

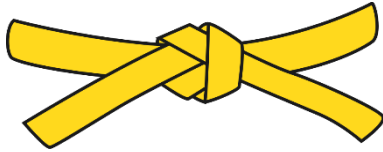


Lecture 2: Measures of relatedness

Magnus Dehli Vigeland

Young Investigator Day 2024

Plan



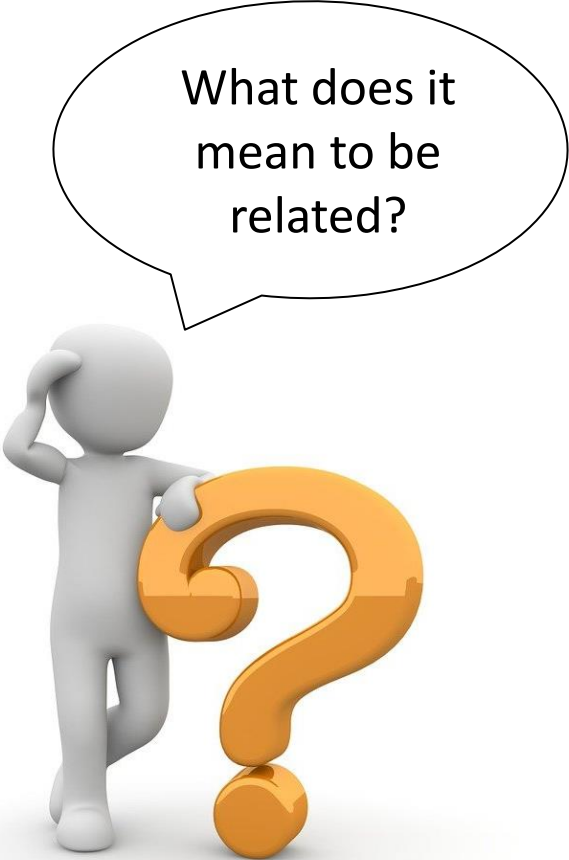
Kinship/inbreeding coefficient



The relatedness triangle



Realised relatedness



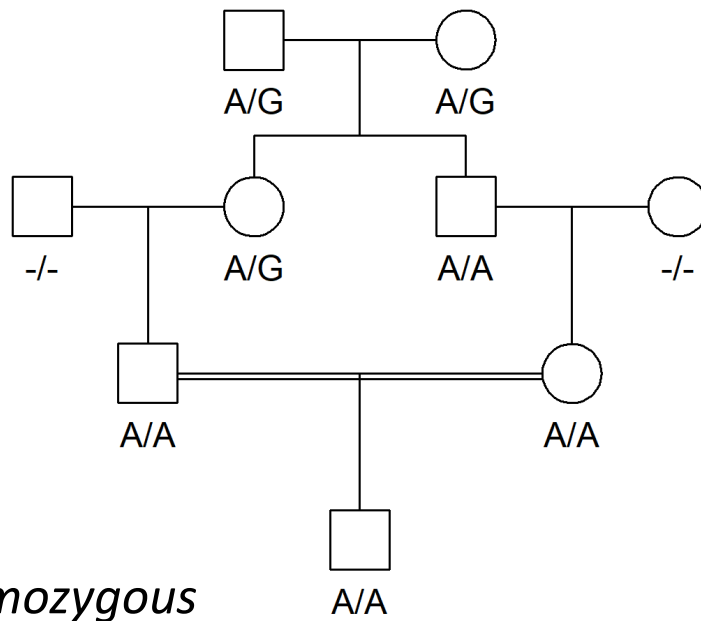
What does it
mean to be
related?

- Attempt 1
 - being connected through a pedigree
 - having a common ancestor...not too far back
- Attempt 2 - genetic
 - sharing DNA?
 - (more than unrelated people)
- To make this precise, we need some terminology!

Identity by descent (IBD)

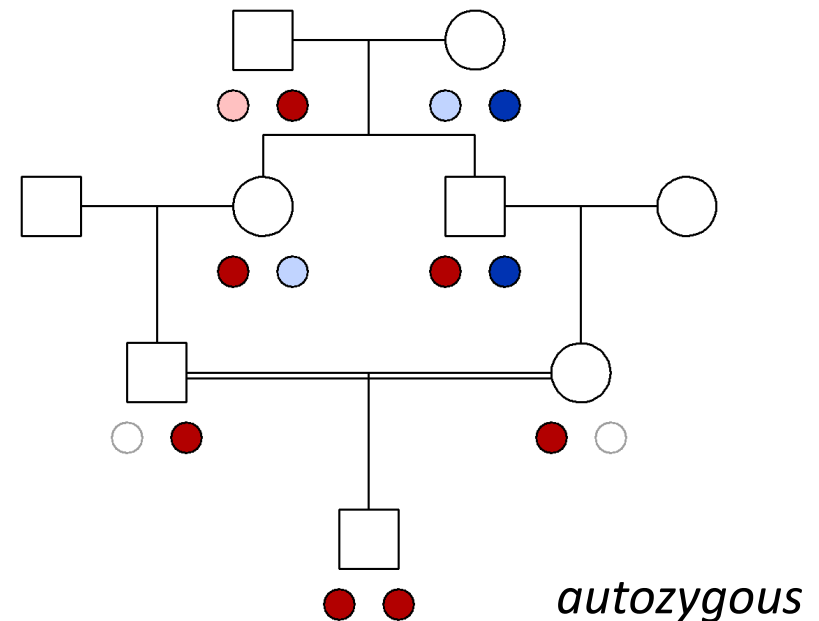
Identity by *state*

= equal alleles



Identity by *descent*

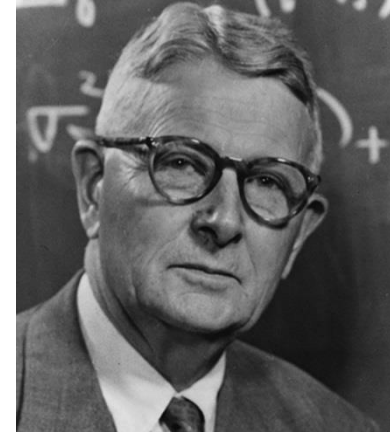
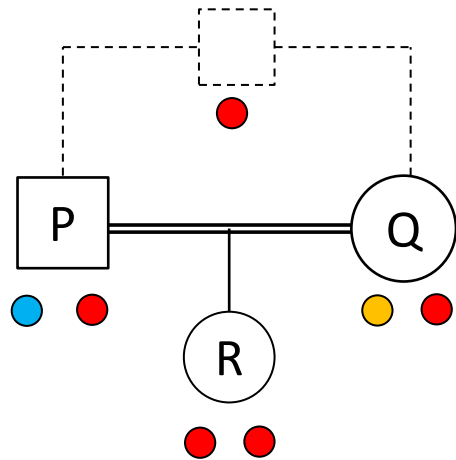
= same origin in the pedigree



Inbreeding coefficient
 $f = Pr(\text{autozygosity})$



Coefficient of kinship/inbreeding



Sewall Wright
(1889 - 1988)

