (ASIP) gene was upregulated in white hares.

How did the warm coastal Pacific Northwest populations evolve to never become white?

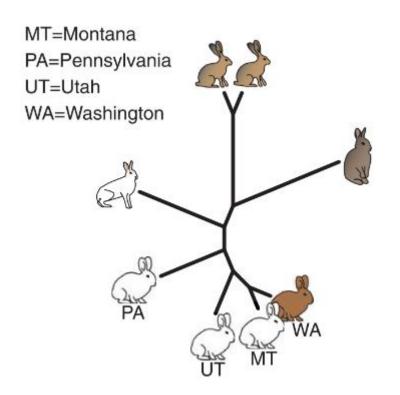
(they moult, but from summer brown to winter brown)

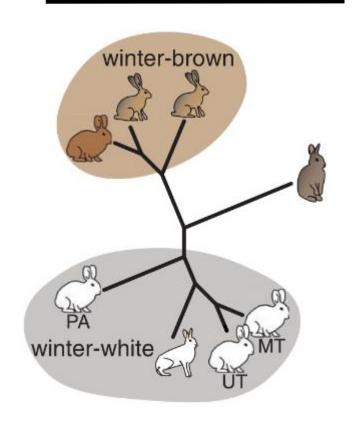


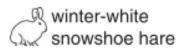
#### Species tree

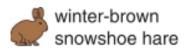
#### Agouti tree

#### Based on whole genome sequencing









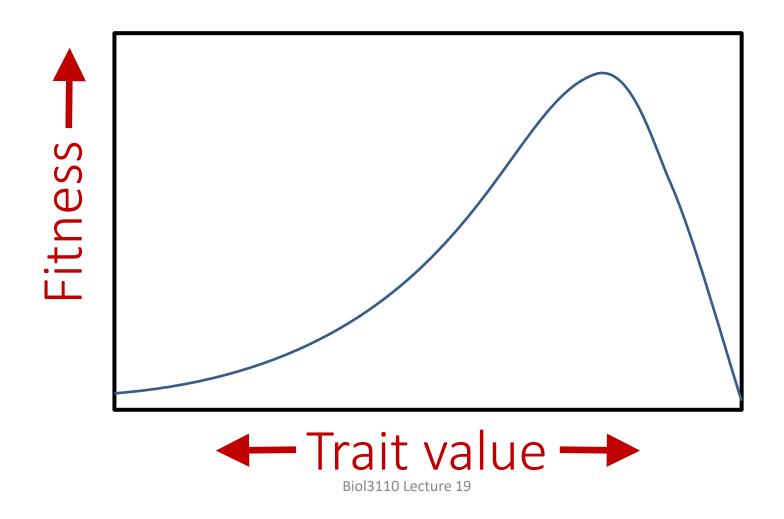




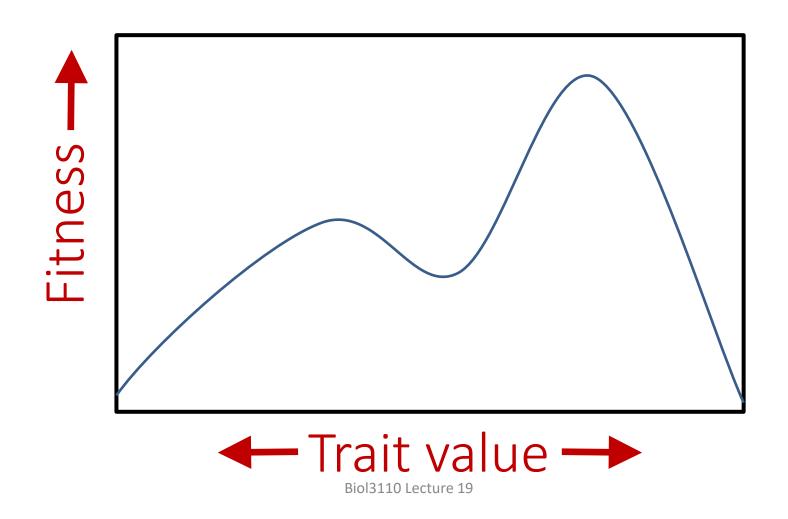


i.e. the 'brown' gene moved by hybridization from Black-tailed jackrabbit into Pacific Northwest Coast hares - adaptive introgression

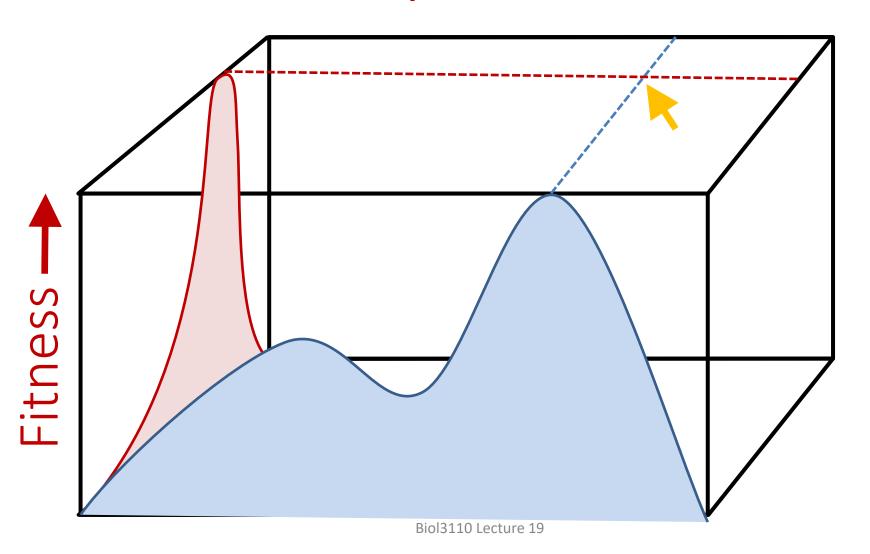
If we think of how a single trait relates to fitness:



