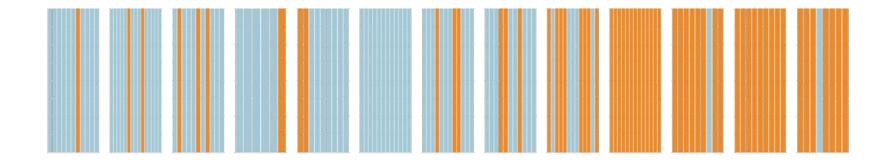


# Genetics in support of fisheries and aquaculture management

17-19 September Faro, Portugal





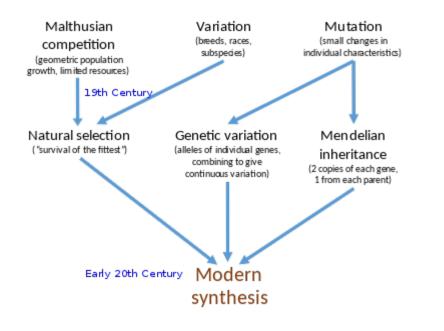
# Background principles of population genetics

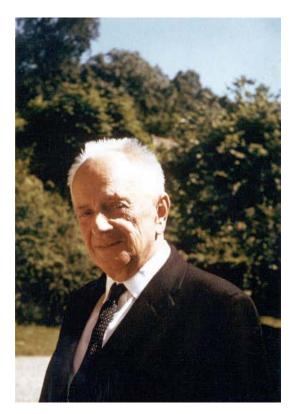
17 September Faro, Portugal



#### The importance of population genetics

- "Nothing in Biology Makes Sense Except in the Light of Evolution"
- Evolution is: "a change in the frequency of an allele within a gene pool"





Theodosius Grygorovych Dobzhansky

# "Nothing in Evolution makes sense except in the light of population genetics" (Michael Lynch)



#### Why is population genetics important?

- What are the risks of getting a child with a genetic disease?
- Has farming or breeding caused a loss in genetic variability in farmed plants and animals?
- Wild rare animals or plants be able to avoid inbreeding and survive?
- How are populations from different parts of the world related to each other?
- Has genomes adapted to different environmental conditions across the globe?
- How do populations and species evolve over time?



# What is a (genetic) population?

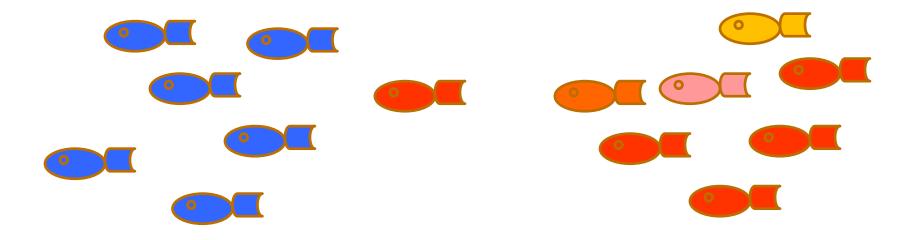
- The individuals of a species living within an country?
- The individuals of a certain species living within a restricted geographical region, e.g. the Scandinavian peninsula?
- Group of individuals within a species who are reproductively isolated (to some degree) from other individuals of the same species?

