

# Lecture 2: Measures of relatedness

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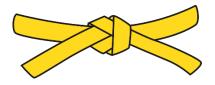
ISFG-GHEP Online School 2024

Kinship and pedigree analysis: Methods and applications





Plan



#### **Kinship/inbreeding coefficient**



**IBD coefficients & IBD triangle** 



#### Jacquard's identity coefficients





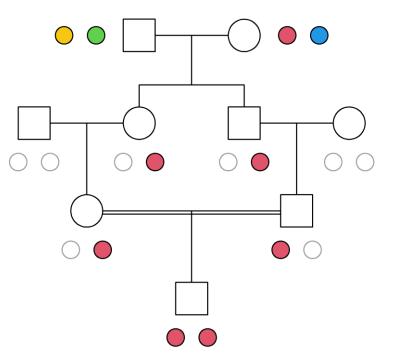


- Attempt 1
  - being connected through a pedigree
  - having a common ancestor...not too far back
- Attempt 2 genetic
  - sharing DNA?
  - (more than unrelated people)
- To make this precise, we need some terminology!



### **IBD and autozygosity**

IBD = Identical by descent
 = identical alleles with a common origin in the given pedigree



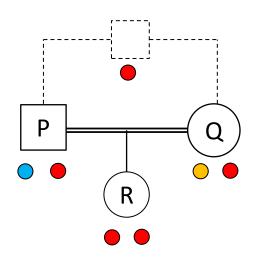
autozygous = homozygous + IBD

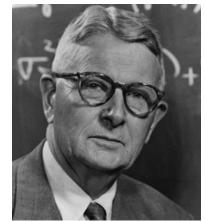
Inbreeding coefficient f = P(autozygosity)





## **Coefficient of kinship/inbreeding**





Sewall Wright (1889 - 1988)

• Wright (1921): The kinship coefficient  $\varphi$  between P and Q

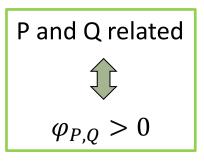
 $\varphi_{P,Q}$  = P( random allele of P is IBD with random allele of Q )

= P( R receive IBD alleles from her parents )

= P( R is autozygous )

