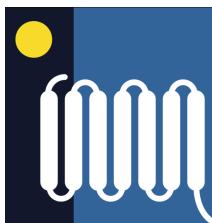


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IMPERIAL

Pedigrees and Risk

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Session Plan

Part 1

- Why draw a pedigree
- How to draw a pedigree

Part 2

- What is risk
- Modification of risk
- How to calculate risk
- Worked examples – mentimeter -



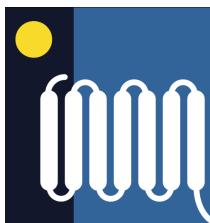
Session Plan

Part 1

- Why draw a pedigree
- How to draw a pedigree

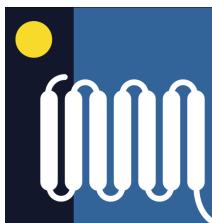
Part 2

- What is risk
- Modification of risk
- How to calculate risk
- Worked examples -



Why draw pedigree

- Provides a clear simple summary of information
- Able to spot patterns easily (quiz examples added to modes insendi)
- Explain pattern to patient
- Identify potential carriers of risk gene
- Calculate risk of passing on disease or being a carrier
- Allows informed choice



Why draw pedigree

- Why not test for disease gene
- Many diseases mix of familial and sporadic
- Many causative gene isn't known



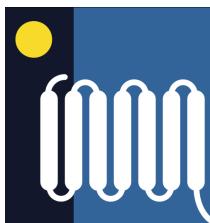
Why draw pedigree

- Amyotrophic lateral sclerosis (ALS, motor neurone disease)



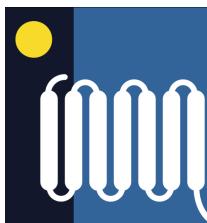
~85% sporadic – cause unclear

~15% familial – four known genes account for 65% of cases



Why draw pedigree

- Why not test for disease gene
- Many diseases mix of familial and sporadic
- Many causative gene isn't known
- Impossible to test everyone for disease genes
- When disease exclusively familial and mutation known many do not want test