# **Concepts and definitions**



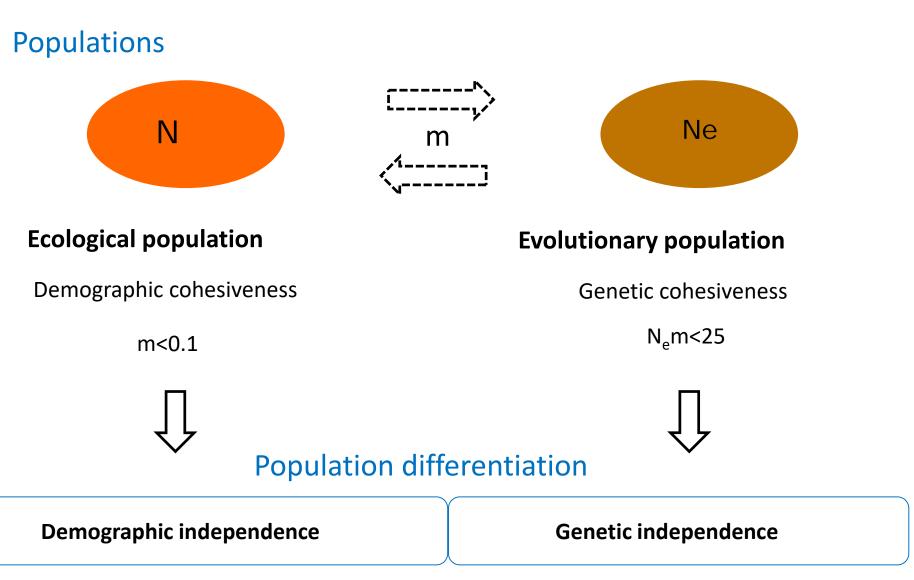
#### **Populations**

A group of individuals of the same species that are reproductively isolated (at least to some extent) from other conspecifics



## **Concepts and definitions**

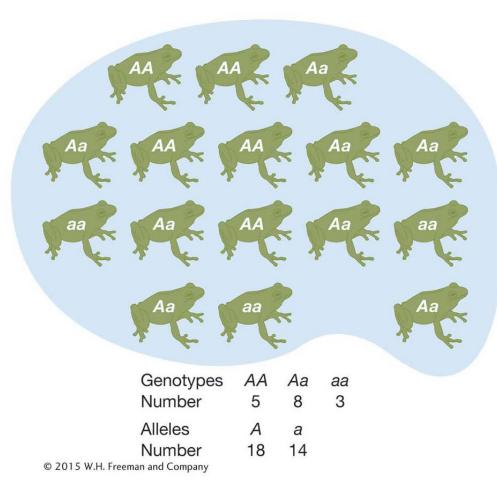






# EVOLVE HAPPY HOUR IN THE GENE POOL

#### **The Gene-Pool Concept**



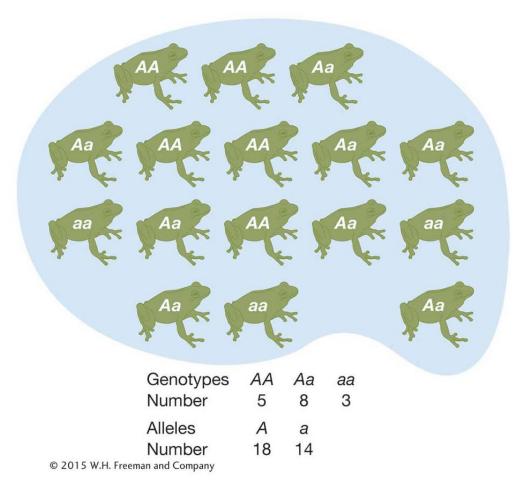
# The gene pool is the sum total of alleles in a population

The gene pool is a fundamental concept for the study of genetic variation in populations: it is the sum total of all alleles in the breeding members of a population at a given time. We can describe the variation in a population in terms of genotype and allele frequencies



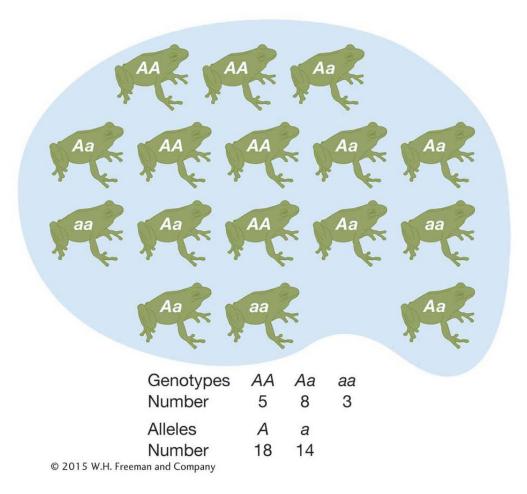
p=? q=?

#### The Hardy Weinberg law





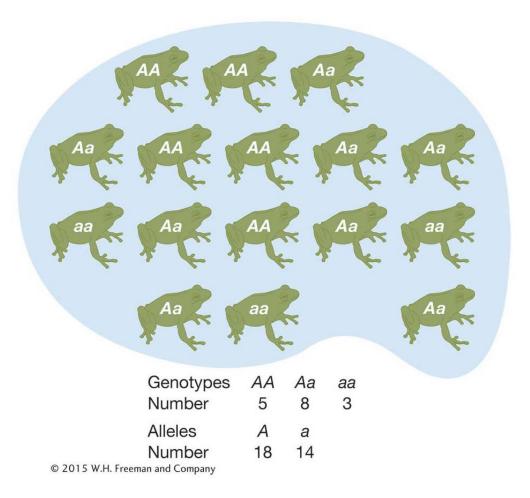
#### The Hardy Weinberg law



p= 0,56 q=0.44

What is the probability of getting an AA homozygote in the next generation?

