

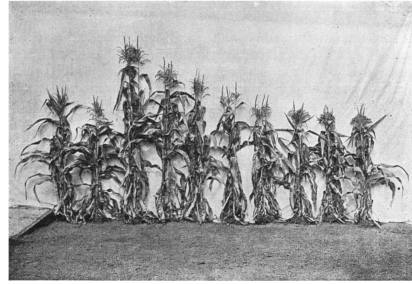
Inbreeding

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March 24, 2020

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Inbreeding depression in corn

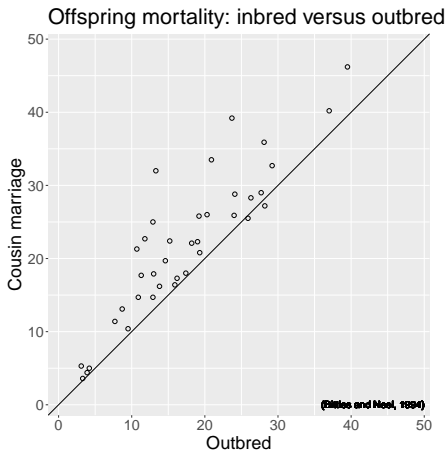


(JONES 1924)

- ▶ Two plants on left are from inbred homozygous strains.
- ▶ Next: the F_1 offspring of these strains
- ▶ Then offspring (F_2) of two F_1 s.
- ▶ Then F_3
- ▶ And so on.

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Inbreeding depression in humans



Offspring of cousin marriages are less likely to survive.
(BITTLES AND NEEL 1994)

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Genotype frequencies without random mating

Describes any bi-allelic locus.

$F = 0$ under random mating.
Reduces to Hardy-Weinberg.

$F > 0$ under inbreeding. Gives excess of homozygotes.

F is the *coefficient of inbreeding*.

Genotype	Frequency
A_1A_1	$p^2 + pqF$
A_1A_2	$2pq(1 - F)$
A_2A_2	$q^2 + pqF$

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Example

Assume $p = 1/2$

Genotype	Frequency	
	$F = 0$	$F = 0.1$
A_1A_1	0.25	0.275
A_1A_2	0.50	0.450
A_2A_2	0.25	0.275

Inbred population has more homozygotes.

Suffers if either

- ▶ Heterozygotes tend to have high fitness
- ▶ Deleterious alleles tend to be recessive.

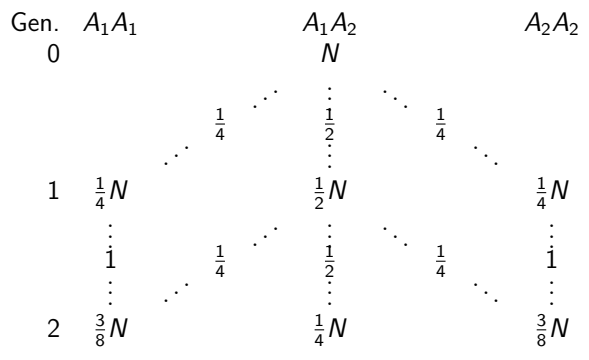
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Outline of theory

Inbreeding increases F , which increases homozygosity, which decreases fitness.

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Decay of heterozygosity under selfing



Decay of heterozygosity under selfing

Half of heterozygosity is lost each generation.

What about cousin mating, or mating between sibs?

It is *extremely* difficult to work this out, using the method we just used.

Between 1903 and 1915, no one could get it right.

Pearl (1913): Only for brother-sister matings does inbreeding reduce heterozygosity. **[Wrong!]**

Solution: build theory looking backwards in time, not forwards.

