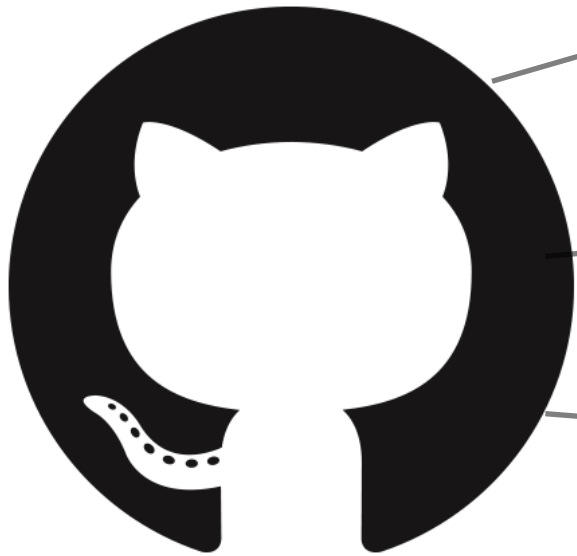


# Leverage proxy-cases in your data!

**Beta version available!**



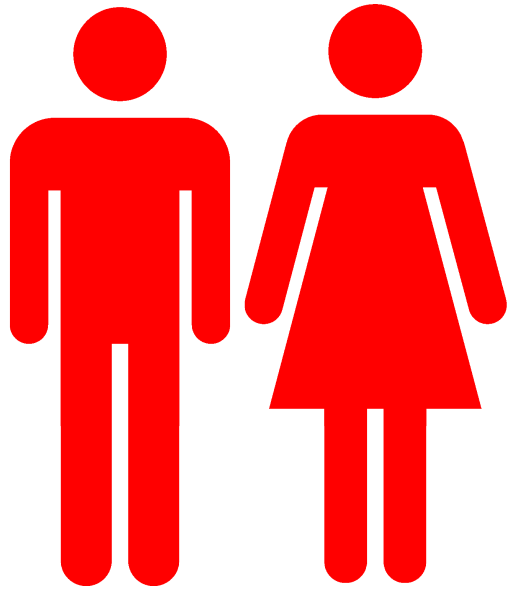
Identify  
unaffected  
relatives of  
cases in your  
data