- Wright (1921): The kinship coefficient φ between P and Q

$$\varphi_{P,Q} = Pr(\text{random allele of P is IBD with random allele of Q})$$

$$= Pr(\text{R receive IBD alleles from her parents})$$

$$= Pr(\text{R is autozygous})$$

$$= f_R$$

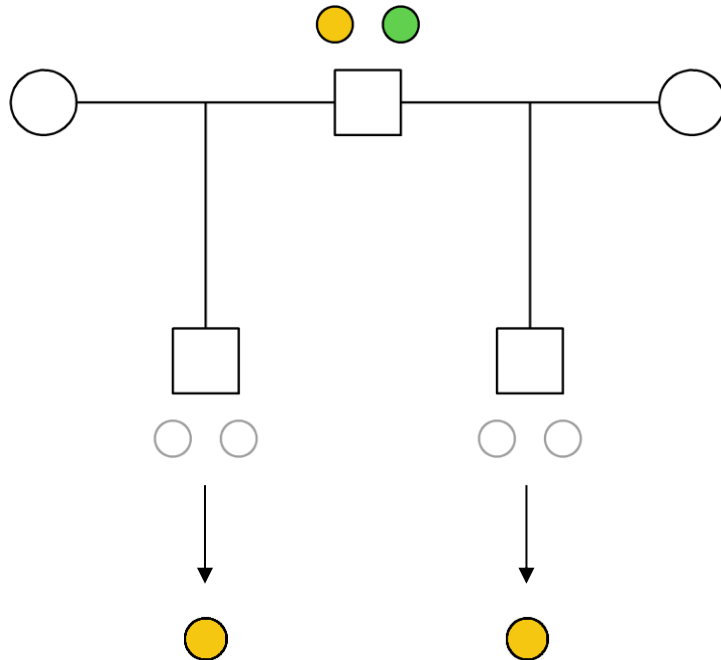
the inbreeding coefficient of R

P and Q related



$$\varphi_{P,Q} > 0$$

Example: Kinship coefficient of half siblings



Kinship coefficient

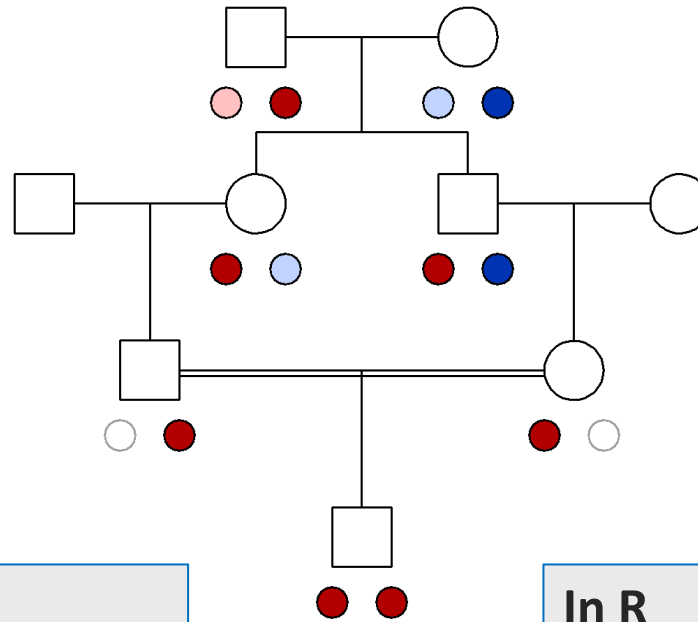
$$\begin{aligned}\varphi &= P(\text{from both}) \cdot 2 \\ &= 0.5^4 \cdot 2 \\ &= 1/8\end{aligned}$$

↑
green

Inbreeding coefficient: Example

Wright's path formula:

$$\varphi_{P,Q} = \sum_A \sum_v \left(\frac{1}{2}\right)^{|v|+1} (1 + f_A)$$



By hand

$$\begin{aligned} f &= P(\text{red circle autozygous}) \cdot 4 \\ &= 0.5^6 \cdot 4 \\ &= 1/16 \end{aligned}$$

↑
other colors

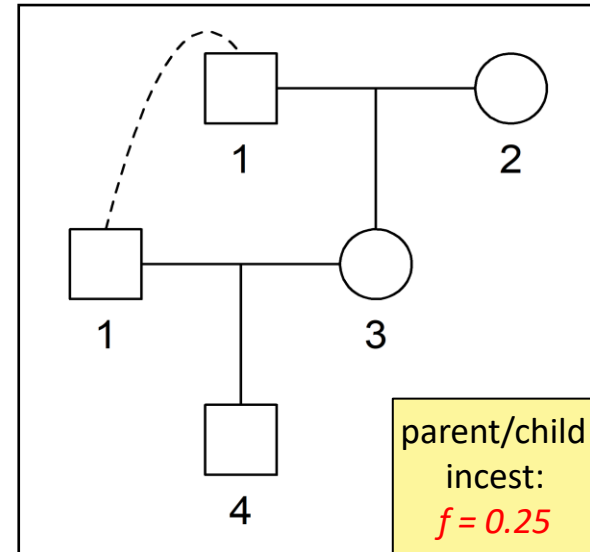
In R

```
> library(pedsuite)
> x = cousinPed(1, child = T)
> inbreeding(x, ids = 9)
[1] 0.0625
```

More kinship & inbreeding coefficients

Relationship	kinship ϕ = f of child
Parent-child	1/4
Full siblings	1/4
Half siblings	1/8
Grandparent-grandchild	1/8
Avuncular (uncle/aunt)	1/8
1st cousins	1/16
2nd cousins	1/64
3rd cousins	1/256

Challenge
Different relationships
with the same kinship!



```
> x = nuclearPed(1, sex = 2) |>
  addSon(parents = c(1, 3))

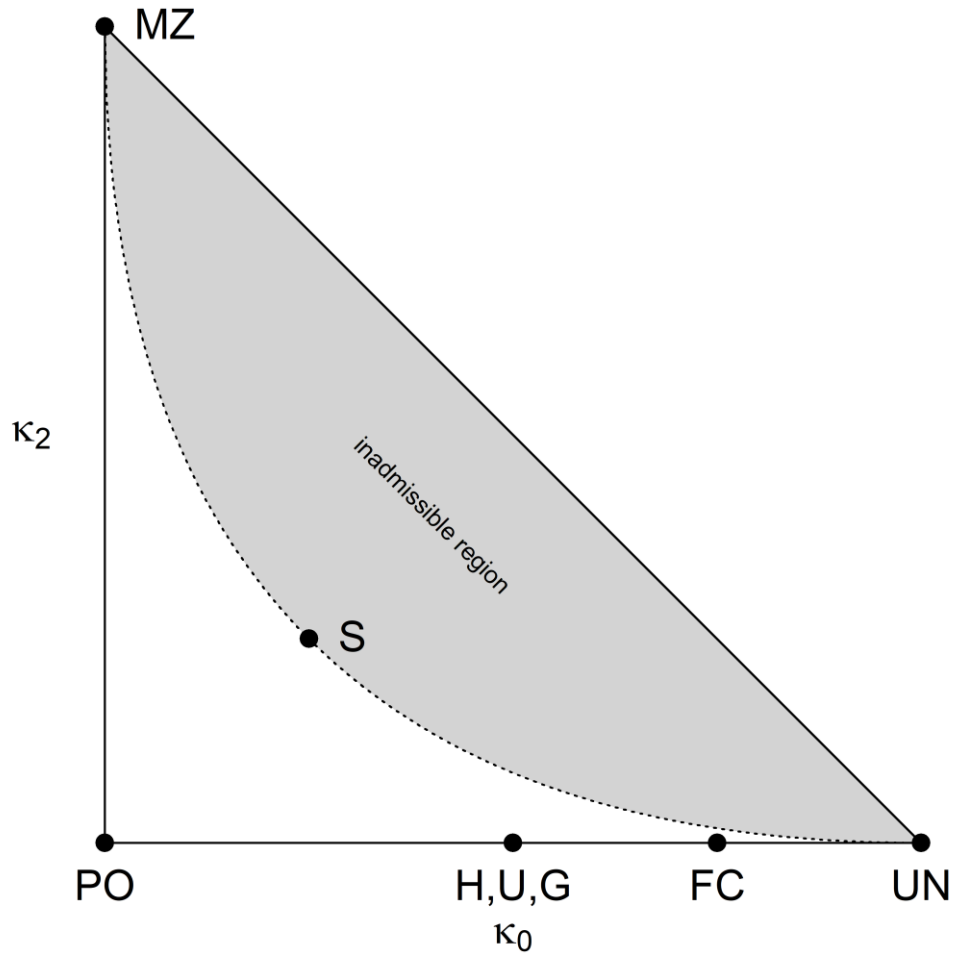
> kinship(x, ids = c(1, 3))
[1] 0.25

> inbreeding(x, id = 4)
[1] 0.25
```




The relatedness triangle

(or: the IBD triangle)



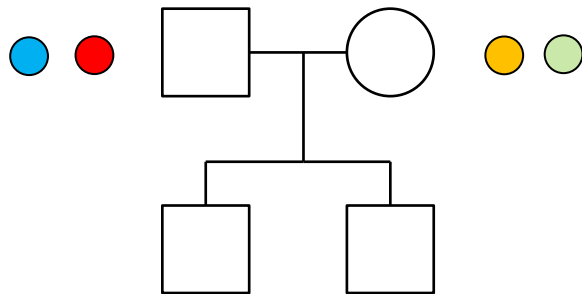
Charles Cotterman
(1914-1989)








Elisabeth Thompson
(1949 -)

IBD coefficients





- Summary so far:
 - Two individuals are related if they can have IBD alleles
 - Their kinship coefficient measures the amount of IBD sharing
- Natural generalisation:
 - How *many* alleles are IBD in each locus?