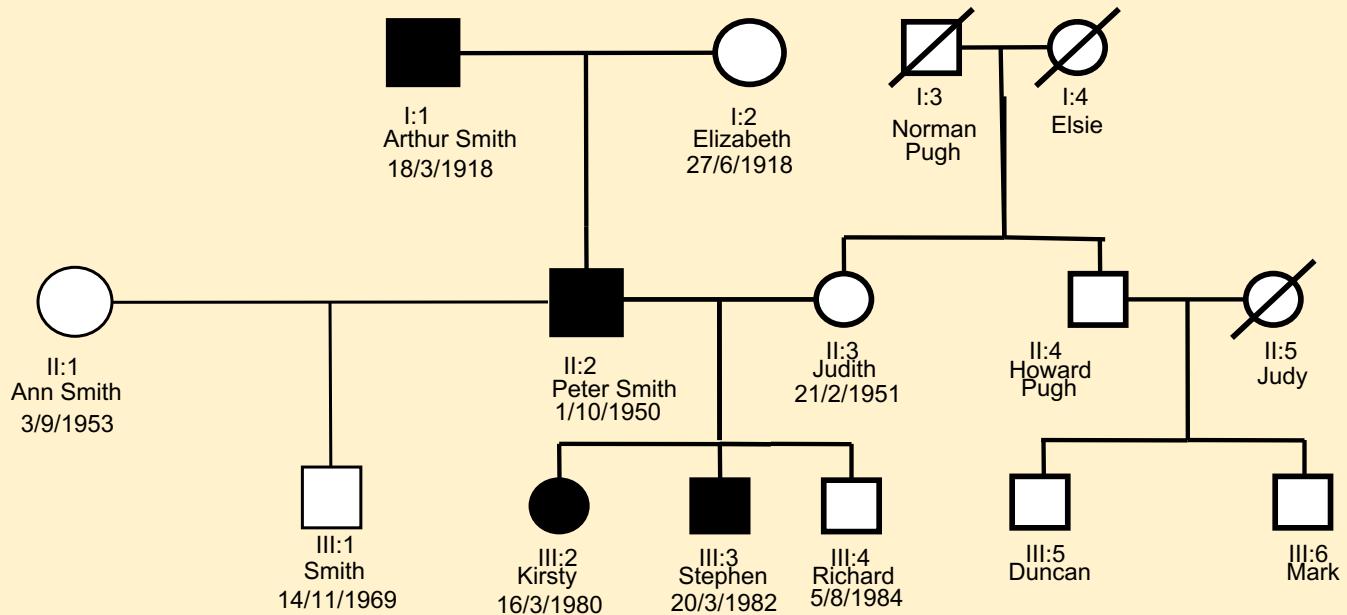
Why draw pedigree

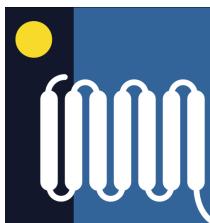
- Why not test for disease gene
- Many diseases mix of familial and sporadic
- Many causative gene isn't known
- Impossible to test everyone for disease genes
- When disease exclusively familial and mutation known many do not want test
 - >80% of people with 50% risk of inheriting Huntingdon's disease decline test



How to draw a pedigree

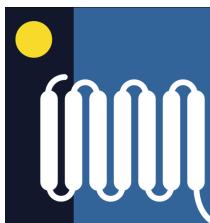
- Start at the bottom with the proband and siblings
- Choose one parent ask about their siblings, children and parents
- Add other side of family
- Ask about children of other partners





Potential difficulties

- Incomplete information
 - May not have information on all or many relatives
- Incorrect information
- Family history may not be correct
- Important in clinical setting



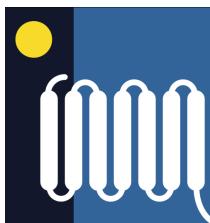
Progress check

Part 1

- Pedigree allows identification of inheritance type if any
- Needed to calculate risk

Part 2

- Risk helps informed decision making
- Can reassure some patients if not affected
- Identify patients who require further support
 - Genetic counselling



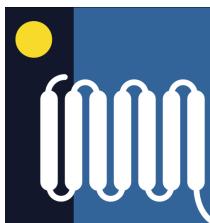
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- Why draw a pedigree
- How to draw a pedigree

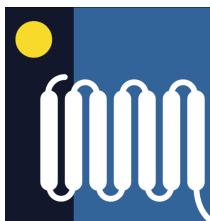
Part 2

- What is risk
- Modification of risk
- How to calculate risk
- Worked examples mentimeter (1937 0915)-



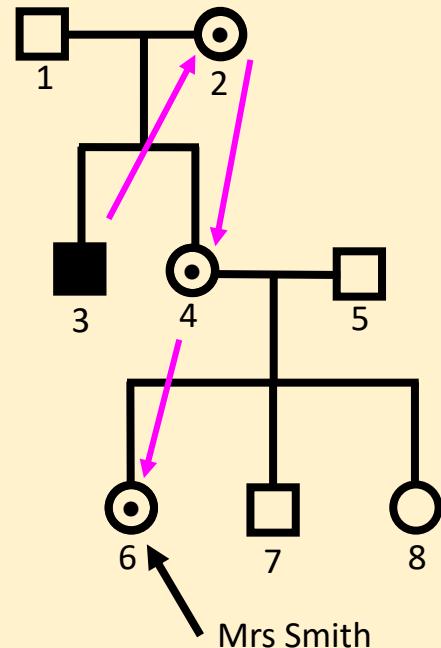
Risk

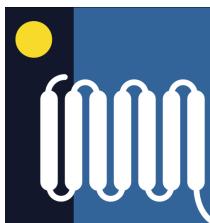
- This is a calculation of the predicated chance of having the disease or being carrier.
- Work from person with known phenotype to subject
- If more than 1 affected individual then start closest relative on each side of family
- Calculate risk for each person on path from the start point to the subject
- Multiply risks together