Perform meta-  
analysis using  
GWAX and  
GWAS with  
shared  
controls

Create  
phenotype  
files for  
BOLT-LMM  
for multiple  
models

<https://github.com/bnwolford/proxyPower>

Proxy-cases are unaffected first degree relatives of cases



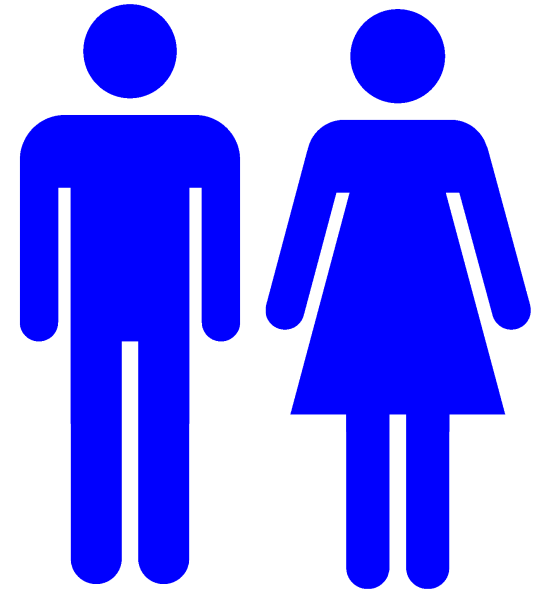
Affected

Case



Unaffected first degree  
relative of affected

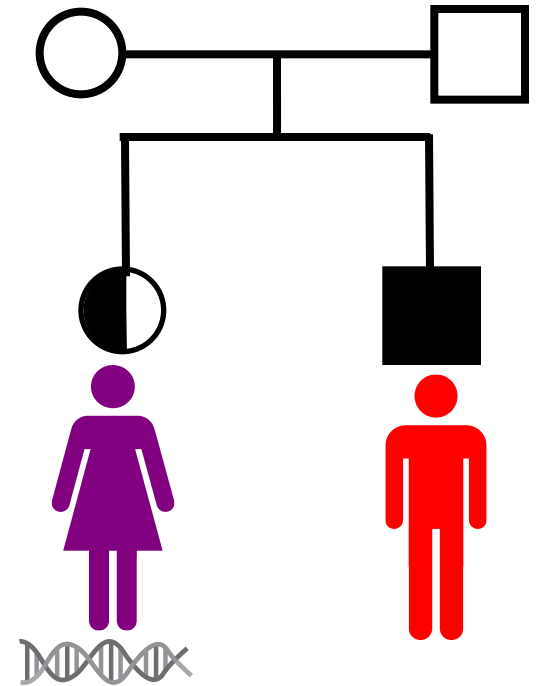
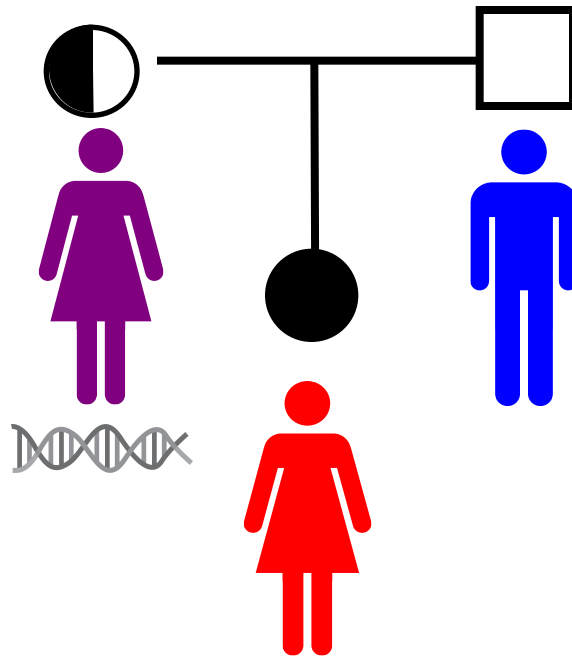
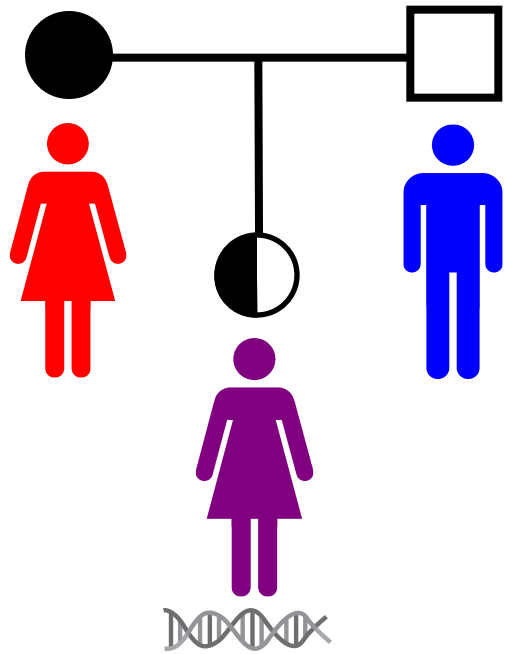
Proxy-case



Unaffected

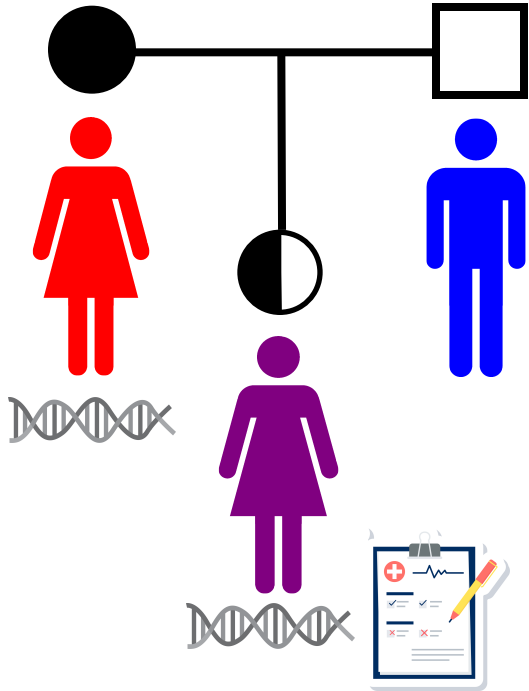
Control

Proxy-case has some combination of affected mother, father, child, sibling depending on epidemiological survey wording