 0 alleles IBD. Prob = 0.25


 1 allele IBD. Prob = 0.25 + 0.25 = 0.5

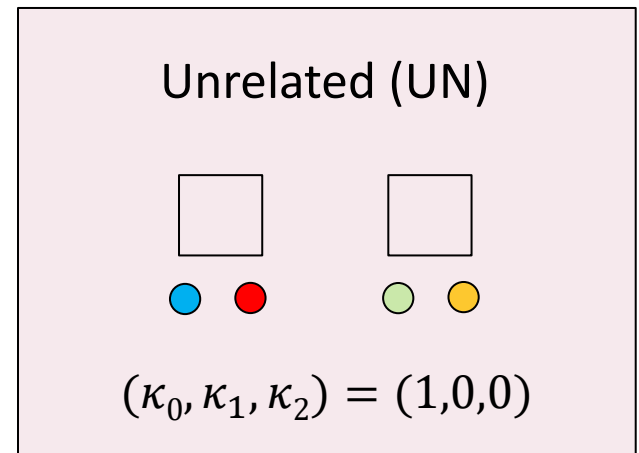
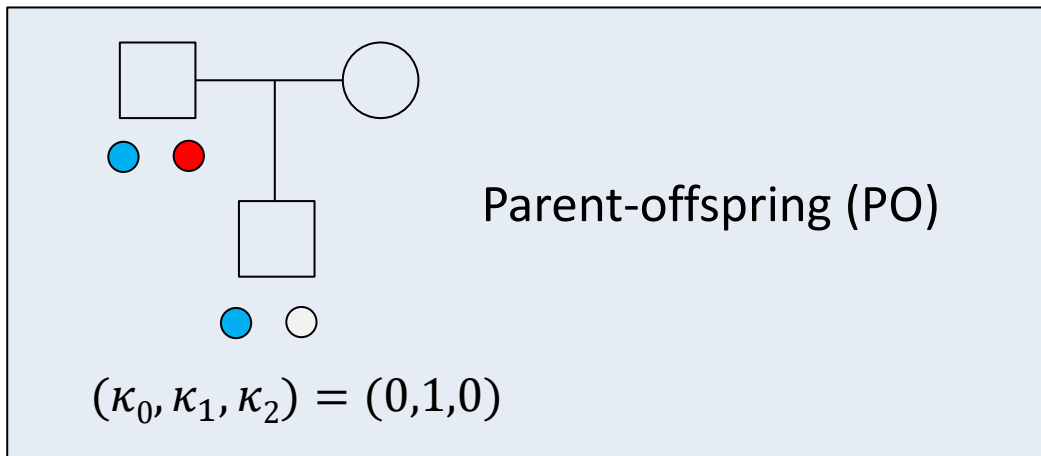
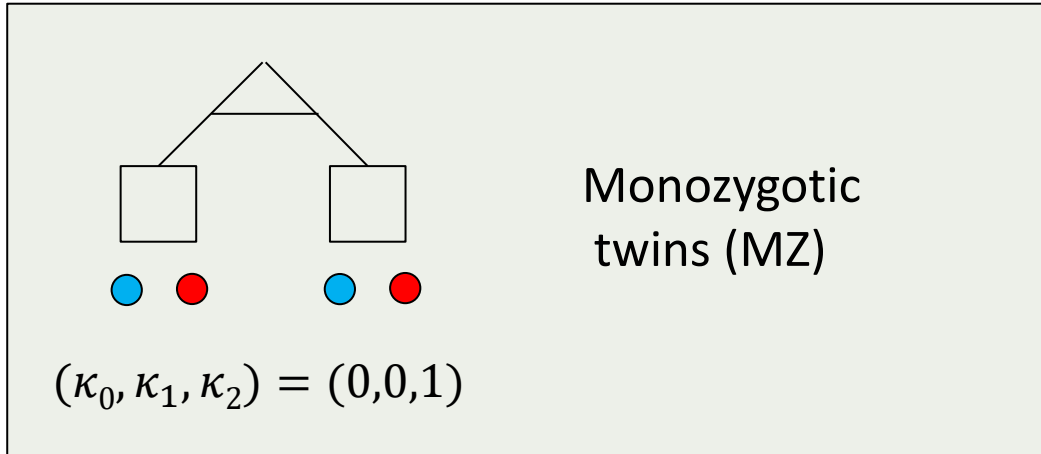




 2 alleles IBD. Prob = 0.25

Definition

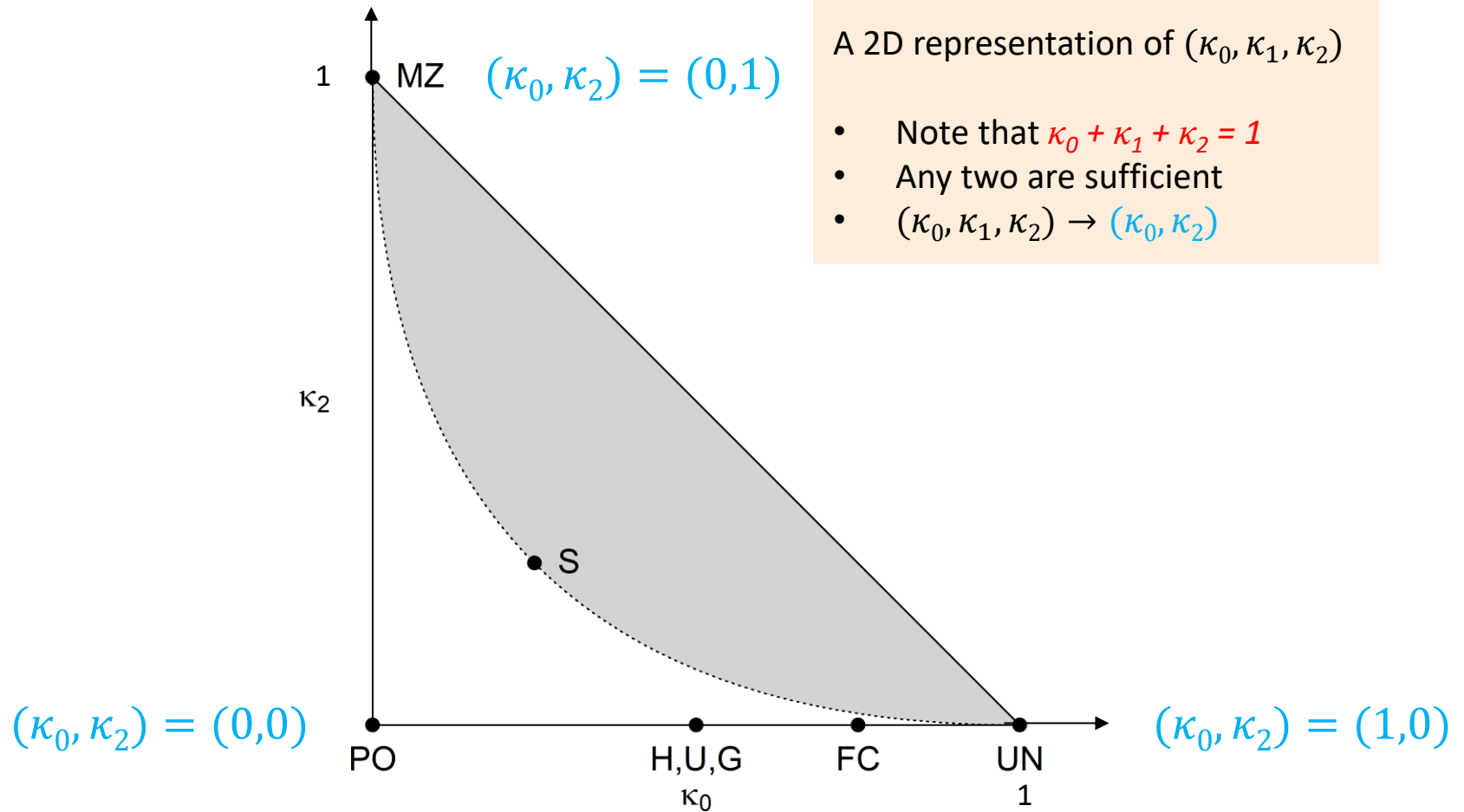
- $\kappa_0 = Pr(\text{IBD} = 0)$
- $\kappa_1 = Pr(\text{IBD} = 1)$
- $\kappa_2 = Pr(\text{IBD} = 2)$

(at random autosomal locus)