Risk

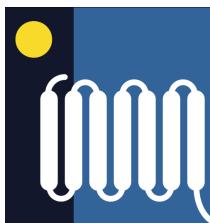
- This is a calculation of the predicated chance of having the disease or being carrier.
- **Mrs Smiths maternal uncle (3) has X linked recessive condition**
- **What is chance Mrs Smith (6) is a carrier**
- **Work from person with known phenotype to subject**
- 2 must be carrier 1 in 1 chance
- Since 2 is carrier 1 in 2 chance 4 inherited mutation
- Assume 4 carrier therefore 1 in 2 chance Mrs Smith (6) inherited mutation
- Multiply risks together $1 \times 1/2 \times 1/2$
- Therefore 1 in 4 chance Mrs Smith (6) carrier





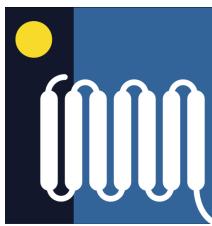
Risk

- Take account of all information
 - Phenotype
 - Disease characteristics
- Family distribution for X-linked and mitochondrial disease
 - which side of family disease is on
 - Which parent has the disease



Risk modifiers

- Which side of family disease is on
 - X-linked, mitochondrial
- Ethnic background
 - Many diseases have different prevalence in different populations eg CF, Sickle cell, Tay-Sachs
 - Heterozygote advantage- sickle cell, CF?
 - Founder effect- Tay-Sachs



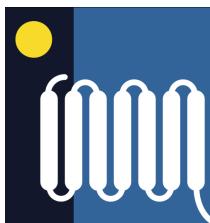
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Risk modifiers

On which continent is the area with the highest prevalence of Sickle cell trait

- Europe
- America
- Africa
- Asia



Risk modifiers

On which continent is the area with the highest prevalence of Sickle cell trait

Europe in parts of Greece 1 in 10

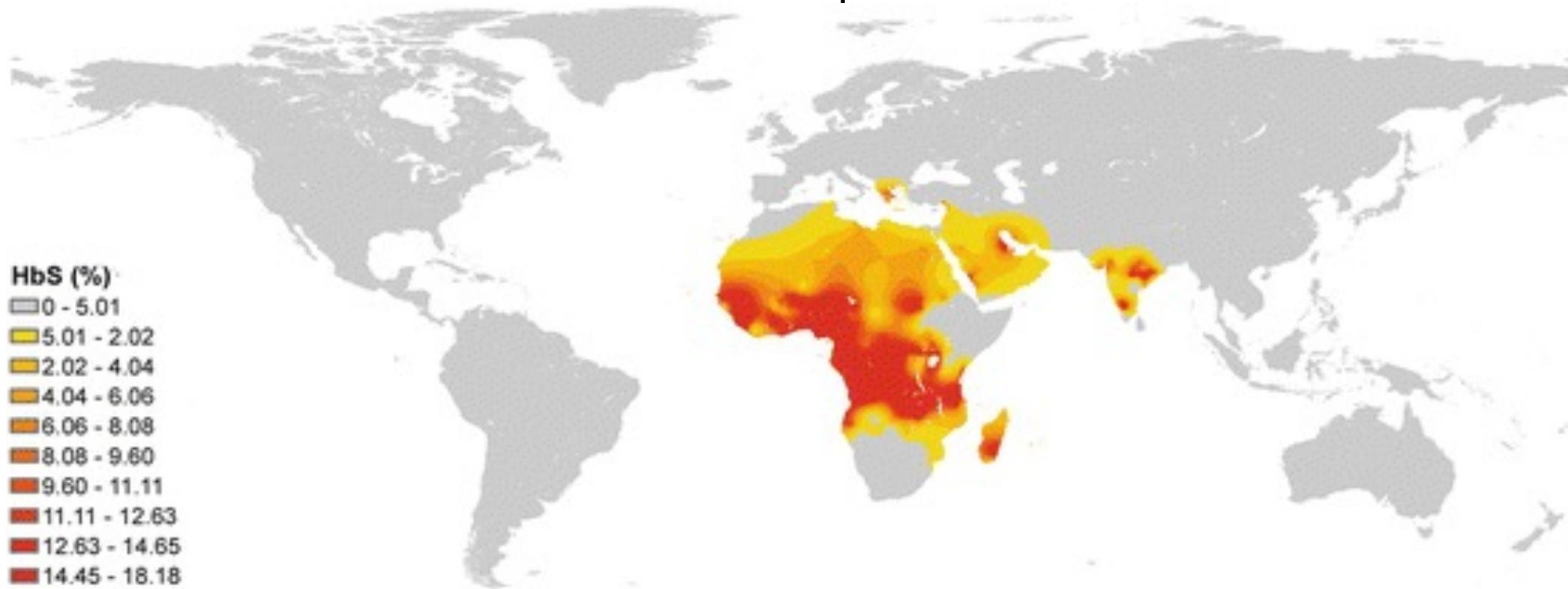
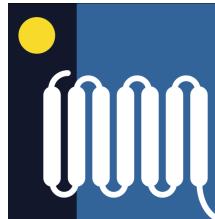
America