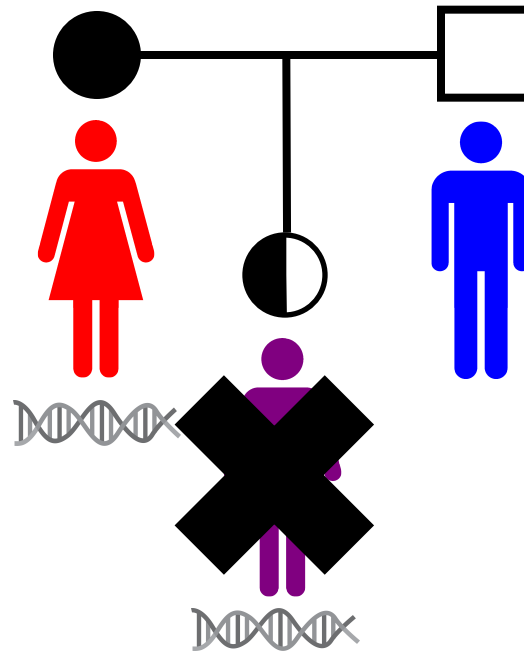
\*double counting genetic info of cases

# Proxy-case definition



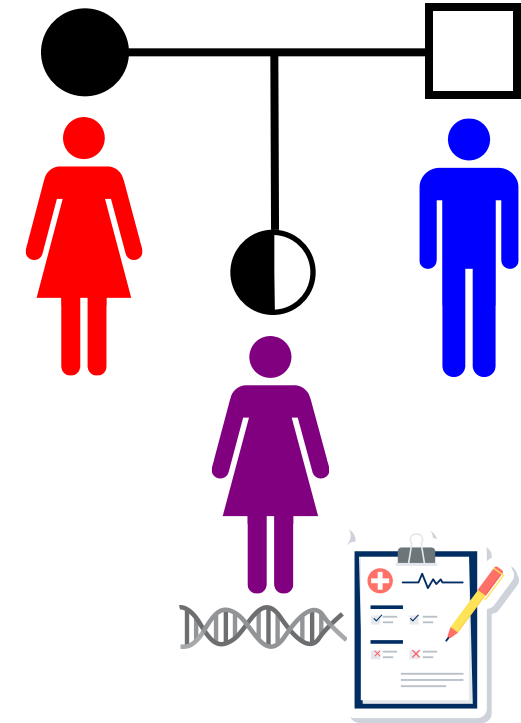
## Self report only:

Identify proxy-cases who report an affected relative on a epidemiological survey\*



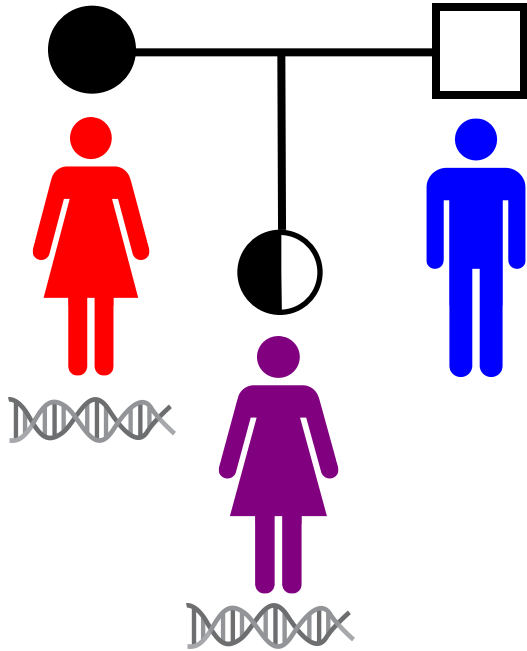
## Self report minus kinship:

Exclude proxy-cases whose affected relative is in dataset, use only proxy-cases whose affected relative is not in dataset

