

Management strategies for inbreeding control in unselected and selected populations

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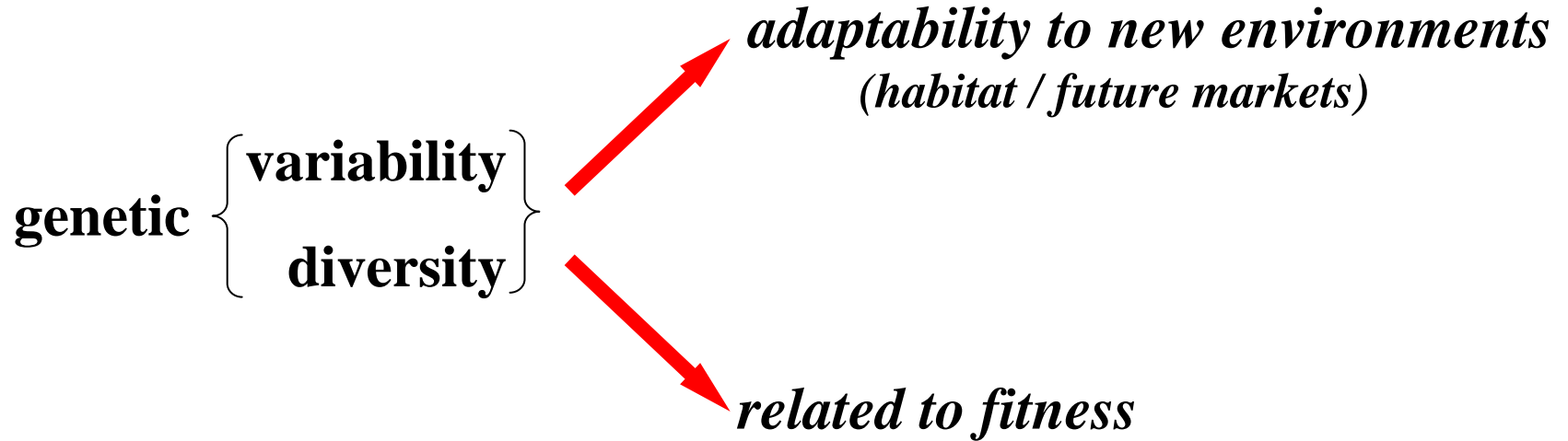
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Preserve

~~NATURAL RESERVE~~



Gary Larson



short-term: inbreeding depression

*long-term: accumulation of deleterious mutations
loss of evolutionary potential*