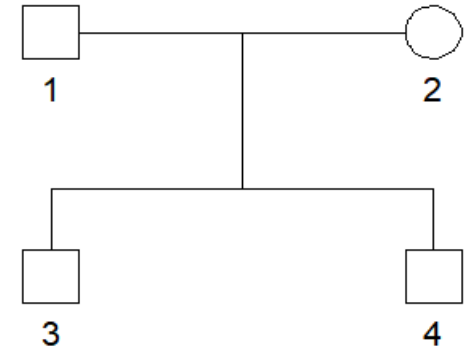
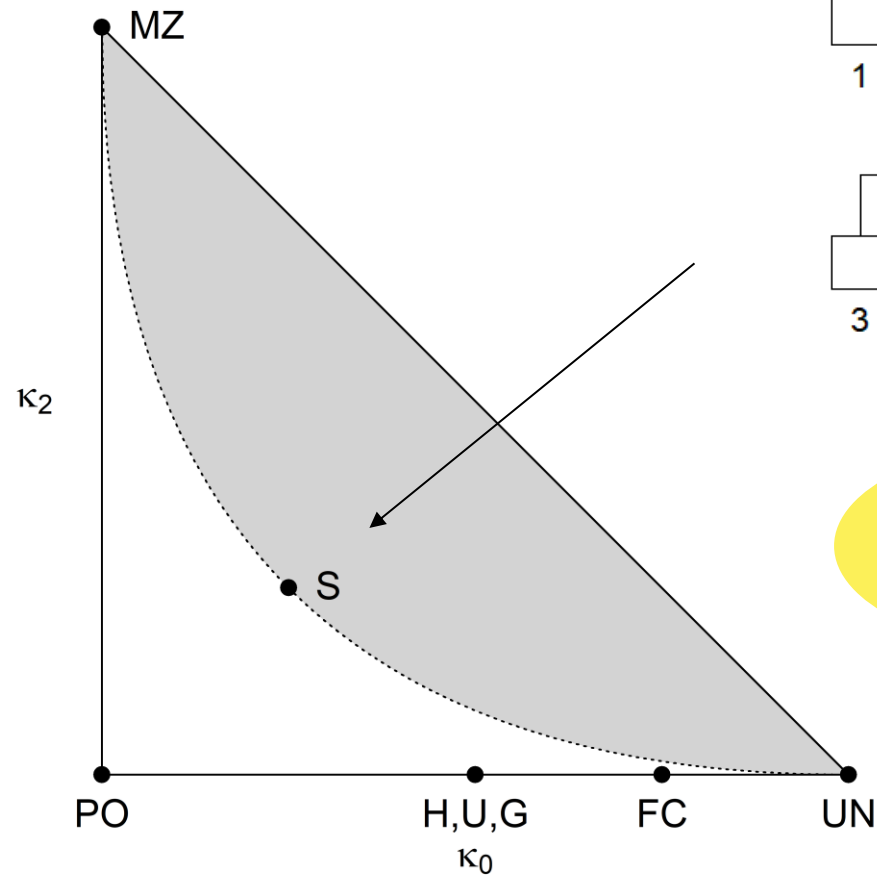
Three trivial relationships



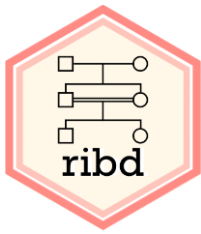
The relatedness triangle



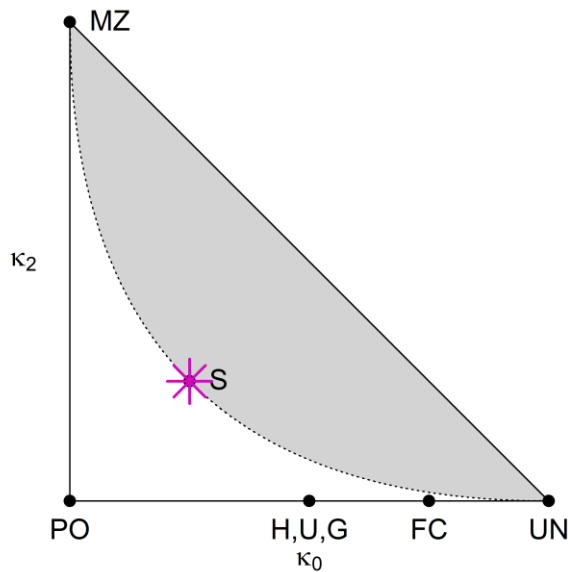
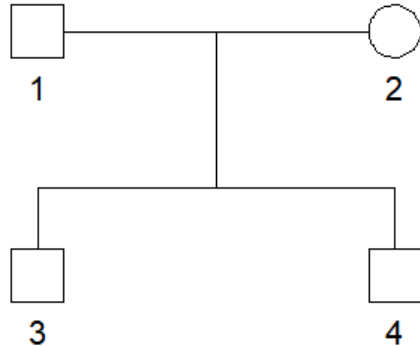
What are the coefficients of full sibs



Let's do this
in R!



ribd: Pedigree-based relatedness coefficients



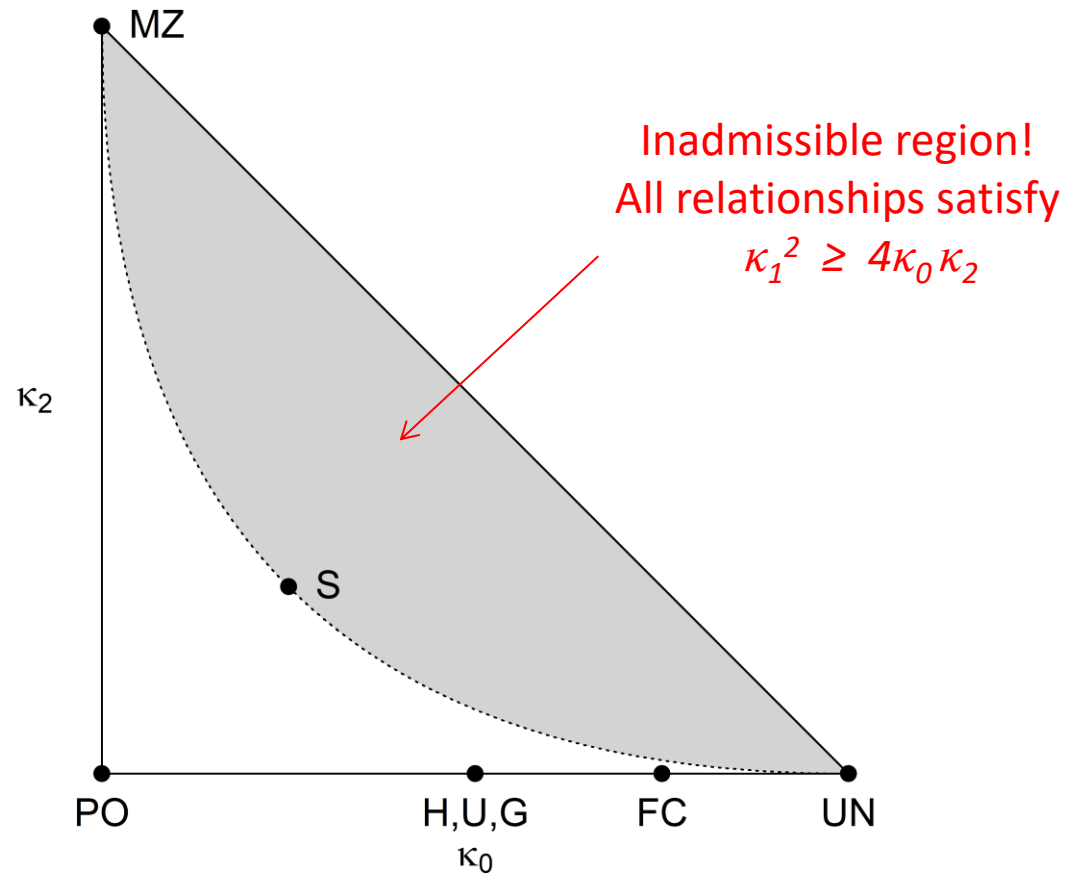
```
> library(pedsuite)
> x = nuclearPed(2)

> kinship(x, ids = 3:4)
[1] 0.25

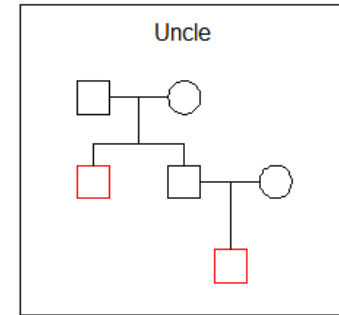
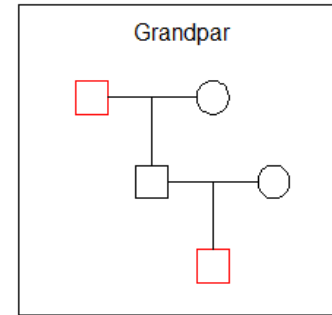
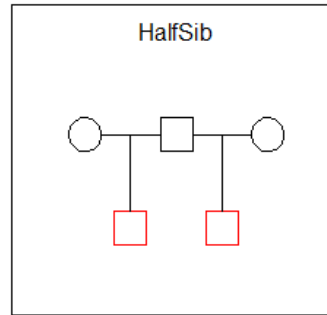
> kappaIBD(x)
id1 id2 kappa0 kappa1 kappa2
1 2 1.00 0.0 0.00
1 3 0.00 1.0 0.00
1 4 0.00 1.0 0.00
2 3 0.00 1.0 0.00
2 4 0.00 1.0 0.00
3 4 0.25 0.5 0.25

> k = kappaIBD(x, ids = 3:4)
> showInTriangle(k)
```