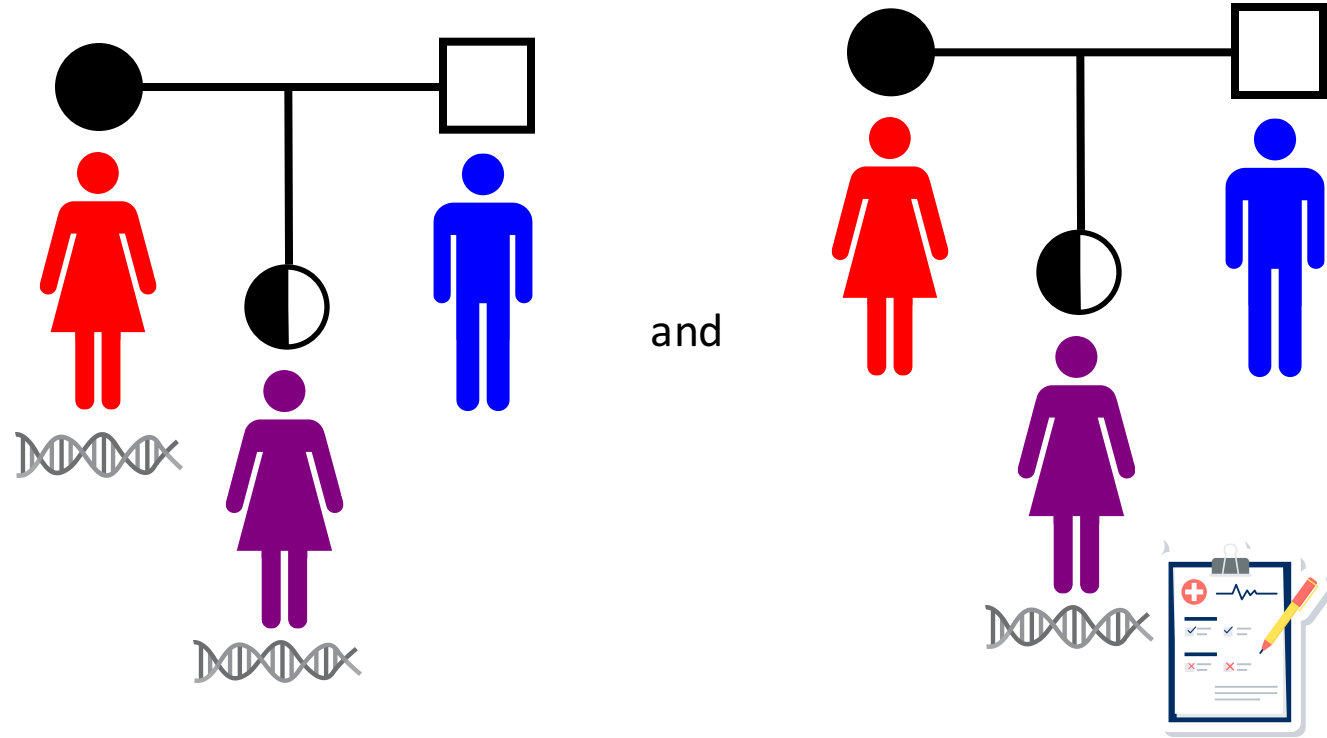


\*double counting genetic info of cases

# Proxy-case definition

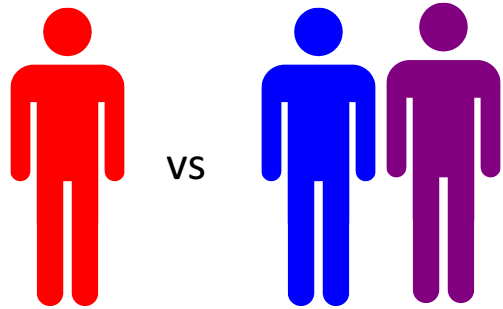


**Kinship only:**  
Identify proxy-cases whose affected relative is in the dataset using kinship matrix



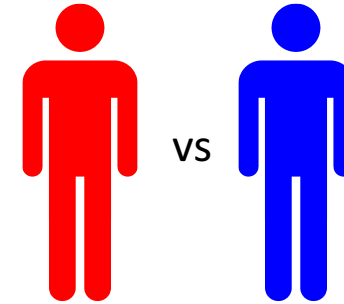
**Self report plus kinship:**  
Identify proxy-cases whose affected relative is in dataset even if the sample did not report having an affected relative in the epidemiological survey\*