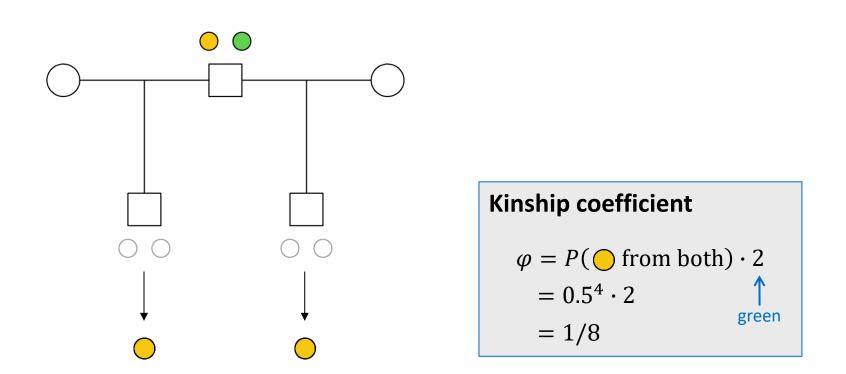


the inbreeding coefficient of R





#### **Example: Kinship coefficient of half siblings**

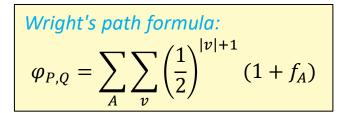


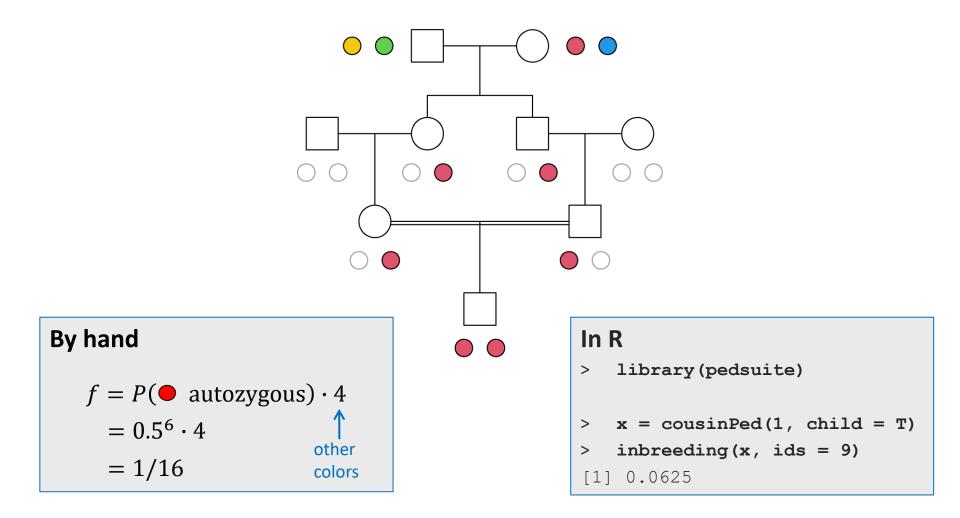


Statistical methods in genetic relatedness and pedigree analysis



#### Inbreding coefficient: Example

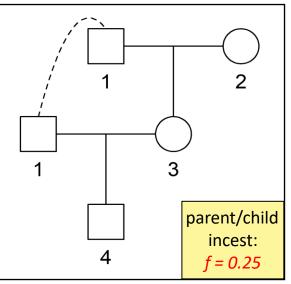






### More kinship & inbreeding coefficients

Relationship	kinship $\boldsymbol{\varphi}$ = $f$ of child	ra eships
Parent-child	1/4 <b>Challe</b>	nge rent relationships The same kinship!
Full siblings	1/4 with	the
Half siblings	1/8	
Grandparent- grandchild	1/8	
Avuncular (uncle/aunt)	1/8	> x =
1st cousins	1/16	ac
2nd cousins	1/64	> kins [1] 0.2
3rd cousins	1/256	> inb: [1] 0.2



- nuclearPed(1, sex = 2) |>ddSon(parents = c(1, 3))
- ship(x, ids = c(1, 3))25
- preeding(x, id = 4)
- 25

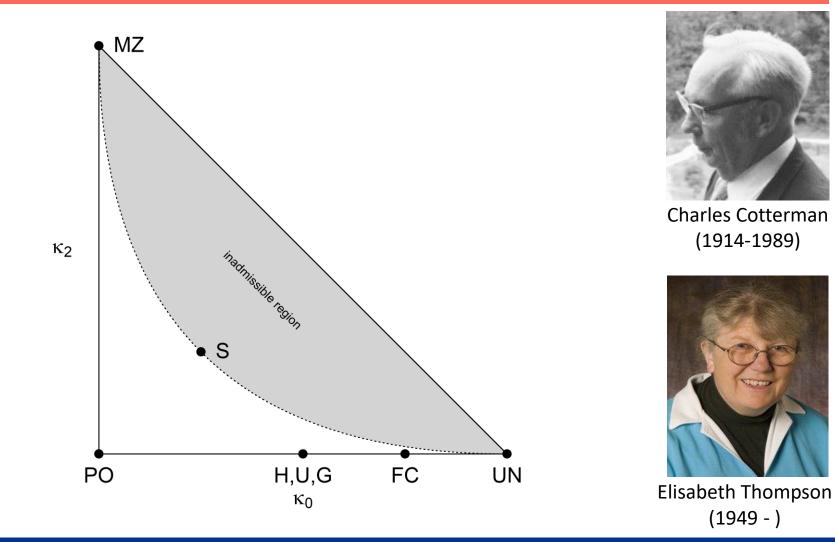








#### **The IBD triangle**

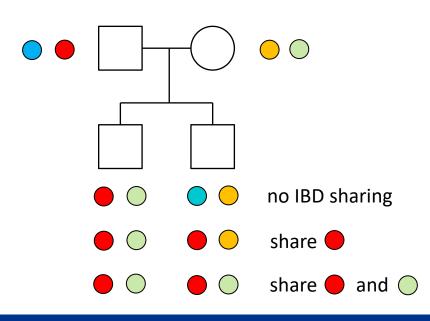






### **IBD coefficients**

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



#### Definition

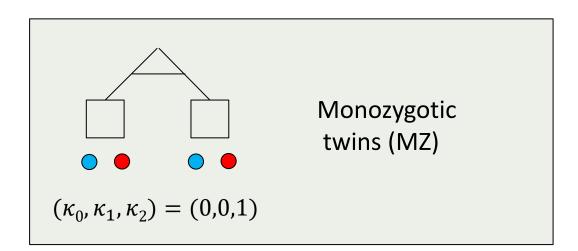
- $\kappa_0 = Pr(0 \text{ alleles IBD})$
- $\kappa_1 = Pr(1 \text{ alleles IBD})$
- $\kappa_2 = Pr(2 \text{ alleles IBD})$