#### The Hardy Weinberg law



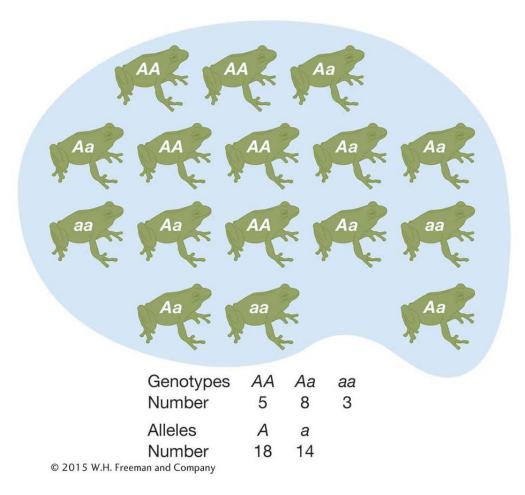
What is the probability of getting an AA homozygote in the next generation?

$$pxp = p^2 = 0.3136$$

aa homozygote:  $qxq = q^2 = 0.1936$ 



#### The Hardy Weinberg law



What is the probability of getting an Aa heterozygote in the next generation?

(pxq)x2 = 2pq = 0.4928

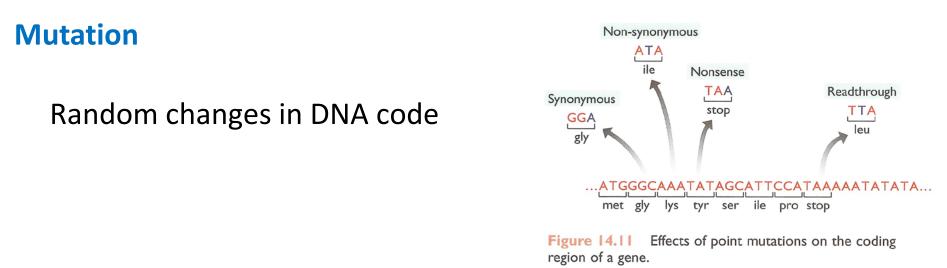


# Factors affecting genetic variation in populations

- Four evolutionary forces:
- Mutation
- Migration
- Random genetic drift
- Selection



"Why Dug! I haven't seen you in years! You haven't evolved a bit!

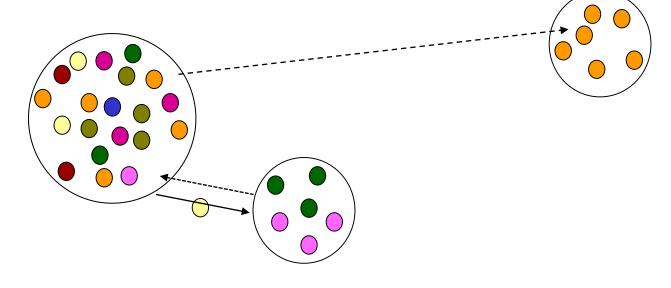


- Most mutations are harmful -> removed through natural selection
- A few mutations are beneficial -> favoured and increase in frequency in the population
- Mutation rates are generally very low, but varies among species and genomic regions



#### **Gene flow**

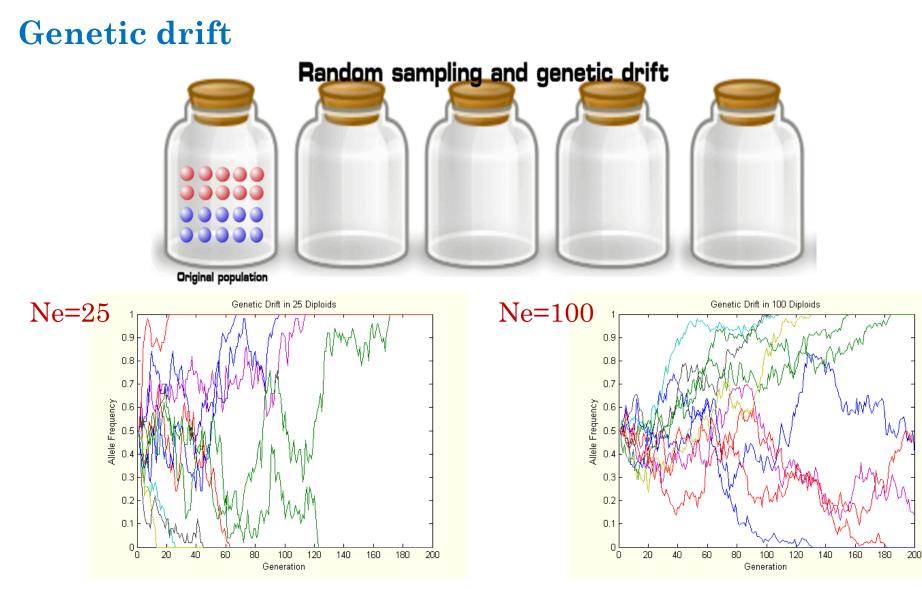
#### Movement of alleles between populations