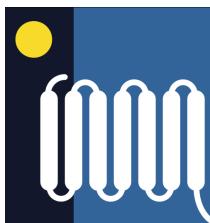
Africa- 1 in 5 in equatorial Africa (Nigeria, Ghana), less than 1 in 100 southern Africa

Asia- certain tribal areas of India as high as 1 in 2

80% of people affected by sickle cell anaemia sub-Saharan African heritage

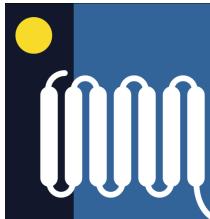
Sickle cell trait distribution



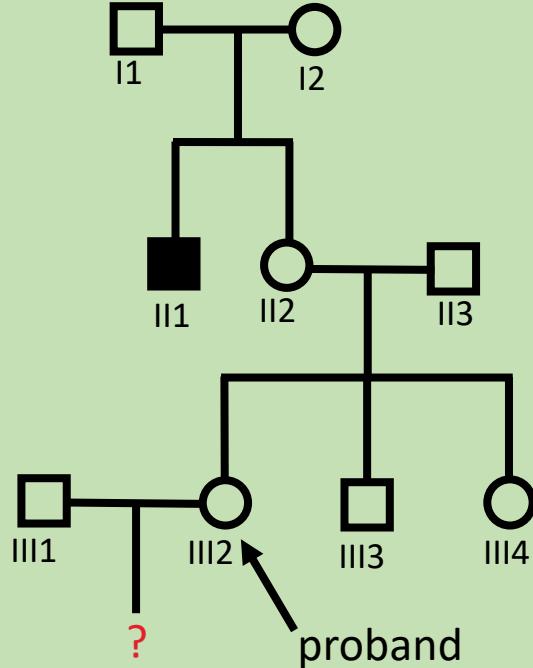


Risk modifiers

- Which side of family disease is on
 - X-linked, mitochondrial
- Ethnic background
 - Many diseases have different prevalence in different populations eg CF, Sickle cell, Tay-Sachs
 - Heterozygote advantage
 - Founder effect
- Information you know about the person
 - Their phenotype
 - Biological sex



