ISFG-GHEP Online School 2024 October 7, 14, **21**, 28

Kinship and pedigree analysis: Methods and applications



Teachers

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Schedule

The course runs each Monday of October 2024, from 16 to 20 (CEST). The following schedule is tentative:

Oct 7: Theory of relatedness

- 16:00–17:00 Introduction to pedigrees, QuickPed and R (MDV)
- 17:00–17:45 Exercises I. (Solutions)
- 17:45-18:00 Break
- 18:00–19:00 Measures of relatedness (MDV)
- 19:00–19:45 Exercises II. (Solutions)
- 19:45-20:00 Wrap-up

Oct 14: Kinship testing

- 16:00-17:00 Introduction to forensic kinship testing (TE)
- 17:00–17:45 Exercises III. (Solutions)
- 17:45–18:00 Break
- 18:00–19:00 Kinship testing with Familias (TE)
- 19:00–19:45 Exercises IV. File needed: kinship-riddle.fam. (Solutions)
- 19:45-20:00 Wrap-up

Oct 21: Relatedness inference

- 16:00-17:00 Realised relatedness: Why are some siblings more alike than others? (MDV)
- 17:00-17:45 Exercises V
- 17:45–18:00 Break
- 18:00-19:00 Pedigree reconstruction (MDV)
- 19:00-19:45 Exercises VI
- 19:45-20:00 Wrap-up

Oct 28: Disaster victim identification

- 16:00-17:00 DNA-based disaster victim identification (TE)
- 17:00-17:45 Exercises VII
- 17:45-18:00 Break
- 18:00-19:00 Practical DVI with Diviana (MDV)
- 19:00–19:45 Exercises VIII
- 19:45–20:00 Wrap-up

Home page

https://magnusdv.github.io/pedsuite/ articles/web_only/course-ghep2024.html





Lecture 5: Realised relatedness or Why are some siblings more alike than others?

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Meiotic recombination







- Genetic distance between two loci:
 - = average # crossovers per meiosis
- Units:
 - 1 Morgan (M) = 1 crossover per meiosis
 - 1 centiMorgan (cM) = 0.01 M
- The human genome: Ca 30 Morgan



Rule of thumb: One crossover per chromosome arm

















Half first cousins, expected sharing:

$$\kappa_1 = 2 \cdot \left(\frac{1}{2}\right)^4 = \frac{1}{8} = 12.5\%$$



Realisert k1





Realised inbreeding



f_R = autozygous fraction of genome



Autozygous segments





Realised IBD coefficients







ibdsim2 Variation in realised IBD coefficients 1000 simulations • MZ Some siblings are more alike than others! k_2 library(pedsuite) > library(ibdsim2) > x = nuclearPed(2)> s = ibdsim(x, N = 1000)> k = realisedKappa(s, ids = 3:4) >PO H,Ŭ,G FC UN k_0 showInTriangle(k) >





Variation depends on the genome







Indistinguishable relationships?









Simulated IBD distributions















The probability of zero IBD



N'th cousins	P(zero IBD)
first	0.0 %
second	0.0 %
third	1.5 %
fourth	28 %
fifth	67 %

Third cousins

Expected fraction with IBD = 1:

$$k_1 = \frac{1}{64}$$

Two individuals can have a common ancestor without being genetically related





Distant cousins share either nothing or quite a bit





Reversely:

Is 100 % inbreeding possible?

- Continuous full-sib mating
- Easy to show:
 - inbreeding coefficient $f \rightarrow 1$
- In finite pedigree:
 - pedigree-based f will never reach 1

After ~30 generations, But: the realised inbreeding typically reaches 1





The importance of sex

- Rule of thumb:
 - $1 \,\mathrm{cM} \approx 1 \,\mathrm{Mb}$
- But: crossover rates vary
 - across the genome
 - males vs. females



Genetic map of chromosome 1

Females have a much longer genome!





Can we separate these??







Yes!







Napoleon Bonaparte (1769 - 1821)











Summary of genetic relatedness

- Pedigree-based measures:
 - the kinship/inbreeding coefficient arphi
 - the kappa coefficients $\kappa = (\kappa_0, \kappa_1, \kappa_2)$
- Each coefficient is
 - the probability of observing a certain IBD pattern in a random locus
 - the expected proportion of the genome in this state
- Realised relatedness:
 - IBD segments determined by meiotic recombination
 - females recombine more than males
 - may separate relationships with equal kappa's
 - 0% and 100% realised identity is possible!





So...what does it mean to be related?

- Pedigree based definition: φ > 0
 <u>potentially</u> having alleles IBD
- Genomic definition (**realised** relatedness): <u>actually</u> having alleles IBD