The relatedness triangle

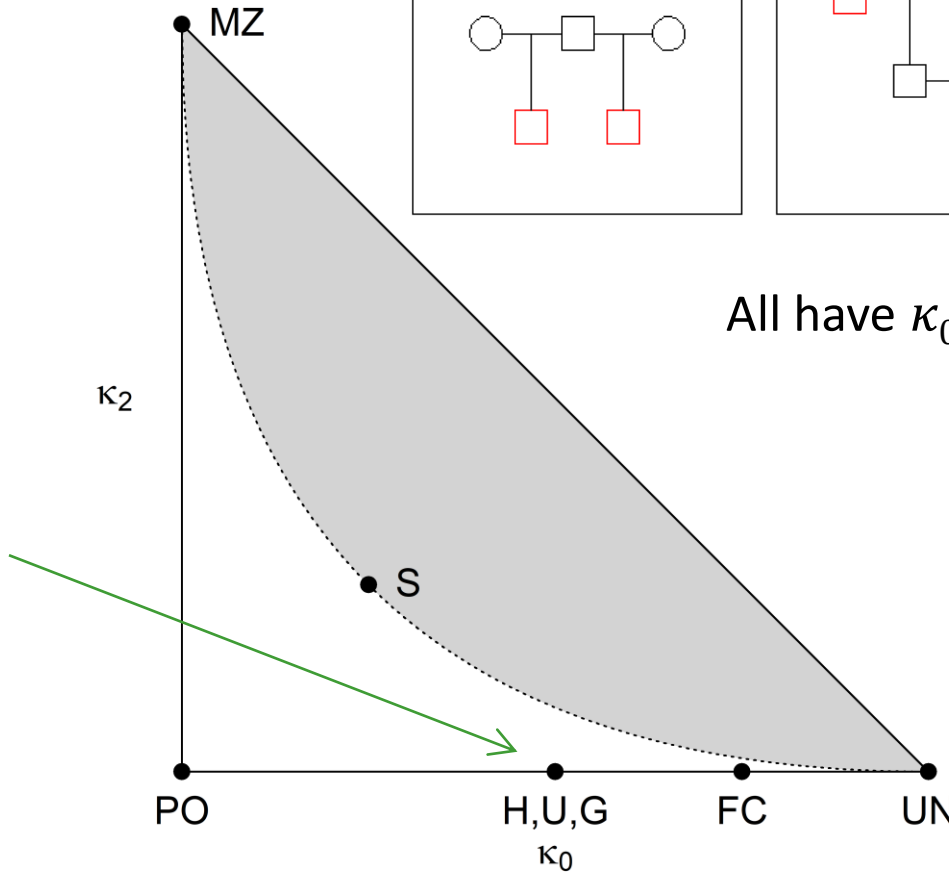


The relatedness triangle



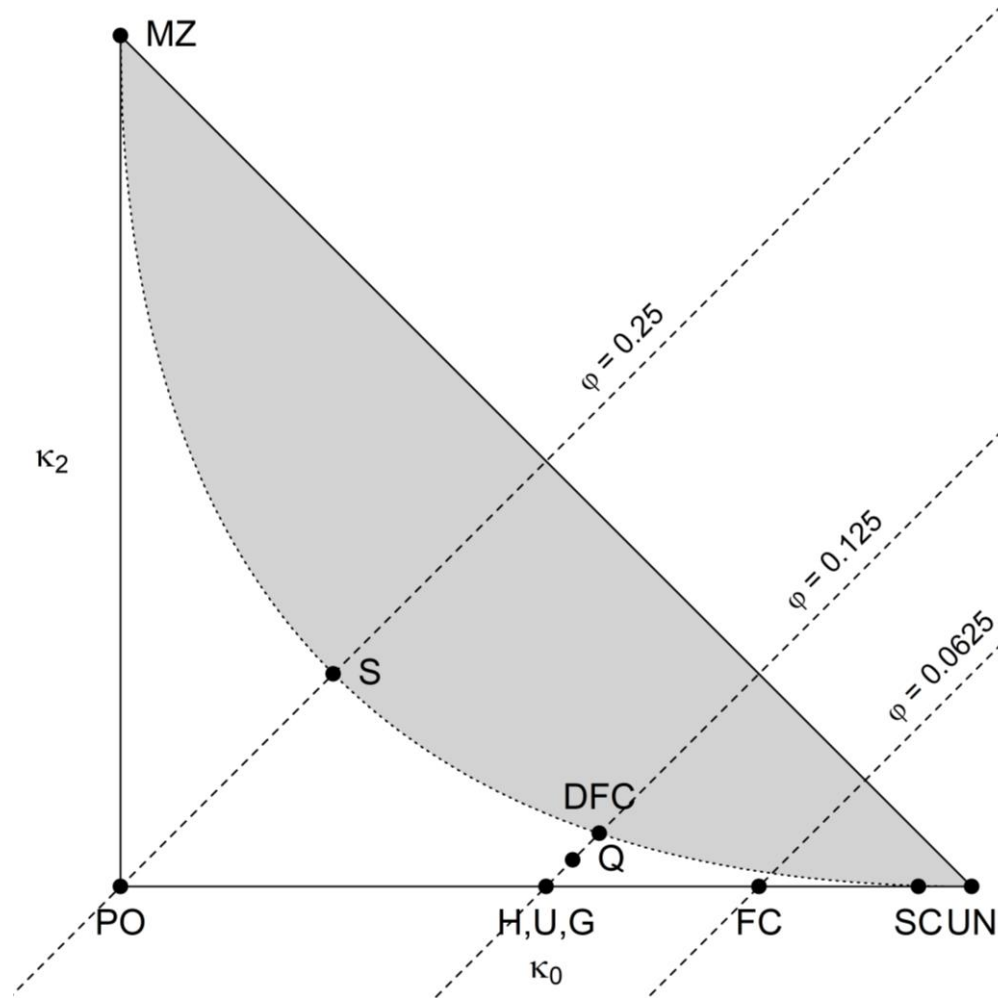
All have $\kappa_0 = \kappa_1 = \frac{1}{2}$

Some relationships coincide!

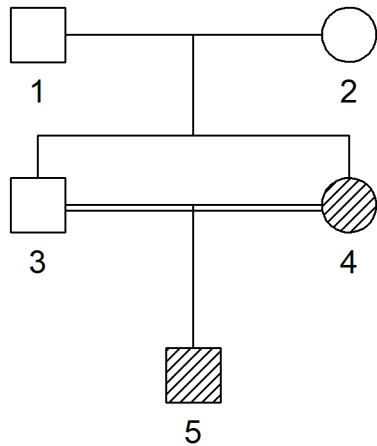


An important identity:

$$\varphi = \frac{1}{4}\kappa_1 + \frac{1}{2}\kappa_2$$



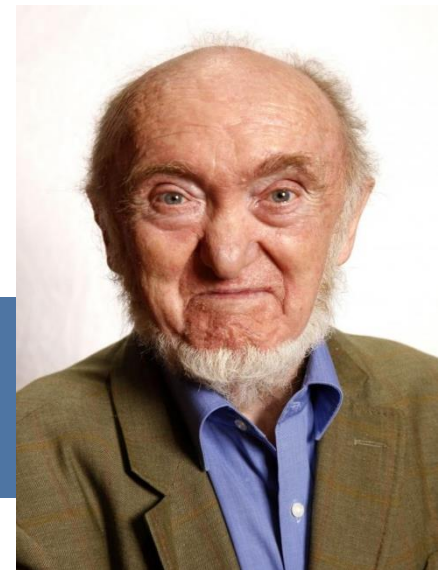
A word of caution



\mathcal{K} is only defined for non-inbred individuals.
For the whole story, we need 9 coefficients!

Jacquard's identity coefficients

Not suitable for a basic course ...




Albert Jacquard
(1925 - 2013)

QuickPed: An Interactive Pedigree Creator

New app design!

Discover the **new features**
Or stay with the old version: QuickPed3

Purpose: QuickPed lets you rapidly create attractive pedigree plots, save them as images or text files, and analyse the relationships within them.

Instructions: Choose a suitable start pedigree and modify it by clicking on individuals and using appropriate buttons. For example, to add a male child, select the parent(s) and press the  icon. Check out the [online user manual](#) for various tips and tricks, including an introduction to relatedness coefficients.

Citation: If you use QuickPed in a publication, please cite this paper: Vigeland MD (2022). QuickPed: an online tool for drawing pedigrees and analysing relatedness. *BMC Bioinformatics*, 23. DOI:10.1186/s12859-022-04759-y.


Quick start

Built-in pedigree

Trio ▼

or

Load a ped file



or

Random pedigree

Reset all

Modify

Add



Sex



Style



Fill



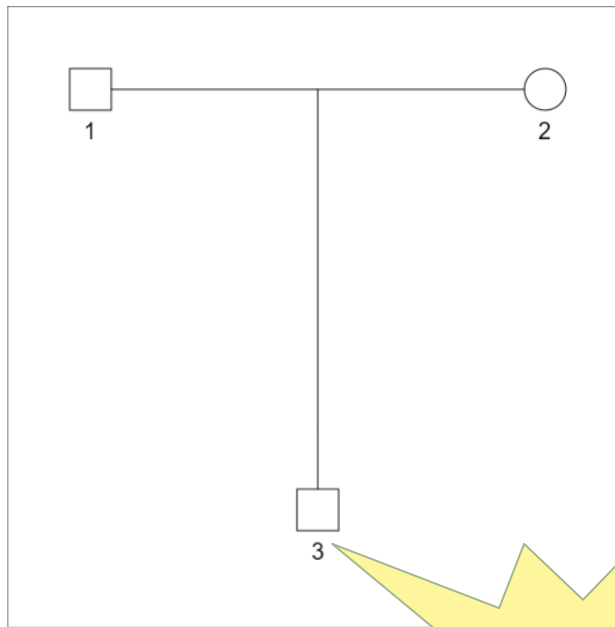
Twins

MZ / DZ

Remove



Undo



Double-click on an individual to add text



Labels

1, 2, 3, .. I-1, I-2, ..