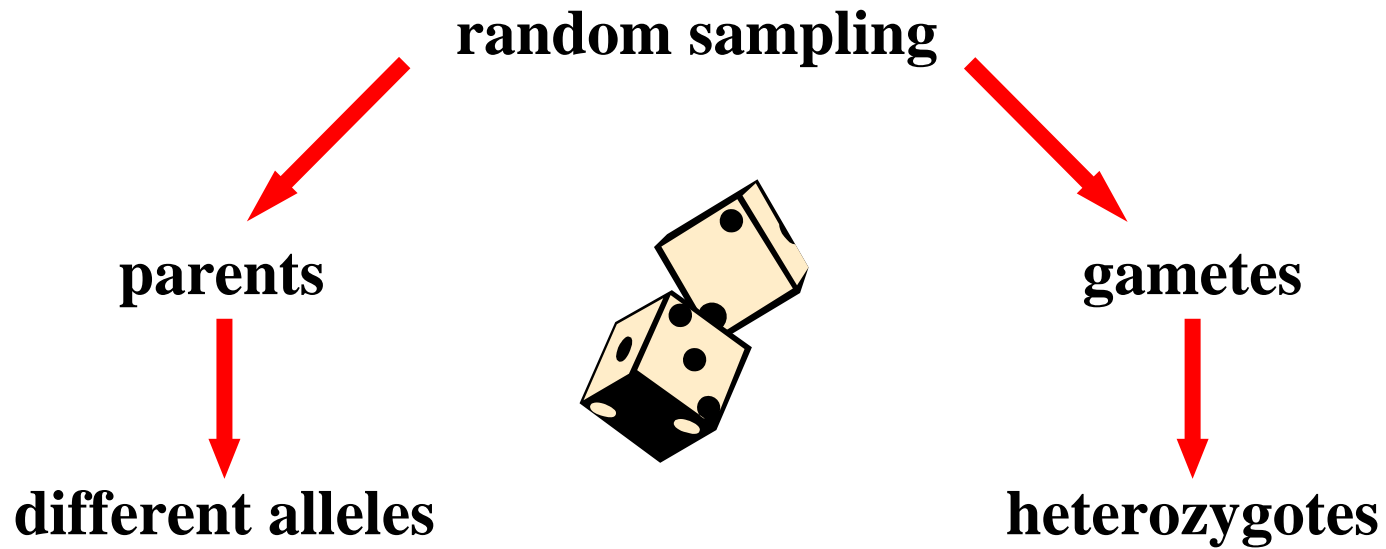


loss of variability

selection \Rightarrow *loss of detrimental alleles*

drift \Rightarrow *random sampling*





Genetic drift

**Variation of allelic
frequencies**

$$\propto 1/N$$

**Increase of
inbreeding (F)**

Effective population size (N_e)

\Rightarrow ideal population with the same $\begin{cases} \Delta F \\ \Delta V(q) \end{cases}$

fluctuating
population size

$$\frac{1}{N_e} \approx \frac{1}{t} \sum_{i=1}^n \frac{1}{N_i}$$

biased sex ratio

$$\frac{1}{N_e} = \frac{1}{4N_f} + \frac{1}{4N_m}$$

differential
contributions

$$N_e \approx \frac{4N}{2 + S_k^2}$$

Recommendations:

keep constant census
1:1 sex ratios
equalise contributions

ESTIMATION OF N_e

✓ demographic data

✓ pedigree analysis

$$\Delta F = \frac{1}{2N_e}$$

✓ molecular data

- *fluctuation of allelic frequencies*
- *decrease in heterozygosity*
- *amount of linkage disequilibrium*

Genetic management of a population

➤ two decisions to take:

✓ *which individuals reproduce?*

1

✓ *how they mate?*

SELECTION

CONTRIBUTIONS

➤ maximise N_e

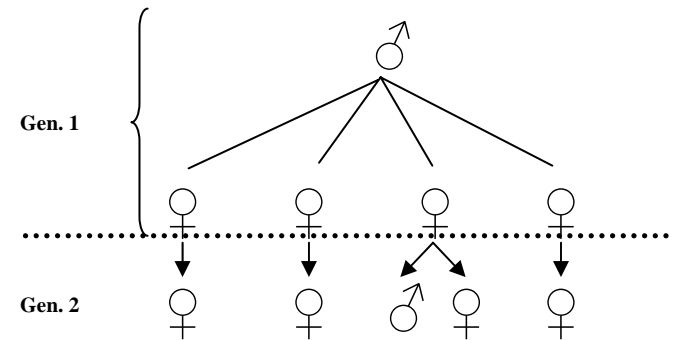
keep constant census

1:1 sex ratios

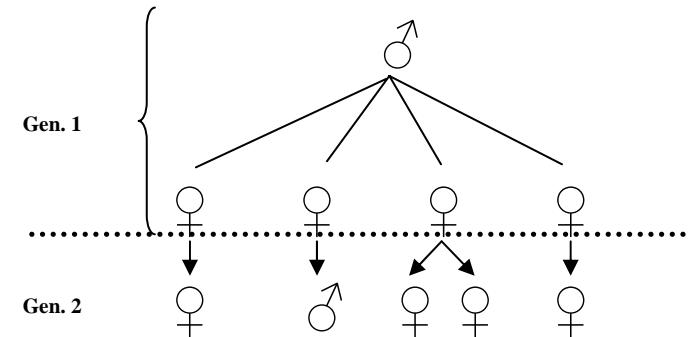
equalise contributions

Hierarchical
regular
methods

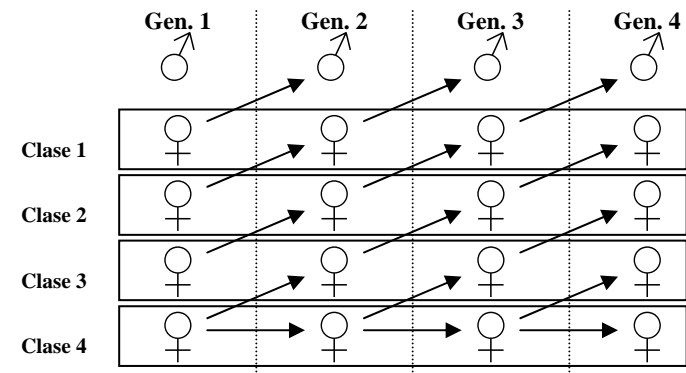
Gowe et al. (1959)



Wang (1997)