# Study designs in cohort-based GWAS



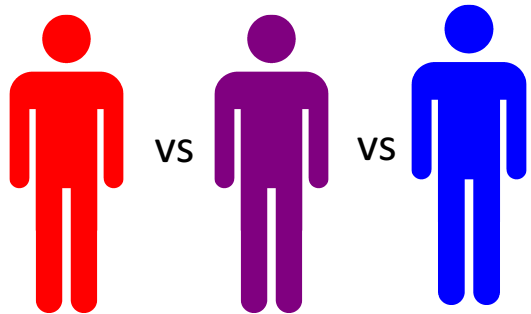
GWAS

1



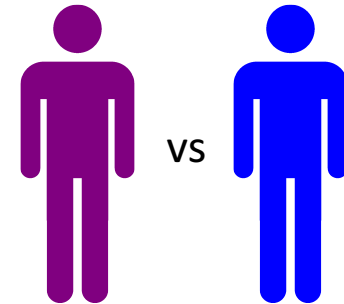
Exclude proxy-cases

2



Model proxy-cases

4

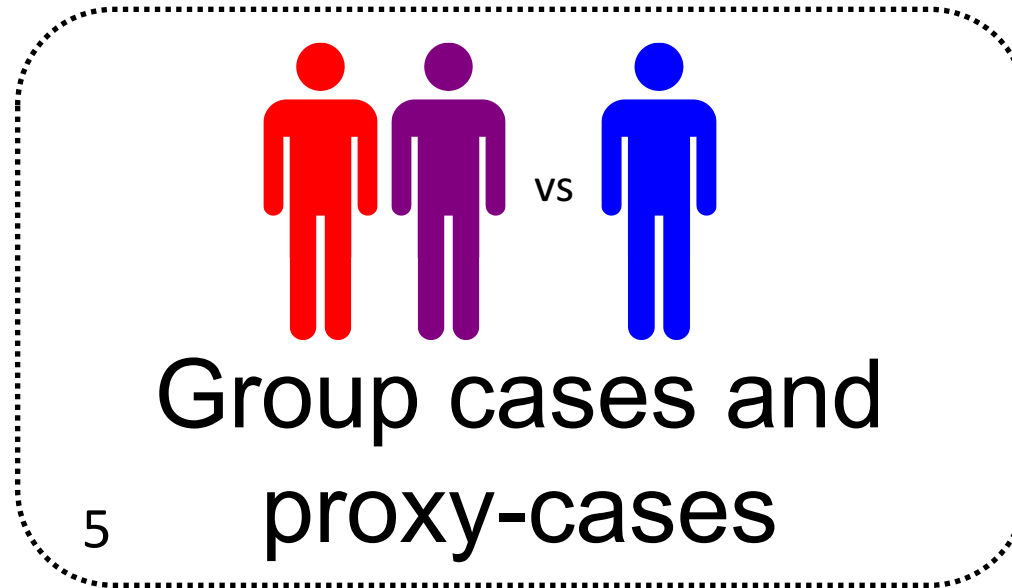


GWAX

*Liu et al, 2017*

3

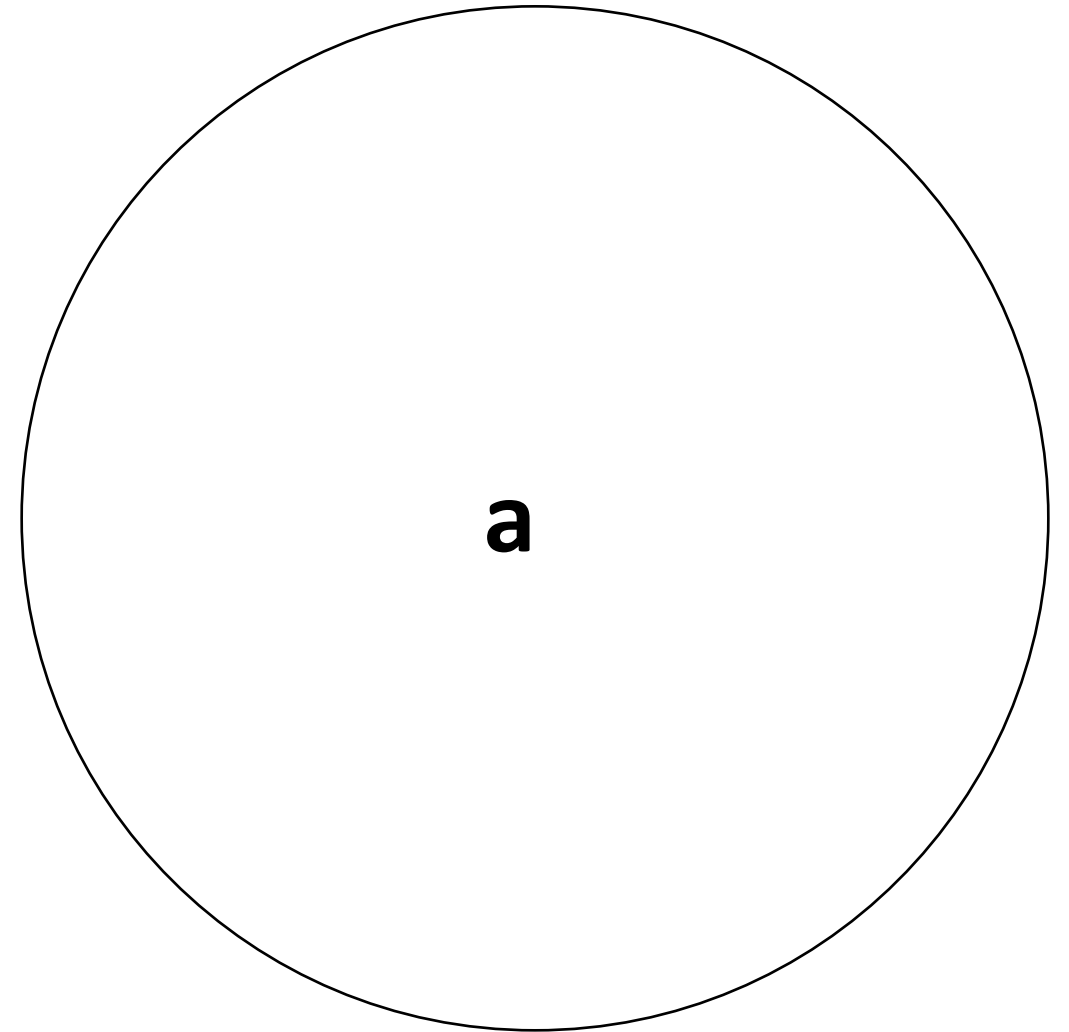
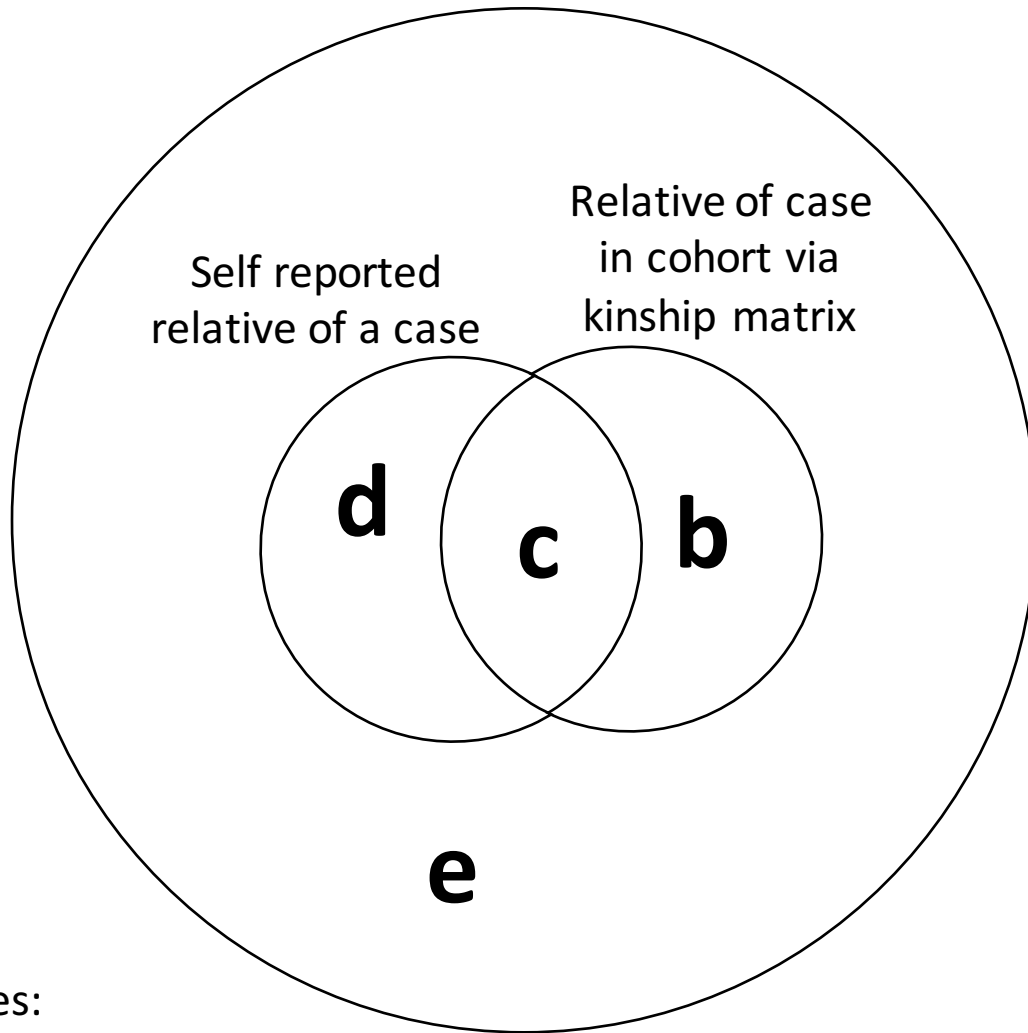
# Study designs in cohort-based GWAS