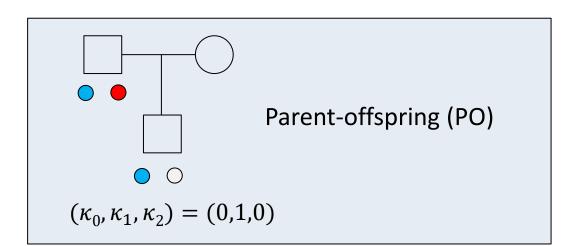
(at random autosomal locus)

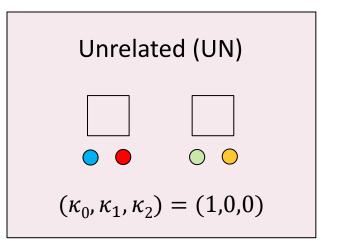




#### **Three trivial relationships**



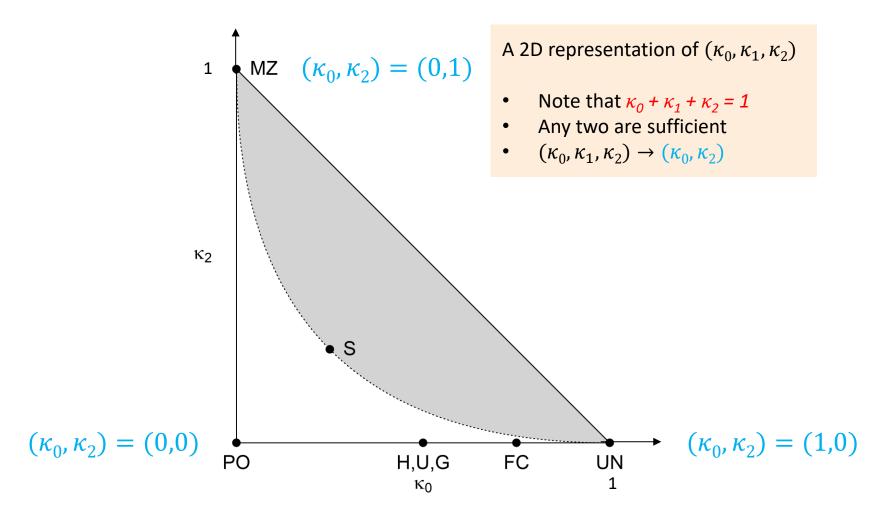






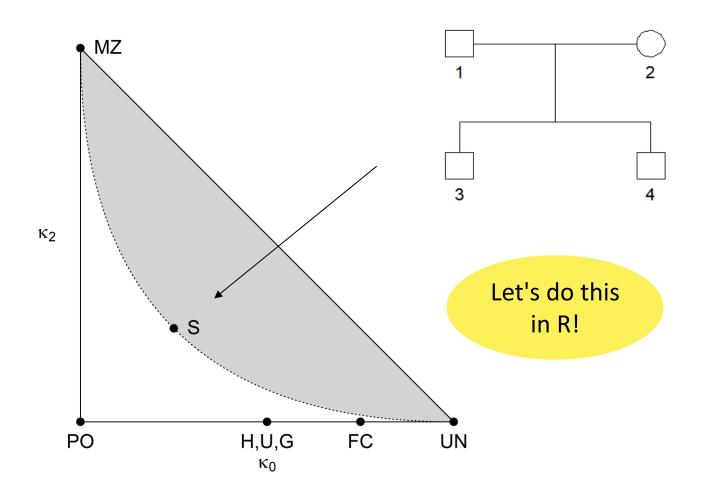


#### The relatedness triangle





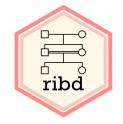
#### What are the coefficients of full sibs