Show all Hide all

1

2

3

Update

Plot settings

Width Height

430 430

Cex Symbols

1,4 1

Margins

3

Other options (beta)

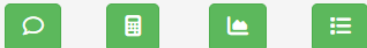
- Straight legs
- Arrows

R code

 PNG

 PDF

Relationships



Coefficients!

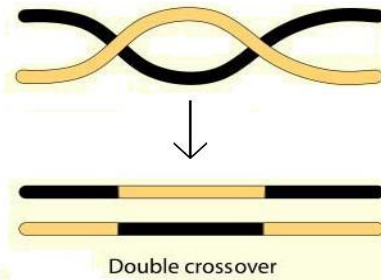
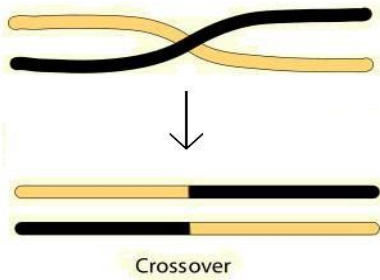
Ped file

Include

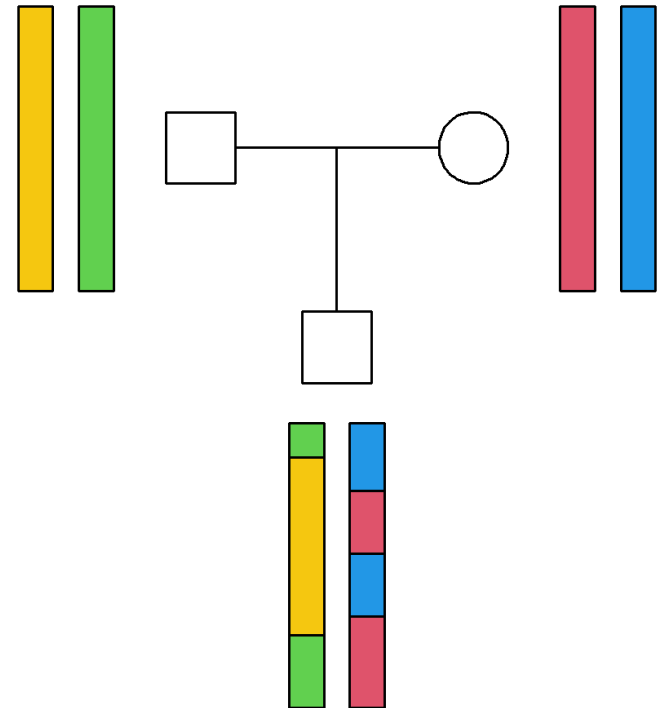
- Headers
- Family ID
- Affection status

 Save ped file

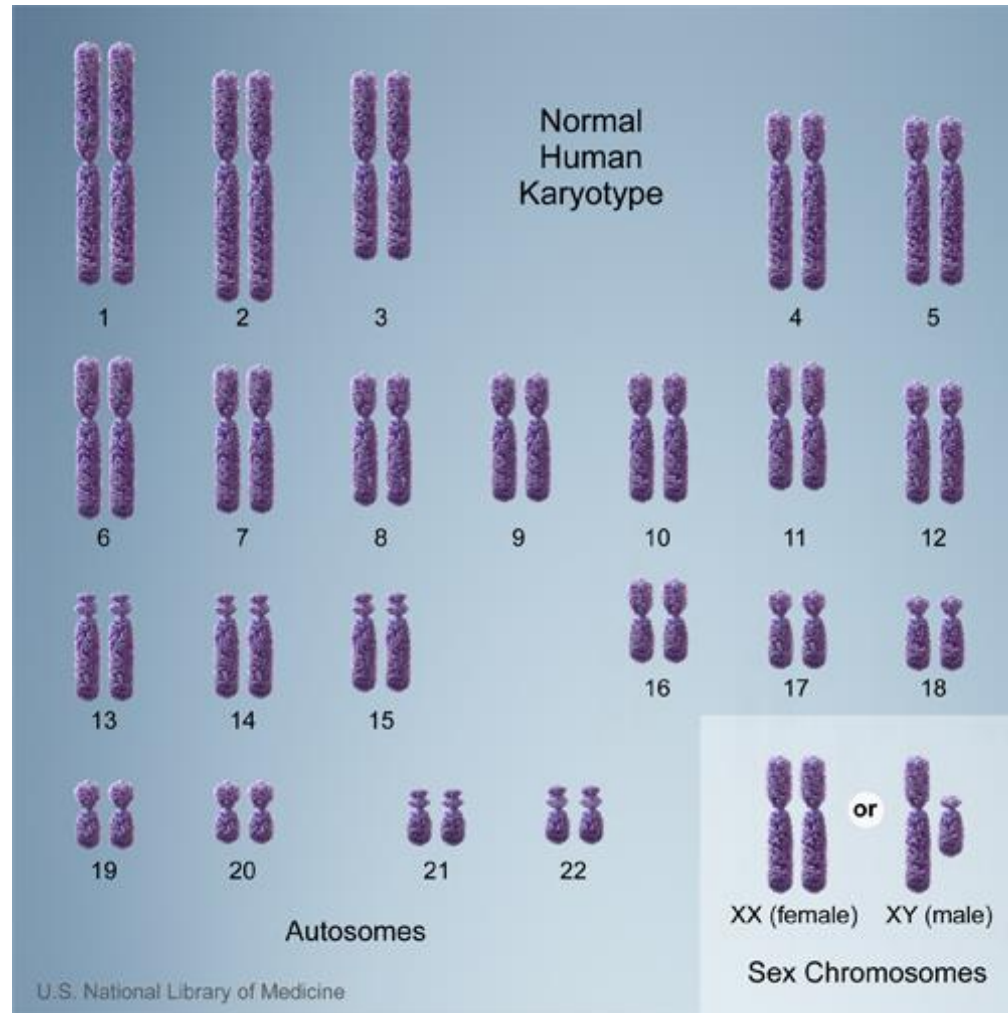
Realised relatedness



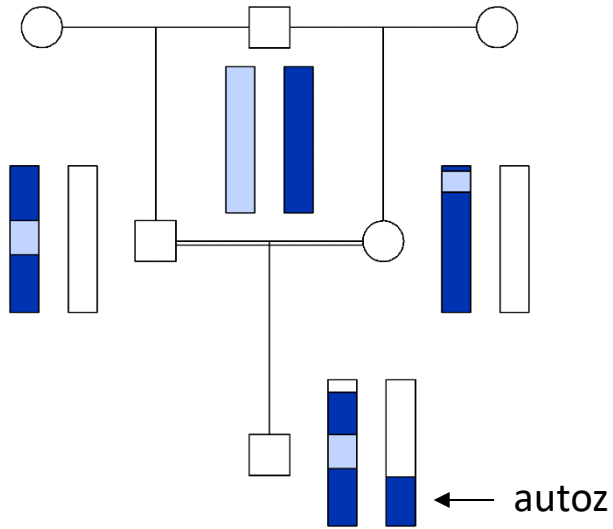
Meiotic recombination



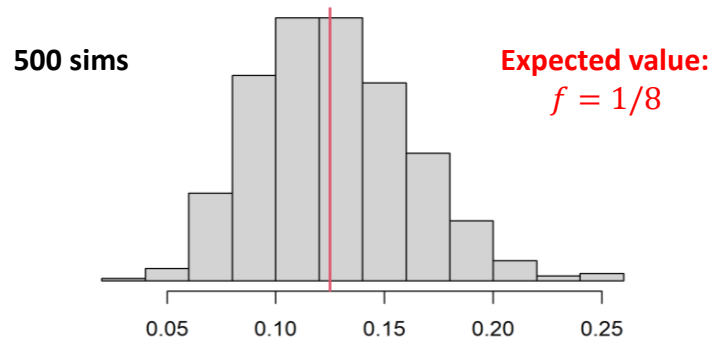
Rule of thumb: One crossover per chromosome arm