#### Is gene flow = migration??

#### No!

Gene flow requires successful reproduction in new population



18 DTU Aqua, Technical University of Denmark

# Effective population size



The **effective population size (Ne)** is the size of an ideal population, which loses genetic variation at the same rate as the actual population

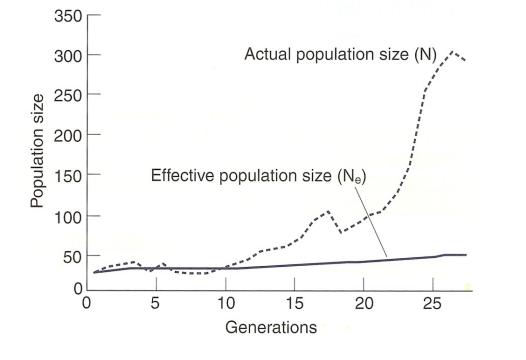
= "how does the population behave genetically?"

So, what is an ideal population?

- 1) Random mating
- 2) 1:1 sex ratio
- 3) Non-overlapping generations
- 4) Random number of offspring (Poisson distribution)

# Effective population size

#### Effective population size (Ne) and census size (N)



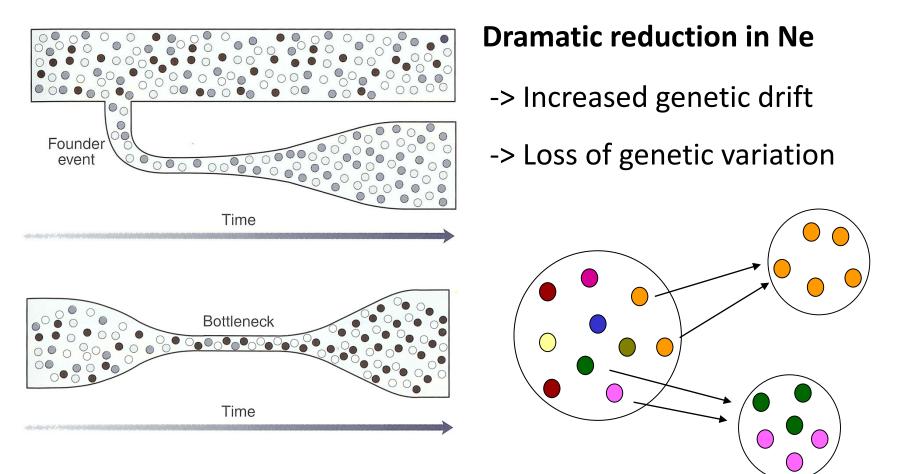
- Ne is often smaller than N
- N<sub>e</sub> is disproportionately affected by the smaller population sizes
- In a recently expanded population, N<sub>e</sub> (and the amount of genetic variation) is still largely determined by the smaller ancestral population size.



