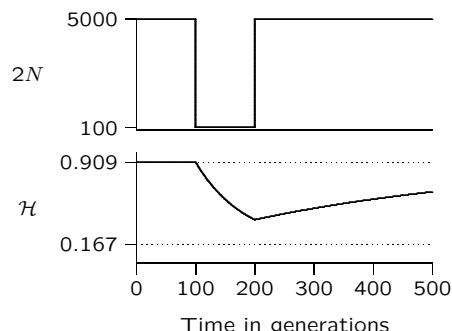


LECTURE: Drift in Populations that Vary in Size

- Heterozygosity is often surprisingly small. Why?
- Urn model assumes N is constant. What if it varies?
- Bottleneck: a temporary reduction in N
- Decline in \mathcal{H} is faster than recovery.
- Effective population size is harmonic mean of N_t

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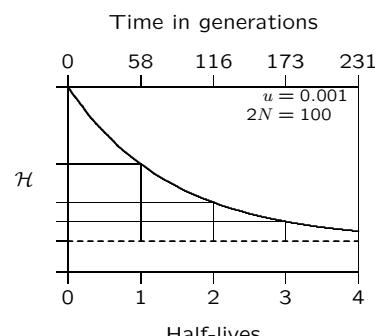
A bottleneck in population size



\mathcal{H} declines rapidly, recovers slowly. Why?

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What is a half-life?



3

Why the decline is faster than the recovery

Gene diversity converges toward its equilibrium with a half-life of

$$t_h = \frac{\ln 2}{2u + 1/2N}$$

Female population size	Half-life of convergence (generations)	(years)
∞	347	1,041
10^6	346	1,038
10^5	345	1,035
10^4	330	990
10^3	231	693
10^2	58	174
10	7	21

(Assumes $u = 0.001$)

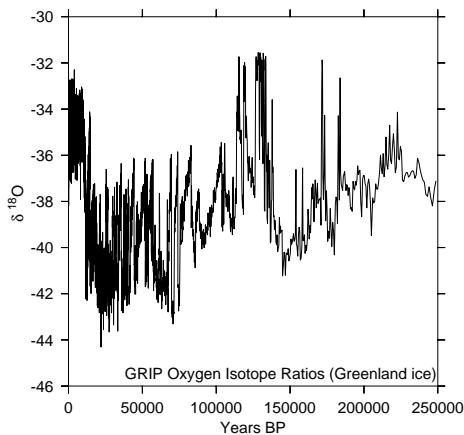
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Another problem

We have been assuming that the Elk population was stationary prior to being depleted by human hunting. Is this a reasonable assumption?

Have a look at the record of climate change during the past 250,000 years.

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These drastic fluctuations in temperature must have caused fluctuations in population size.

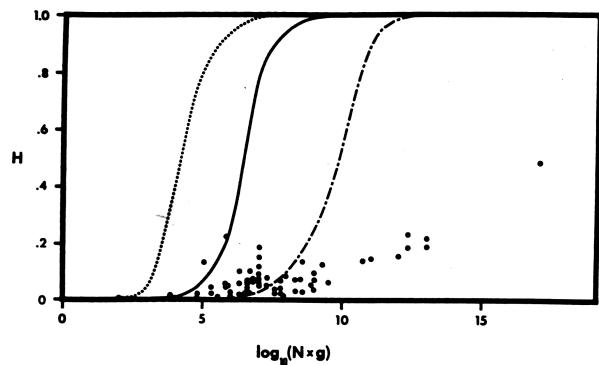


FIG. 1. Relationship between average gene diversity H and Ng . Dots represent observed values. The number of dots is less than 72, because some observed values overlap. Solid line: expected relationship for neutral alleles with a mutation rate of $v = 10^{-7}$. Dotted line: expected relationship for overdominant alleles with $v = 10^{-7}$ and $s = 0.001$. Dot-dashed line: expected relationship for slightly deleterious alleles with $v = 10^{-7}$ and $s = 0.002$. This relationship was obtained by using Kimura's (1979) formulas (23) and (24) with $\beta = 0.5$.

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For many species, H is much smaller than would be expected on the basis of their population sizes. Could this be a result of population size bottlenecks during the Pleistocene?

Effective population size, N_e

Goal: Find a value of N that makes our idealized population behave like a more complicated one.