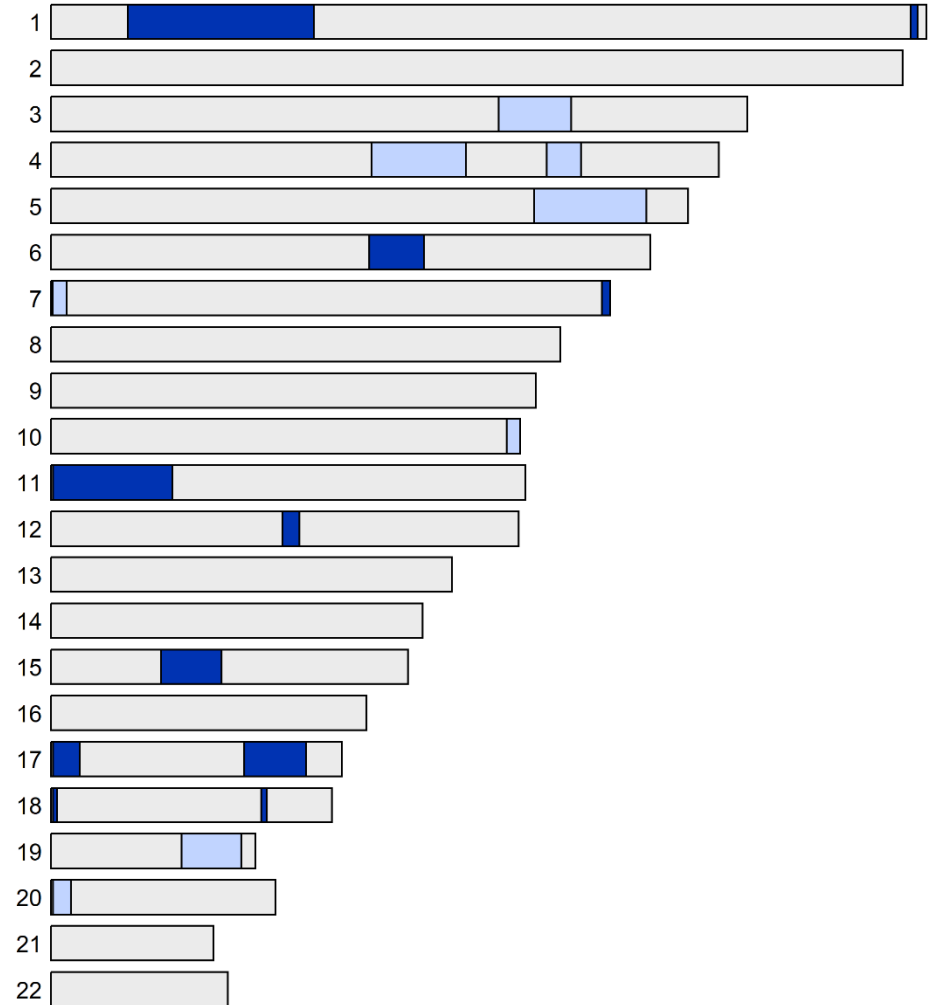
Realised inbreeding



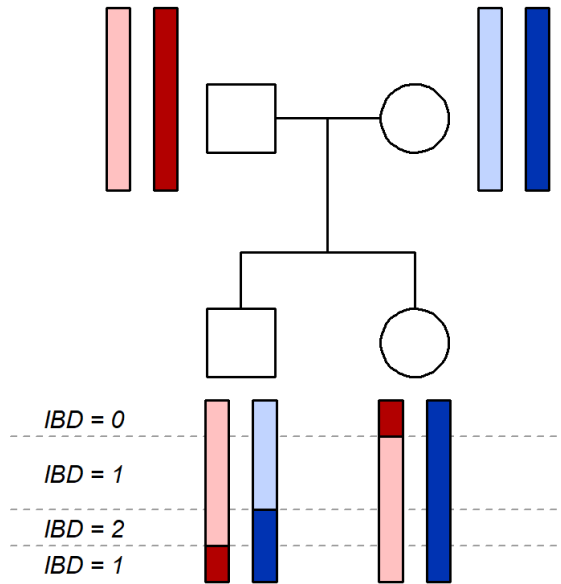
f_R = autozygous fraction of genome



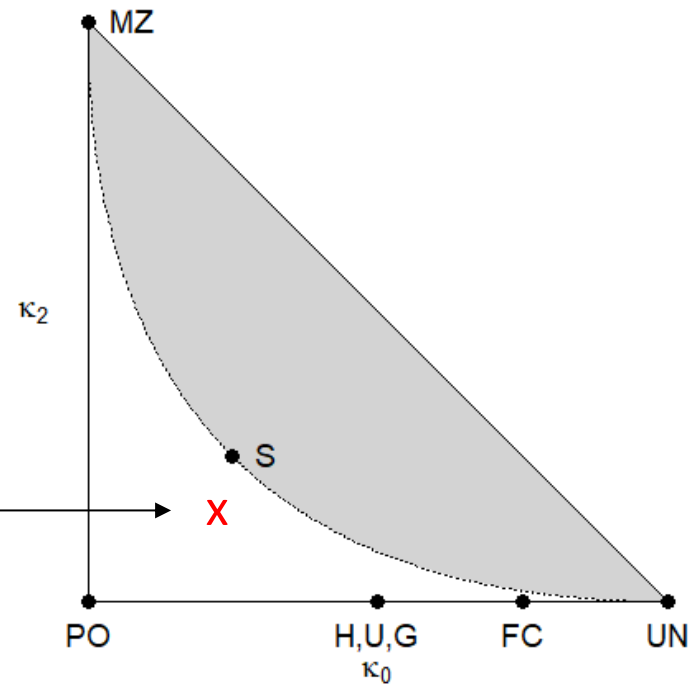
Autozygous segments



Realised IBD coefficients



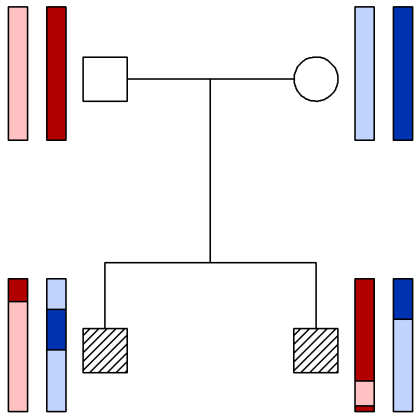
Realised IBD coefficients:
Proportions of genome with $IBD = 0, 1, 2$



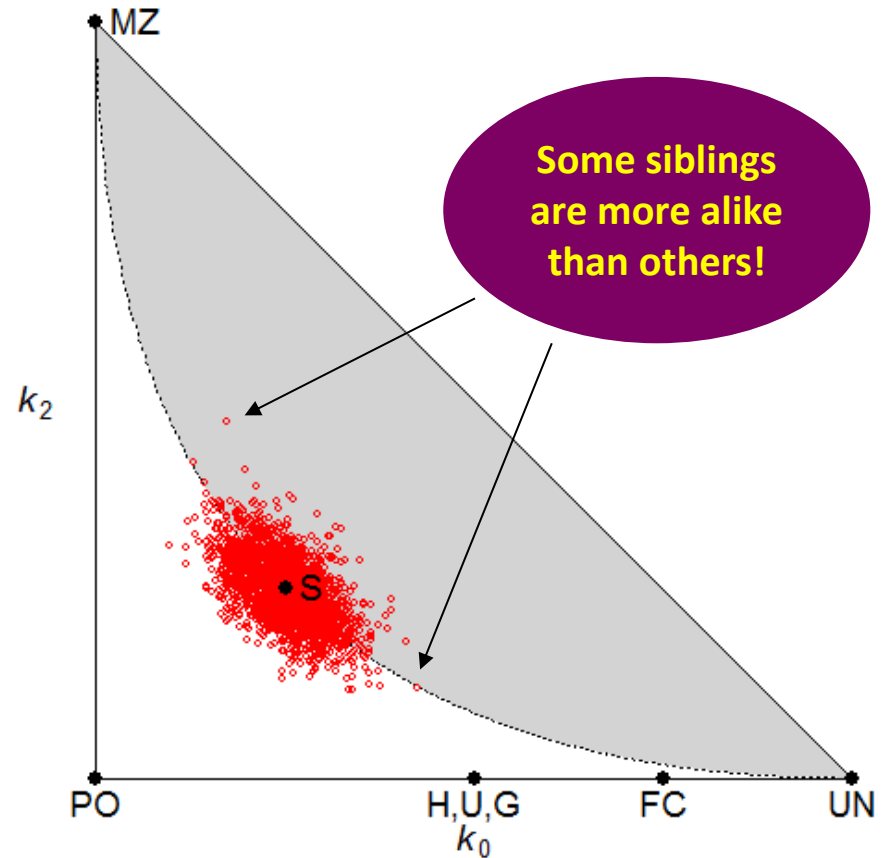
Variation in realised IBD coefficients



1000 simulations



```
> library(ibdsim2)
> x = nuclearPed(2)
> s = ibdsim(x, N = 1000)
> k = realisedKappa(s, ids = 3:4)
> ribd::showInTriangle(k)
```