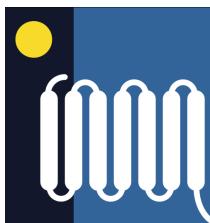
Question Time:1



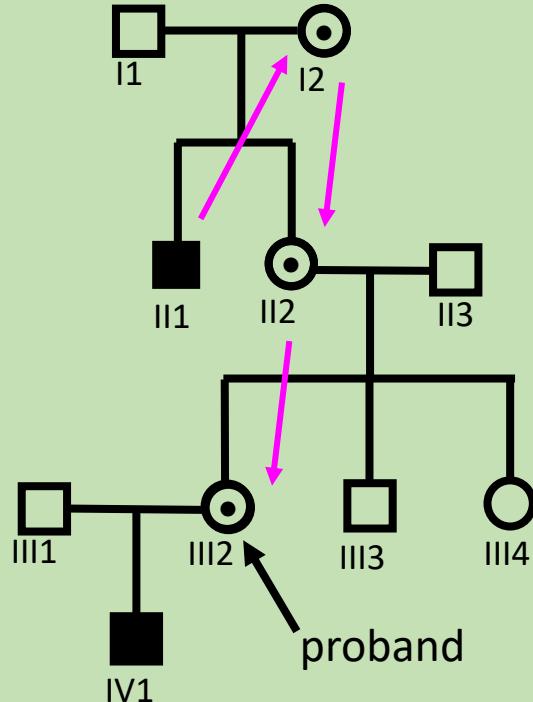
Mrs Smith consults you
Her maternal uncle has Haemophilia A an X-linked recessive disease (incidence 1 in 5000)

Want to know risk of having affected son?

- a. 1 in 4
- b. 1 in 8
- c. 1 in 16
- d. 1 in 2
- e. Population risk

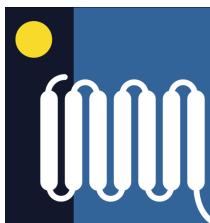


Question Time:1

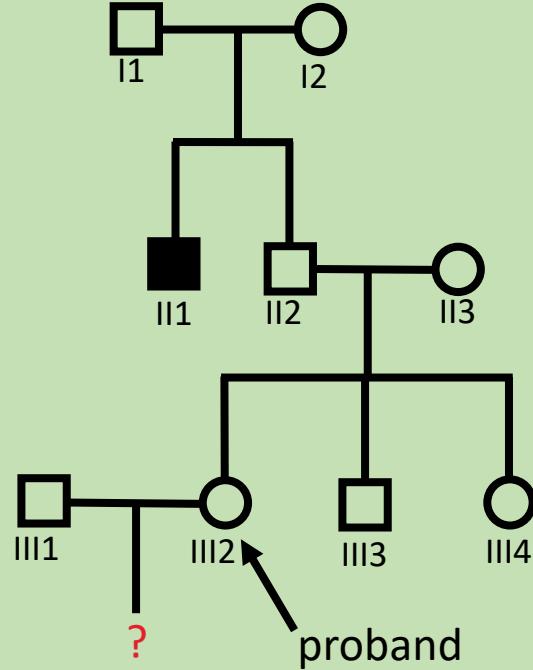


Mrs Smith consults you
Her maternal uncle has Haemophilia A an X-linked recessive disease (incidence 1 in 5000)
Want to know risk of having affected son? 1 in 16

Maternal grandmother must be carrier
1 in 2 chance mother carrier
If mother is carrier 1 in 2 chance she inherited gene
Therefore 1 in 4 chance she is carrier
If she is carrier 1 in 2 chance of passing gene onto son
However only 50% children male therefore 1 in 16
Also has 1:16 chance having carrier daughter