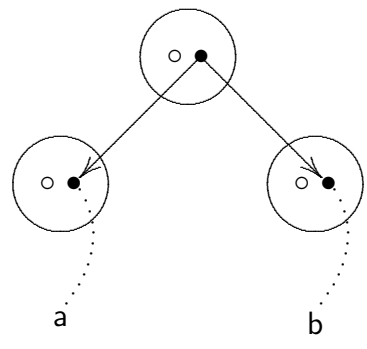
Kinds of gene identity

There are two senses in which a pair of gene copies may be "identical:"

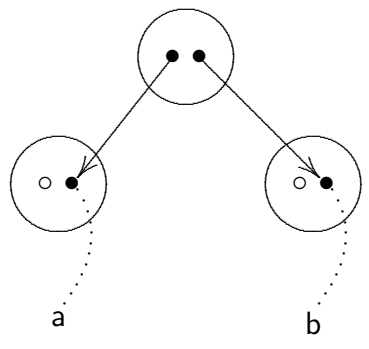
- identity in state : copies of same allele
- identity by descent : copies of same gene copy in an ancestor

Abbreviation: IBD = Identity by Descent

Gametes *a* and *b* are identical by descent



Identical in state, not by descent

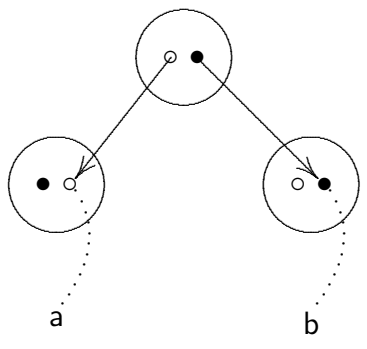


May be IBD relative to an earlier generation.

Not IBD relative to the pedigree shown here.

IBD is always relative to a particular generation.

Identical neither in state nor by descent



Uniting gametes

- Consider the two gametes that unite to form an individual.
- ▶ F , is the probability that they are IBD
- ▶ What is the probability that they both carry A_1 ?

Event	Probability
IBD from A_1 -bearing ancestor	Fp
Descend from two random ancestors who both carry A_1	$(1 - F)p^2$

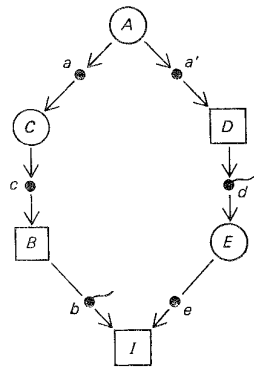
$$P_{11} = Fp + (1 - F)p^2 = p^2 + pqF$$

All three genotypes

Genotype	Frequency
A_1A_1	$p^2 + pqF$
A_1A_2	$2pq(1 - F)$
A_2A_2	$q^2 + pqF$

Same formulas as before.
 F is no longer arbitrary.
 F is probability that uniting gametes are IBD.

Calculating F from a pedigree



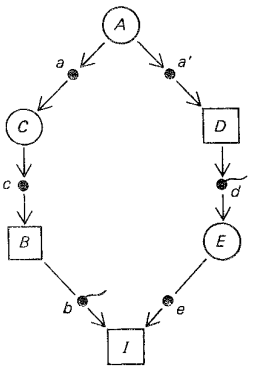
F is probability that b and e are IBD, which I abbreviate as $\Pr[b = e]$.

$$F = \Pr[b = c] \times \Pr[c = a] \times \Pr[a = a'] \times \Pr[a' = d] \times \Pr[d = e]$$

$\Pr[b = c] = 1/2$, because b has an equal chance of coming from C or from B 's other parent.

Ditto $\Pr[c = a]$, $\Pr[a' = d]$, & $\Pr[d = e]$.

$\Pr[a = a']$ is also 1/2

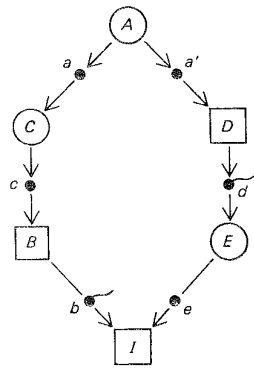


Let \oplus and \otimes represent the two gene copies in A . There are four equally likely possibilities for gametes a and a' : (\oplus, \oplus) , (\oplus, \otimes) , (\otimes, \oplus) , or (\otimes, \otimes) .

$a = a'$ in half of these possibilities, so $\Pr[a = a'] = 1/2$.

Regardless of which gene copy A contributes to C , there is a 50% chance that he contributes the same one to D .