# Your cohort

**Control**

**Case**



Notes:  
Using first degree relative information, ignoring second degree relatives  
Non conservative control  
Group c is self reported proxy-cases who are relatives of cases in the cohort



Proxy-case definition

		Model			
	Proxy-case	GWAS (1)	Exclude proxy-cases (2)	GWAX (3)	Model proxy-cases (4)
Self report only (SR)	cd	a vs bcde	a vs be	cd vs be	a vs cd vs be
Self report plus kinship (SPK)	bcd	a vs bcde	a vs e	bcd vs e	a vs bcd vs e
Self report minus kinship (SMK)	d	a vs bcde	a vs e	d vs e	a vs d vs e
Kinship only (K)	bc	a vs bcde	a vs de	bc vs cd	a vs bc vs de

Notes:

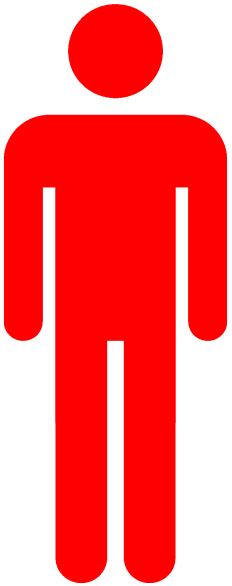
Group a is NA for Model 3

Proxy-case group is NA for Model 2

For SMK definition, group bc is NA for Models 2, 3, 4

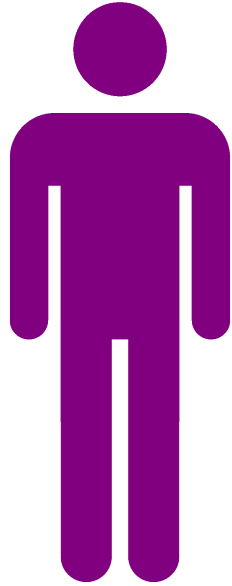
# Modeling coefficient of relationship to a case