Question Time:2

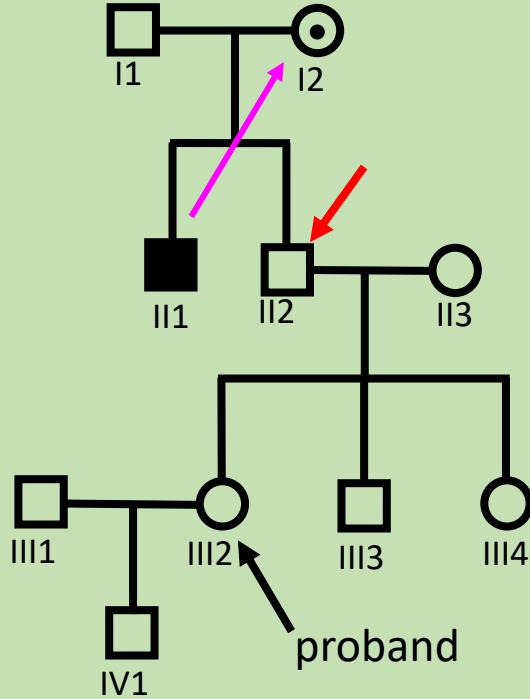


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Want to know risk of having affected son?

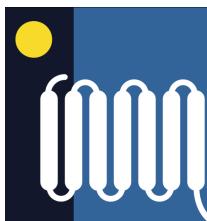
- a. 1 in 4
- b. 1 in 8
- c. 1 in 16
- d. 1 in 2
- e. Population risk

Question Time:2

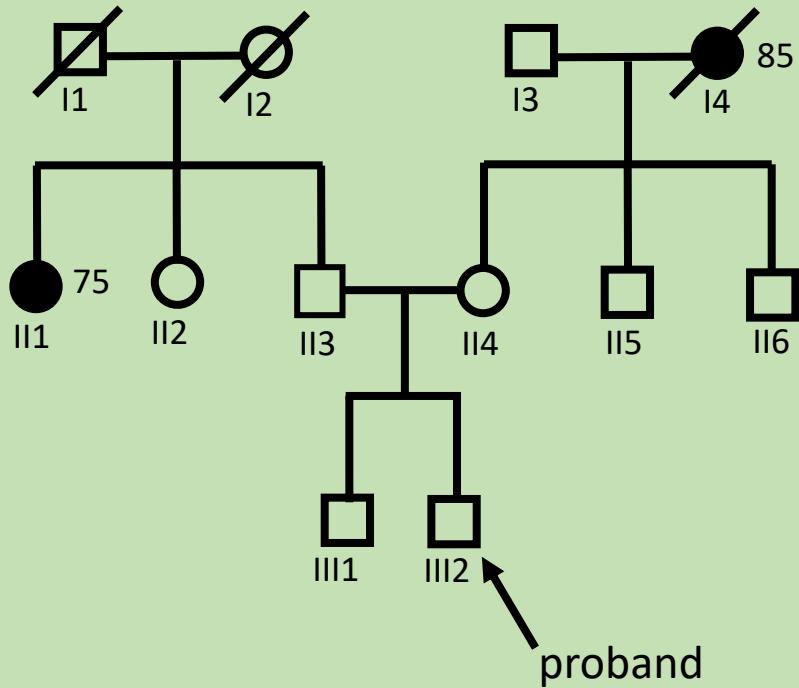


Mrs Smith consults you
Her paternal uncle has Haemophilia A an X-linked recessive disease (incidence 1 in 5000)

Want to know risk of having affected son? Population risk 1 in 5000
X-linked
Paternal grandmother carrier
father not affected
No one affected in maternal family