# Effective population size

# DTU

#### Founder events and bottlenecks





#### Selection (natural and artificial)

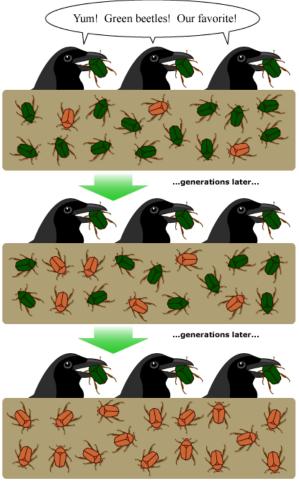
Selection requires

1) Genetic variation between individuals

2) Inheritance from parents to offspring

3) Fitness differences between genetic variants

#### Natural selection, in a nutshell:

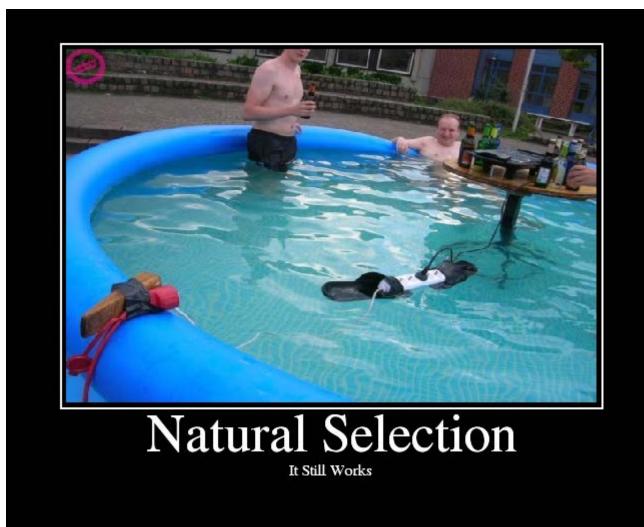


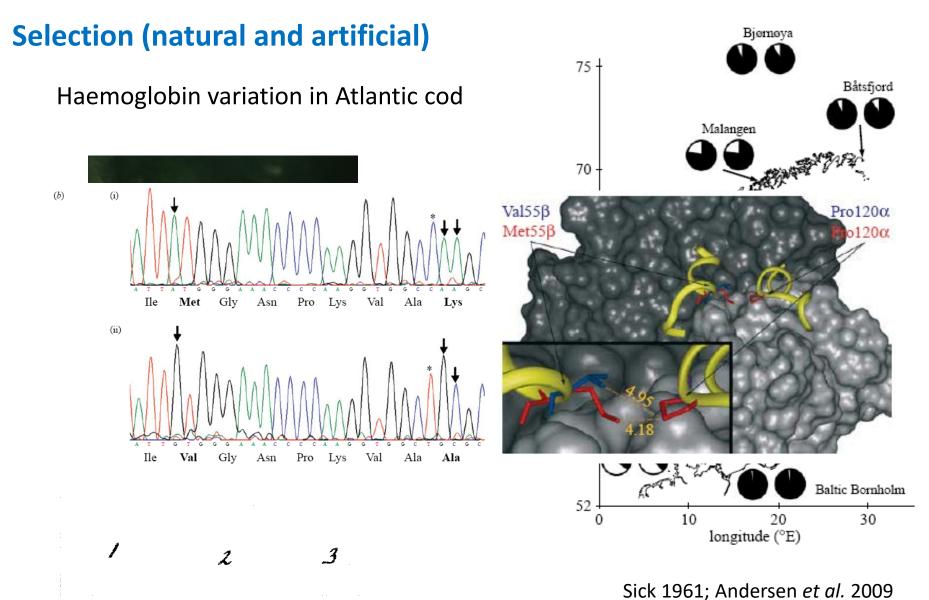
Green beetles have been selected against, and brown beetles have flourished.

http://evolution.berkeley.edu/evolibrary/arti cle/bergstrom\_02



#### **Selection (natural and artificial)**

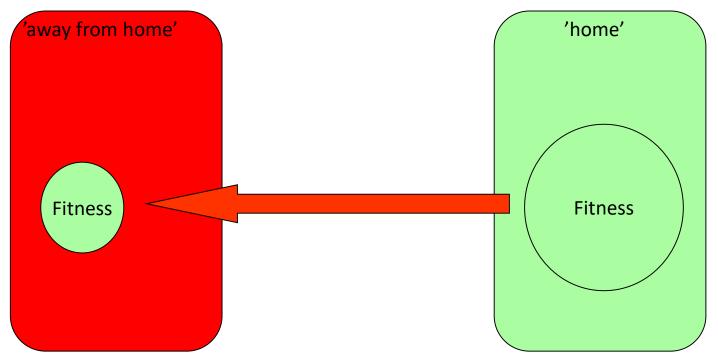




#### **Selection (natural and artificial)**

#### Local adaptation - 'Home' versus 'away'

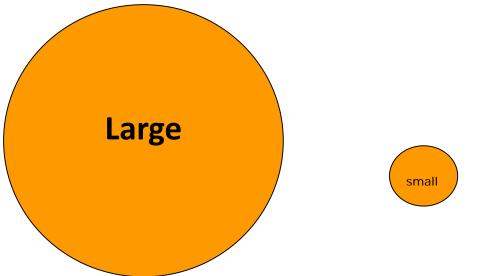
 Individual has higher fitness 'at home' compared to in a different environment





Balance between genetic drift and selection

Where is genetic drift and selection most important?