Variation depends on the genome



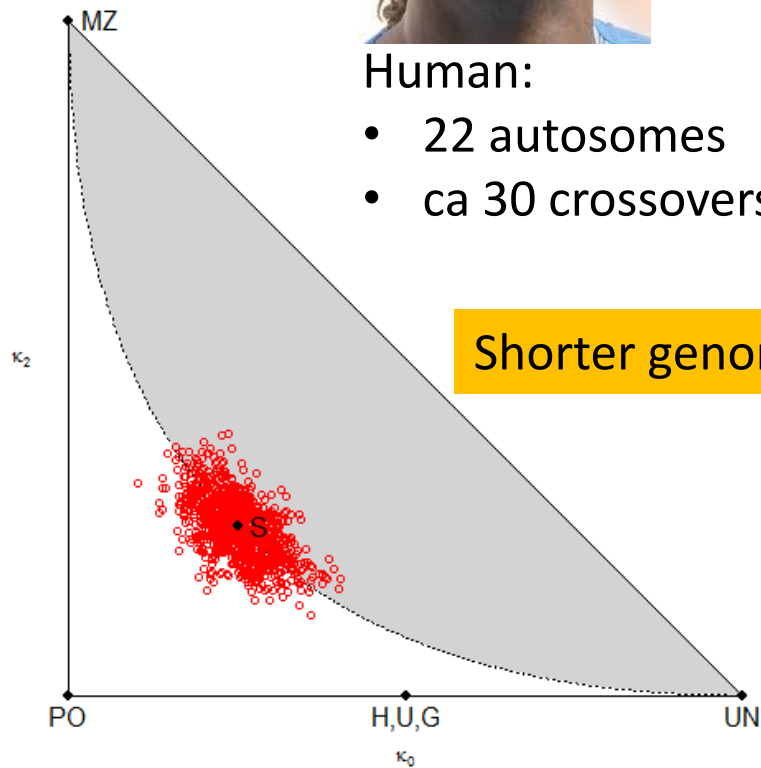
Human:

- 22 autosomes
- ca 30 crossovers

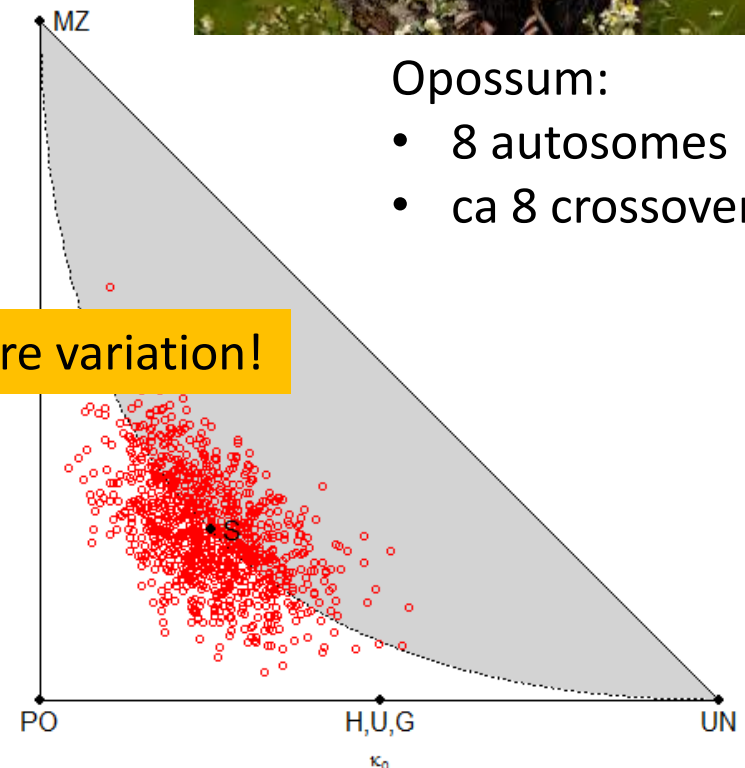


Opossum:

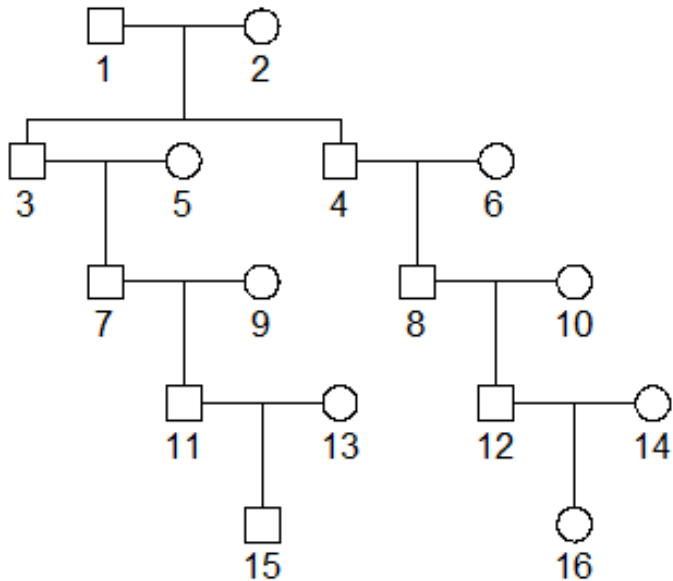
- 8 autosomes
- ca 8 crossovers



Shorter genome = more variation!



The probability of zero IBD



N'th cousins	$P(\text{zero IBD})$
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!

METRO

HOME NEWS SPORT ENTERTAINMENT LIFESTYLE
UK WORLD WEIRD TECH

Let's not forget – Prince Harry and Meghan Markle are actually (very distant) cousins

Richard Hartley-Parkinson for Metro.co.uk Monday 27 Nov 2017 11:35 am 18.8k



FASHION CULTURE

Ou, Awkward... Meghan Markle and Prince Harry are Apparently Related

By Meghan McKenna Date November 2, 2017

FASHION BEAUTY GIFT GUIDE

Brigitte

SPIELE NEWSLETTER VIDEO GEWINNSPIELE FORUM F-MAG ACADEMY SHOPPING ABC
Aktuell Mode Beauty Rezepte Gesund Liebe Familie Leben Horor

Stammbaum erforscht: Prinz Harry und Meghan Markle sind Cousins!

Brigitte → Aktuell → Stars und TV → Meghan Markle und Prinz Harry sind "verwandt"



STJERNER: Meghan Markle og prins Harry viser endelig kjærligheten sin offentlig, etter å ha holdt forholdet svært privat i lang tid. Nå kommer det frem at paret, som er fra to forskjellige kontinenter, faktisk er i slekt. Foto: NTB scanpix

Prins Harry er i slekt med kjæresten

SE OG HØR

universitetssykehus

1.1K shares

View c

(Picture: Mail Online)

Ralph BOWES
(1480–1516)
of Streatlam, Co Durham
High Sheriff

GRANDCHILDREN

Sir George BOWES
Loyal to Queen Elizabeth I
during the rising of the North, 1596

Bridget BOWES
Married John Hussey
of Dorking

GREAT-GRANDSON

Sir William BOWES MP
(1657–1707)
During reign of Charles II
Royalist

GRANDSON

Captain Christopher HUSSEY
(1598/9–1686)
A founder of Nantucket, Massachusetts

Sir George BOWES MP

Huldah HUSSEY
(1643–1740)
Married Lieutenant John SMITH

$P(\text{any IBD}) \approx 0$

Claude George BOWES
(1854–1908)
14th Earl of Strathmore and Kinghorne

George David MERRILL
(1861–1924)

Lady Elizabeth BOWES LYON
(1900–2001)
HM Queen Elizabeth
The Queen Mother

Gertrude May MERRILL
(1887–1938)
Married Frederick George SANDERS

HM The Queen
(1926–)

Doris SANDERS
(1921–)
Married Gordon Arnold Markle

Lady Diana Spencer
(1961–1997)

HRH Prince of Wales
(1948–)

Thomas Wayne Markle
(1944–)

Doria L. Ragland
(1956–)

PRINCE HARRY

(RACHEL) MEGHAN MARKLE

~~15th Cousins~~



13th cousins once removed

- 14
- 13
- 10
- 9
- 8
- 4
- 3
- 2
- 1

- 15
- 13
- 11
- 10
- 9
- 5
- 4
- 3
- 2
- 1

Relatedness: Summary

- Measuring relatedness with increasing precision:
 - the kinship/inbreeding coefficient φ
 - the IBD 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
- IBD is not a pointwise phenomenon: Always in segments
 - determined by meiotic crossovers
 - consequence: Variation in the *realised* IBD!
- Family relation \nRightarrow genetic relation

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

- Pedigree based definition: $\varphi > 0$
potentially having alleles IBD
- Genomic definition (**realised** relatedness):
actually having alleles IBD