Sánchez-Rodríguez et al. (2003)



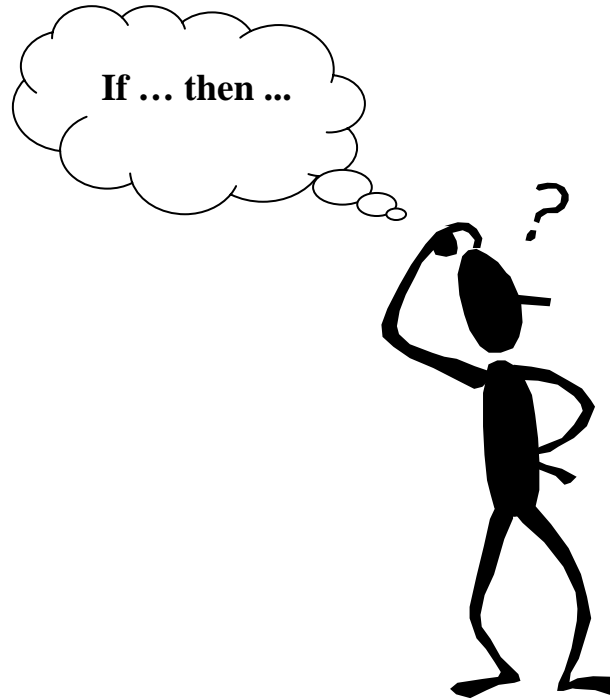
close relatives



**share genetic
information**



low diversity



**mating between
relatives**



**increase of
inbreeding**



**inbreeding
depression**



Let's minimise coancestry !!!

f_{ij} = coancestry coefficient

*... probability of two individuals carrying alleles
identical by descent in a random locus ...*

Malecot (1948)

⇒ from pedigrees or from markers

**OPTIMAL
CONTRIBUTIONS**

- overrepresented individuals are penalised
- loosely related individuals are favoured

Ballou & Lacy (1995)

$$\min \sum_{i=1}^N \sum_{j=1}^N c_i c_j f_{ij}$$

- ✓ equalises ancestral contributions
- ✓ maximises Ne
- ✓ maximises gene diversity (Exp. Het.) $\bar{f} = 1 - GD$
- ✓ minimises ΔF $\Delta F = \frac{1}{2Ne}$
- ✓ flexible and robust

Animal Breeding \Rightarrow selection

\Rightarrow *improving a particular trait*

$$\Delta G = i \rho_{AC} \sigma_A$$

$$\uparrow i \Rightarrow \uparrow \Delta G, \uparrow \Delta F$$

$$\uparrow \rho \Rightarrow \uparrow \Delta G, \uparrow \Delta F$$

**Response
/ Gain**



**Inbreeding /
Genetic diversity**

➤ **Decrease importance of relatives' information**

✓ **Inflated heritability**

✓ **Suboptimal familiar indices**

➤ **Allow for differential contributions**

✓ **proportional to breeding value**

✓ **more selected \Rightarrow same i with more Ne**

OPTIMAL CONTRIBUTIONS

(Wray & Goddard 1994, Meuwissen 1997)

$$\max \left(\sum_{i=1}^N c_i EBV_i \right)$$

contributions proportional to breeding value ...

$$s.t. \left(\sum_{i=1}^N \sum_{j=1}^N c_i c_j f_{ij} \right) \leq F_{t+1}$$

... but also to average relationship

Genetic management of a population

- two decisions to take:
 - ✓ *which individuals reproduce?*
 - ✓ *how they mate?* 2

- less important than selection
 - ⇒ *little margin for improvement*

➤ **Factorial mating**

- ✓ several partners per individual
- ✓ HS families instead of FS families


➤ **Compensatory mating**

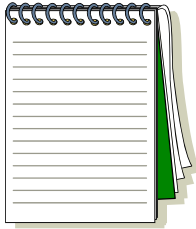
- ✓ mix overrepresented lineages with rare ones

➤ **Minimum coancestry mating**

- ✓ avoid mating between close relatives
- ✓ delays inbreeding (but not ΔF)

CRYOCONSERVATION

- ✓ **use of post-reproductive individuals**
- ✓ **increases census**
- ✓ **increases generation interval** 
- ✓ **reduces drift**



TO TAKE HOME

✓ **N_e is a key parameter**

⇒ management and monitoring

✓ **OC controls the rise of inbreeding**

⇒ with and without selection

⇒ also reduces loss of diversity

✓ **mating less important than selection**

⇒ but *mcm* could be advisable