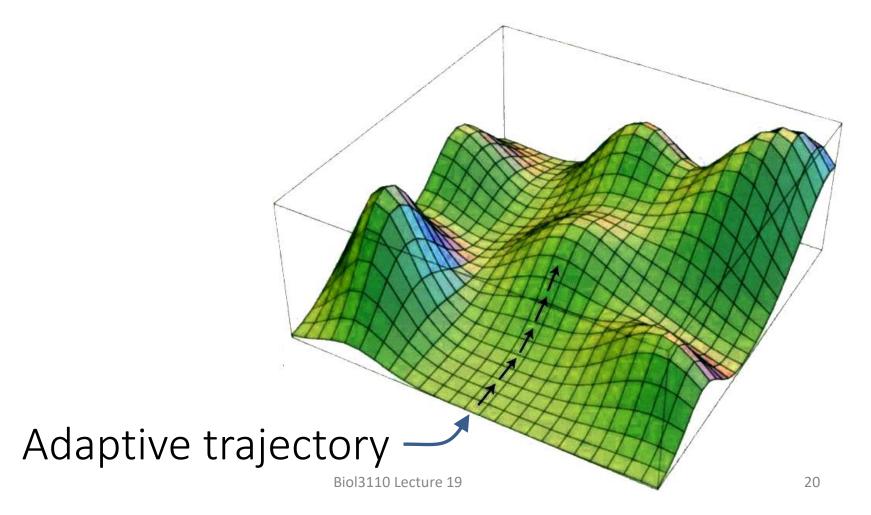
If we think of how a single trait relates to fitness:



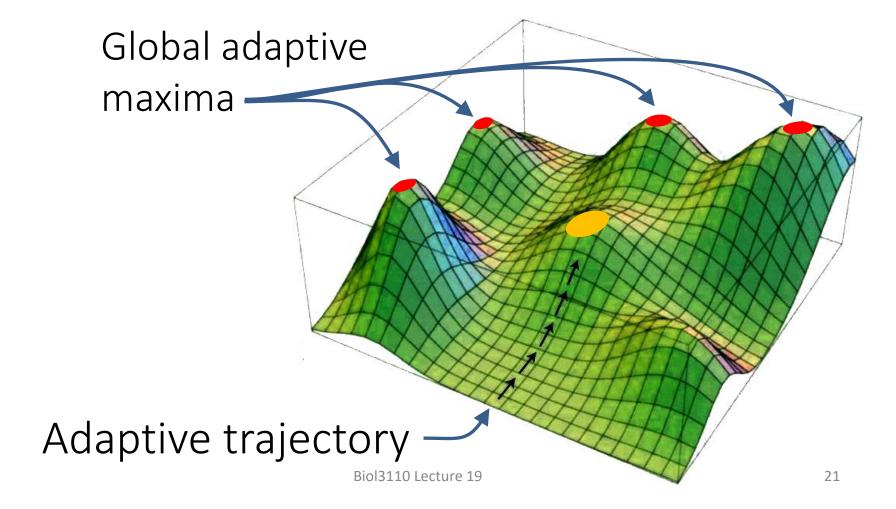
If we think of how multiple traits relate to fitness:



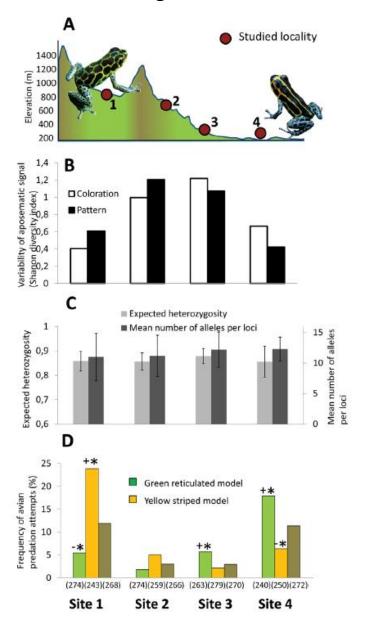
Sewall Wright: Species evolve to localised adaptive "peaks" in an n-dimensional "landscape"



Sewall Wright: Species evolve to localised adaptive "peaks" in an n-dimensional "landscape"



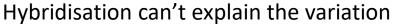
#### Choteau & Angers PloS One 2012

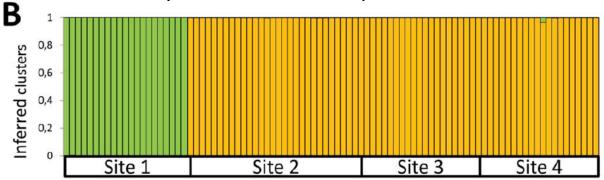


Wrights shifting balance theory – relaxation of selection + drift can allow for polymorphisms

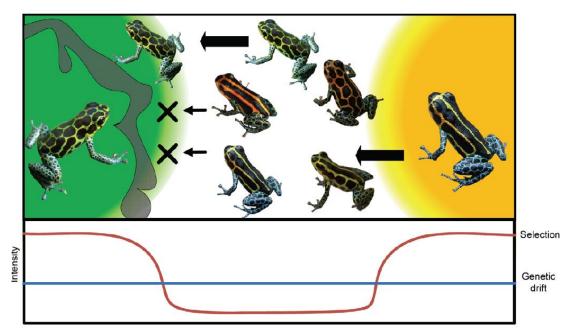
High variability sites 2 & 3

Low rates of predation sites 2 & 3





Wrights shifting balance theory – relaxation of selection + drift can allow for polymorphisms that then 'compete' across the fitness landscape



### **Next Lecture**

Social influences on connectivity