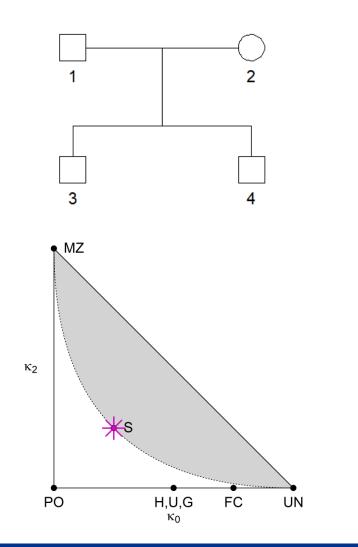






#### ribd: Pedigree-based relatedness coefficients

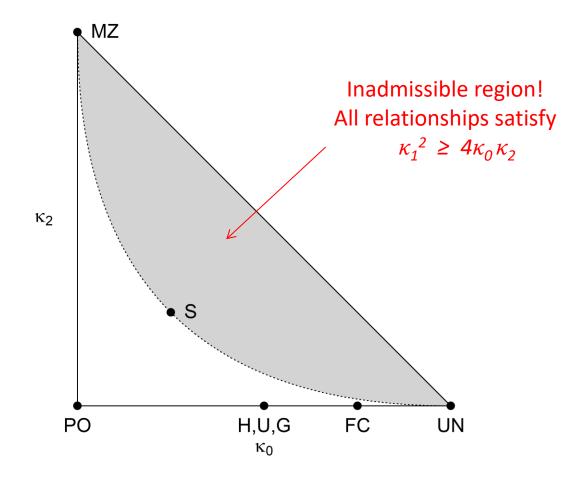




<pre>&gt; library(pedsuite) &gt; x = nuclearPed(2)</pre>							
	<b>kins</b> 0.25	-	ids = 3	:4)			
> 3	kappa	aIBD(x)					
id1	id2	kappa0	kappa1	kappa2			
1	2	1.00	0.0	0.00			
1	3	0.00	1.0	0.00			
1	4	0.00	1.0	0.00			
2	3	0.00	1.0	0.00			
2	4	0.00	1.0	0.00			
3	4	0.25	0.5	0.25			
> :	<b>k</b> = 1	kappaIB	D(x, id:	s = 3:4)			
>	show	InTrian	gle(k)				