Calculating F from a pedigree (conclusion)

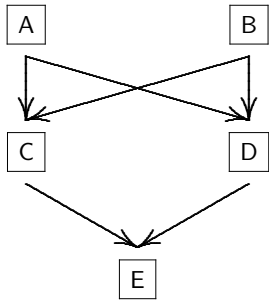


$$F = \Pr[b = c] \times \Pr[c = a] \times \Pr[a = a'] \times \Pr[a' = d] \times \Pr[d = e]$$

Each of the probabilities in this product equals 1/2, so $F = 1/2^5$.

In general, each loop in the pedigree contributes $1/2^n$, where n is the number of ancestors in the loop.

Mating between full siblings



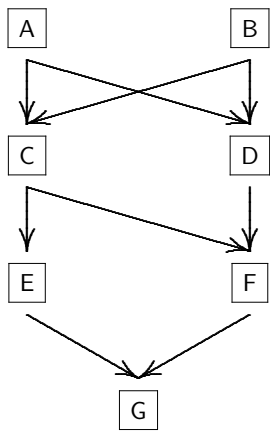
The two loops are mutually exclusive: the two gene copies in *E* cannot be IBD from *A* and also from *B*.

F is the sum of the probabilities of the 2 loops.

Each loop contributes $1/2^3$.

$$F = 2 \times \frac{1}{2^3} = 1/2^2$$

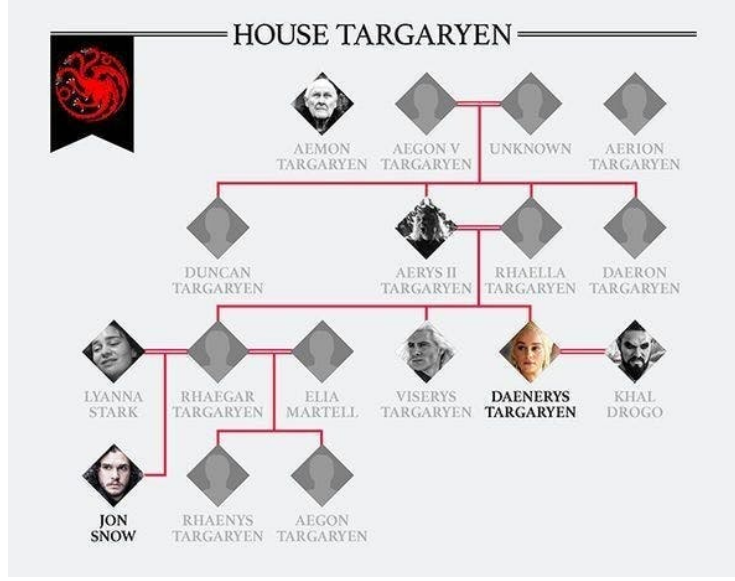
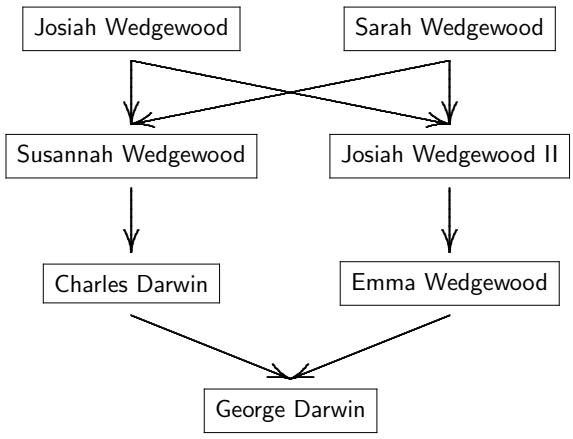
A complex pedigree



Three loops with probabilities $1/2^3$, $1/2^5$, and $1/2^5$.

$$F = 1/2^3 + 2 \times \frac{1}{2^5} = 3/2^4$$

Darwin-Wedgewood Genealogy



Summary

- ▶ The inbreeding coefficient, *F*, is the probability that the two gene copies in an individual are IBD from some given generation in the past.
- ▶ Inbreeding subtracts $2pqF$ from heterozygosity and adds pqF to the frequency of each of the two homozygous genotypes.
- ▶ Each loop in a pedigree contributes $1/2^n$ to *F*, where *n* is the number of ancestors in the loop.
- ▶ Sum across loops to calculate *F*.