$$F = 1$$



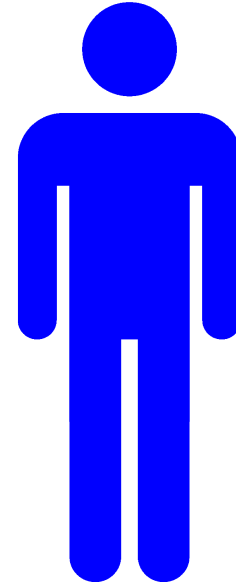
VS

$$F = 0.5$$

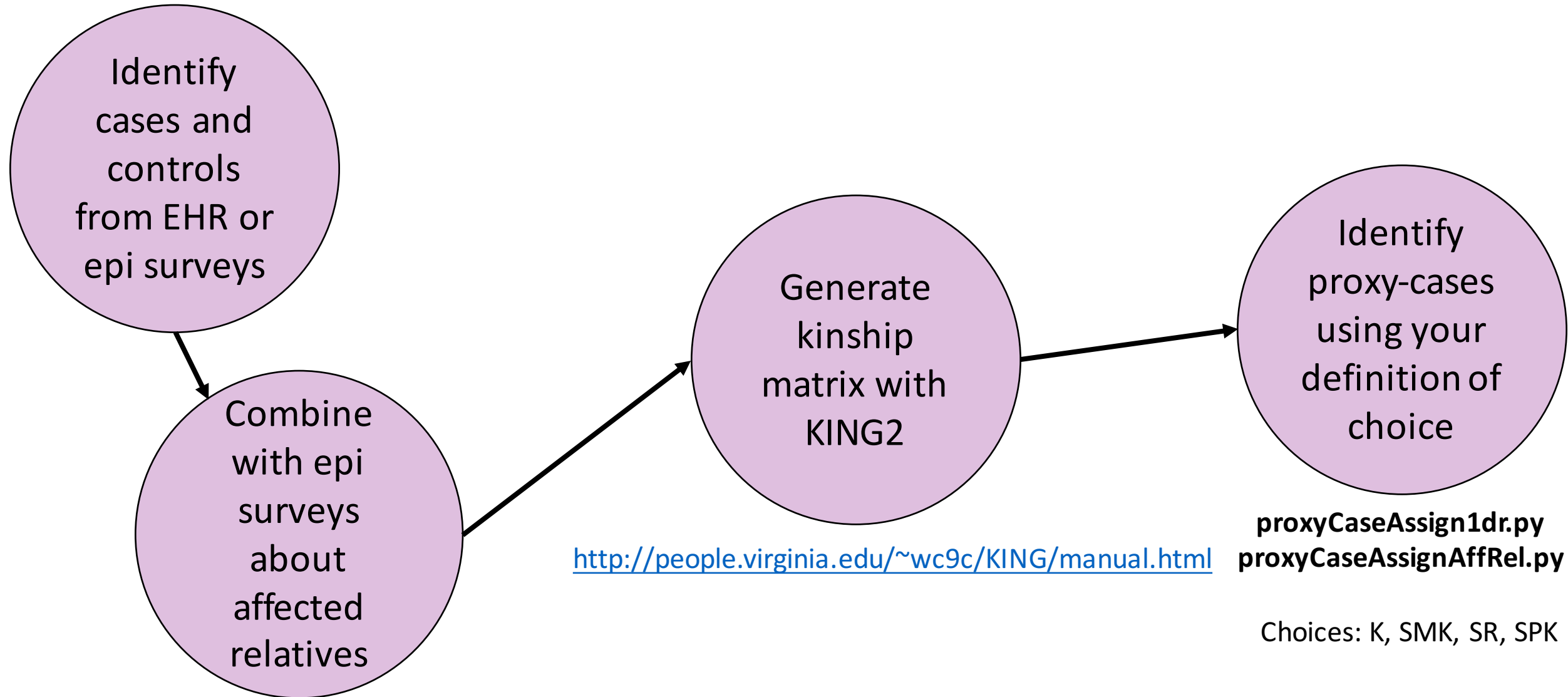


VS

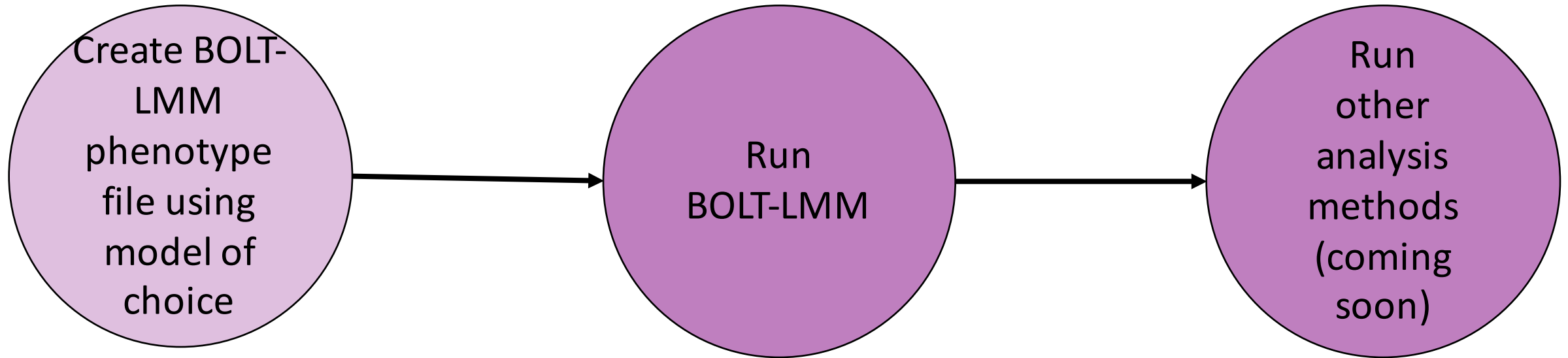
$$F = 0$$



# proxyPower workflow



# proxyPower workflow



**proxyModel.py**

<https://data.broadinstitute.org/alkesgroup/BOLT-LMM/>

Choices: 1, 2, 3, 4, 5