- Selection operates most efficiently in large populations
- In small populations, even disadvantageous alleles may become fixed due to **genetic drift**

# Inbreeding



- **Definition**: Reproduction between (genetic) relatives
- **Result**: Increased homozygosity, reduced heterozygosity
  - difference between expected (He) and observed (Ho) heterozygosity
- Effect: May lead to inbreeding depression through exposure of deleterious recessive alleles to selection

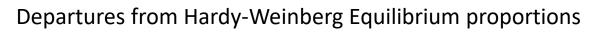
Relevance to wild populations?

-> Poor growth and survival, deformities, reduced fertility

-> Largest effect in small populations

# Inbreeding





3	Genotype	Random mating	Inbreeding (=non-random mating)
	AA	p <sup>2</sup>	p² + Fpq
	Aa	2pq	2pq - F2pq
	aa	q <sup>2</sup>	q <sup>2</sup> + Fpq

**Inbreeding coefficient** Fis = (Hs - Hi) / Hs - the mean reduction in H of a locus within an individual due to non-random mating within a subpopulation

E

 $\frac{1}{F_A}$ 

1/2

<u>1</u>2

 $A_1A_2$ 

붛

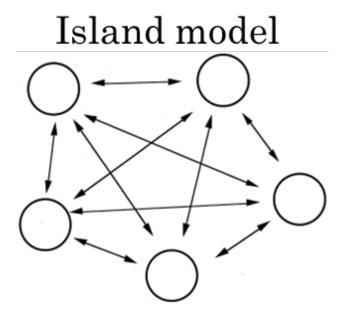
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## Genetic variation and population structure



#### Wright's F statistics

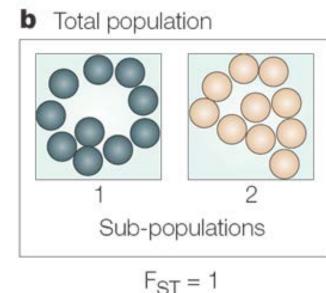
- Inbreeding/co-ancestry coefficients
- Measures how the genetic variation is distributed between
  - Individuals (I)
  - Subpopulations (S)
  - Total population (T)





#### Wright's F statistics

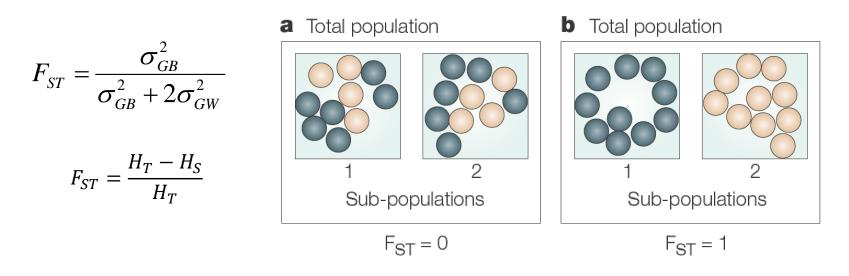
- Inbreeding/co-ancestry coefficients
- Measures the reduction in heterozygosity (H) due to random genetic drift



## Genetic variation and population structure

#### Wright's F statistics

# Proportion of total genetic variance caused by differences among populations



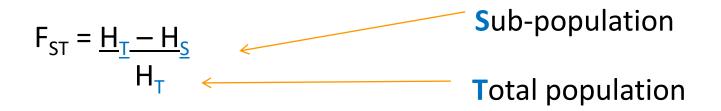
**F**<sub>ST</sub> = **1**: Populations fixed for different alleles

**F**<sub>sT</sub> = **0**: Allele frequencies are the same across populations



#### Genetic variation and population structure





Two alleles, two populations:

	$A_1$	$A_2$	$H_{\tau} = 2*0.5*0.5 = 0.5$
Freq. in pop 1	0.3	0.7	$H_{s} = ((2*0.3*0.7) + (2*0.7*0.3))/2$
Freq. in pop 2	0.7	0.3	= 0.42
Overall freq	0.5	0.5	

 $F_{ST} = (0.5-0.42)/0.5 = 0.16$ 