Example: When population size varies, what value of N makes the gene diversity equal to

$$H = \frac{4N_e u}{4N_e u + 1} ?$$

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Review: H in a population of constant size

$$H_1 = H_0 \left(1 - \frac{1}{2N}\right)$$

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Another generation

$$\begin{aligned}\mathcal{H}_1 &= \mathcal{H}_0 \left(1 - \frac{1}{2N}\right) \\ \mathcal{H}_2 &= \mathcal{H}_1 \left(1 - \frac{1}{2N}\right)\end{aligned}$$

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$$\begin{aligned}\mathcal{H}_1 &= \mathcal{H}_0 \left(1 - \frac{1}{2N}\right) \\ \mathcal{H}_2 &= \mathcal{H}_1 \left(1 - \frac{1}{2N}\right) \\ &= \mathcal{H}_0 \left(1 - \frac{1}{2N}\right)^2\end{aligned}$$

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General form

$$\begin{aligned}\mathcal{H}_t &= \mathcal{H}_0 \left(1 - \frac{1}{2N}\right)^t \\ &\approx \mathcal{H}_0 \exp[-t/2N]\end{aligned}$$

Note approximation: $1 - 1/2N \approx e^{-1/2N}$ when N is large.

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 \mathcal{H} in a population of varying size

$$\mathcal{H}_1 = \mathcal{H}_0 \left(1 - \frac{1}{2N_0}\right)$$

where N_0 is population size in generation 0.

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Another generation

$$\begin{aligned}\mathcal{H}_1 &= \mathcal{H}_0 \left(1 - \frac{1}{2N_0}\right) \\ \mathcal{H}_2 &= \mathcal{H}_1 \left(1 - \frac{1}{2N_1}\right)\end{aligned}$$

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Two generations

$$\mathcal{H}_2 = \mathcal{H}_0 \left(1 - \frac{1}{2N_0}\right) \left(1 - \frac{1}{2N_1}\right)$$

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Two generations again

$$\begin{aligned}\mathcal{H}_2 &= \mathcal{H}_0 \left(1 - \frac{1}{2N_0}\right) \left(1 - \frac{1}{2N_1}\right) \\ &= \mathcal{H}_0 \prod_{i=0}^1 \left(1 - \frac{1}{2N_i}\right)\end{aligned}$$

where \prod is the product operator.

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General form

$$\mathcal{H}_t = \mathcal{H}_0 \prod_{i=0}^t \left(1 - \frac{1}{2N_i}\right)$$

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Approximation

$$\begin{aligned}\mathcal{H}_t &= \mathcal{H}_0 \prod_{i=0}^t \left(1 - \frac{1}{2N_i}\right) \\ &\approx \mathcal{H}_0 \prod_{i=0}^{t-1} \exp[-1/2N_i]\end{aligned}$$

How do you simplify a product of exponentials?

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Remember rule for product of exponentials

$$e^a e^b = \exp[a + b]$$

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Product of exponentials is exponential of sum

$$\begin{aligned}\mathcal{H}_t &\approx \mathcal{H}_0 \prod_{i=0}^{t-1} \exp[-1/2N_i] \\ &= \mathcal{H}_0 \exp \left[- \sum_{i=0}^{t-1} \frac{1}{2N_i} \right]\end{aligned}$$

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Compare results for fixed and varying N

$$\begin{array}{lll} \text{Fixed } N & & \text{Varying } N \\ \mathcal{H}_t \approx \mathcal{H}_0 \exp[-t/2N_e] & = & \mathcal{H}_0 \exp \left[- \sum_{i=0}^{t-1} \frac{1}{2N_i} \right] \end{array}$$

- N_e is called effective population size.
- It is the constant population size that makes the two sides equal.

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The two sides are equal when

$$1/N_e = \frac{1}{t} \sum_{i=0}^{t-1} \frac{1}{N_i}$$

The effective population size, N_e , is the “harmonic mean” of N_0, N_1, \dots, N_{t-1} .

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What is N_e good for?

In a population of varying size, heterozygosity at neutral loci is

$$\mathcal{H} = \frac{4N_e u}{4N_e u + 1}?$$

where N_e is the effective population size.

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Example

- What is the arithmetic mean of 1, 50, and 100?
- What is the harmonic mean?

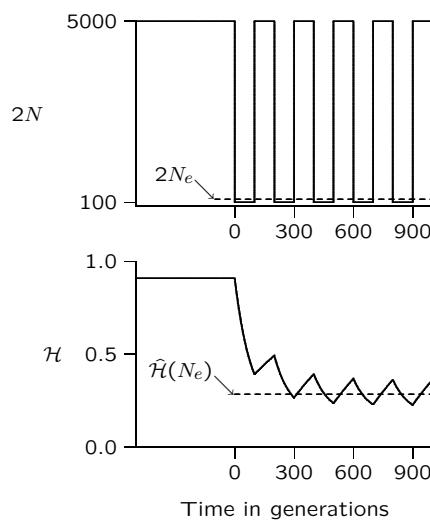
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Answer

- Arithmetic mean: $(1 + 50 + 100)/3 = 50.3333$.
- Harmonic mean: $1/((1.0 + 1.0/50 + 1.0/100)/3) = 2.9126$.

Harmonic mean is *much* smaller than arithmetic mean.

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What do we expect of the Emeryville elk? We would need to look at \mathcal{H} in other populations of large mammals.

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