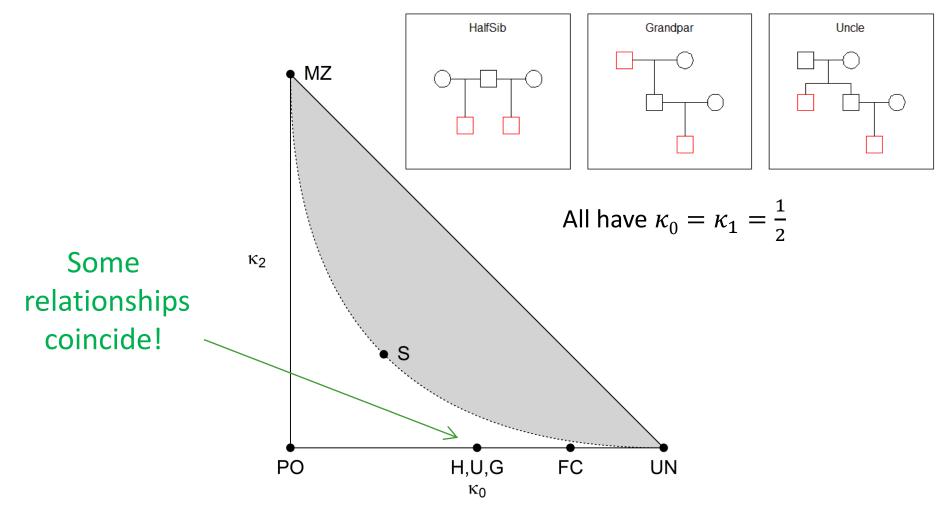
#### The relatedness triangle



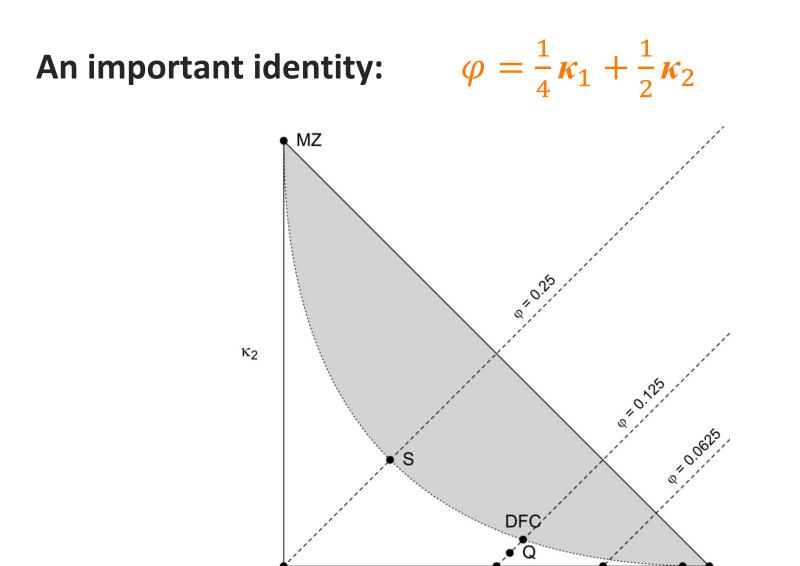




#### The relatedness triangle







H,́U,G

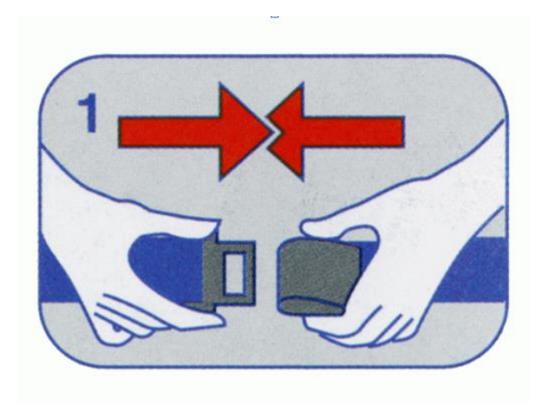
κ<sub>0</sub>

ŕC

SCUN







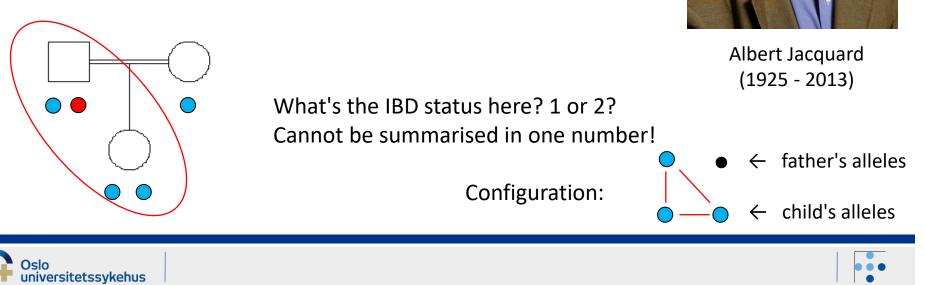


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## Jacquard's identity coefficients

- Jacquard (1970):
  - Structures Génétiques des Populations
- Motivation: Inbred relationships
  - $-\kappa_0, \kappa_1, \kappa_2$  are not well defined
- Example:







### Jacquard's 9 coefficients

9 possible IBD configurations:

• Two individuals, two alleles each:

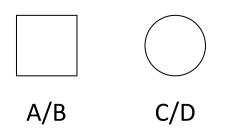
- ← alleles of individual 1
- ← alleles of individual 2

• Any pairwise relationship can be summarised by the relative frequencies of these

Jacquard's *condensed identity* coefficients:  $\Delta_{1,} \Delta_{2}, ..., \Delta_{9}$ 



## If no inbreeding: kappa!



$\square$	•••	arbox.	• •		•••	II	1:	••
0	0	0	0	0	0	κ2	ĸı	κ <sub>0</sub>



● ← alleles of individual 2





 $\uparrow \quad \uparrow \quad \uparrow$ 

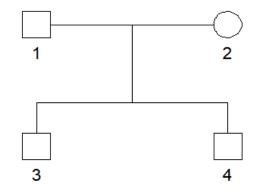
#### Try it out!



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

> plot(x)

>	inbre	eding	(x)
1	2	3	4
0	0	0	0



>	kinship(x)							
	1	2	3	4				
1	0.50	0.00	0.25	0.25				
2	0.00	0.50	0.25	0.25				
3	0.25	0.25	0.50	0.25				
4	0.25	0.25	0.25	0.50				

> 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				

<pre>&gt; identityCoefs(x)</pre>										
id1	id2	D1	D2	D3	D4	D5	D6	D7	D8	D9
1	2	0	0	0	0	0	0	0.00	0.0	1.00
1	3	0	0	0	0	0	0	0.00	1.0	0.00
1	4	0	0	0	0	0	0	0.00	1.0	0.00
2	3	0	0	0	0	0	0	0.00	1.0	0.00
2	4	0	0	0	0	0	0	0.00	1.0	0.00
3	4	0	0	0	0	0	0	0.25	0.5	0.25





#### https://magnusdv.shinyapps.io/quickped/

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#### QuickPed: An Interactive Pedigree Creator

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 income content on line 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.