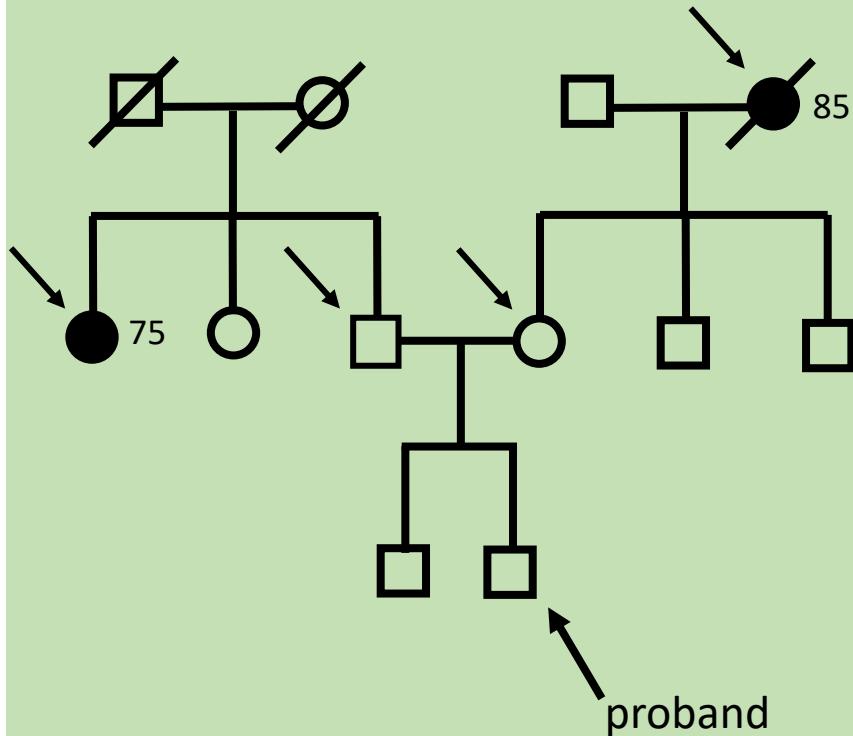
Question Time:3



Mr Jones (age 30) consults you regarding Alzheimer's disease
Mr Jones is worried as he has recently read about an inherited form of AD
AD (~3% familial ~97% sporadic) overall prevalence ~1.6%
Age onset FAD 40-50
FAD autosomal dominant disease
His paternal aunt who is now 80 developed symptoms of dementia at 75
His maternal grandmother developed symptoms at 85
No one else affected (mother is 58; father 60)
Worried he may have disease

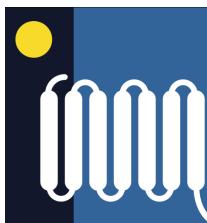
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Question Time:3

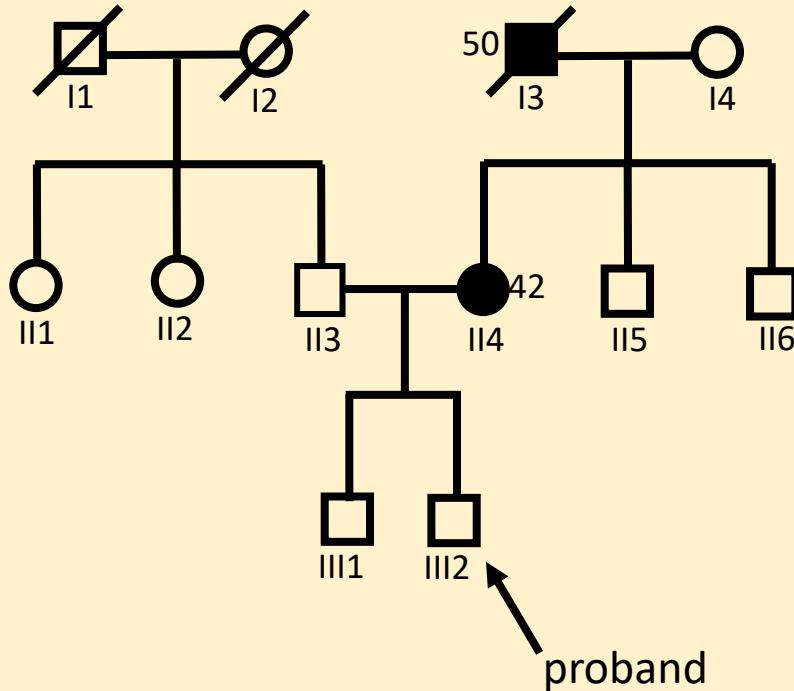


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No one else affected (mother is 58; father 60)
Worried he may have disease

Risk is population risk
Autosomal dominant neither parent affected
Relatives likely to have sporadic form – age of onset, no family affected

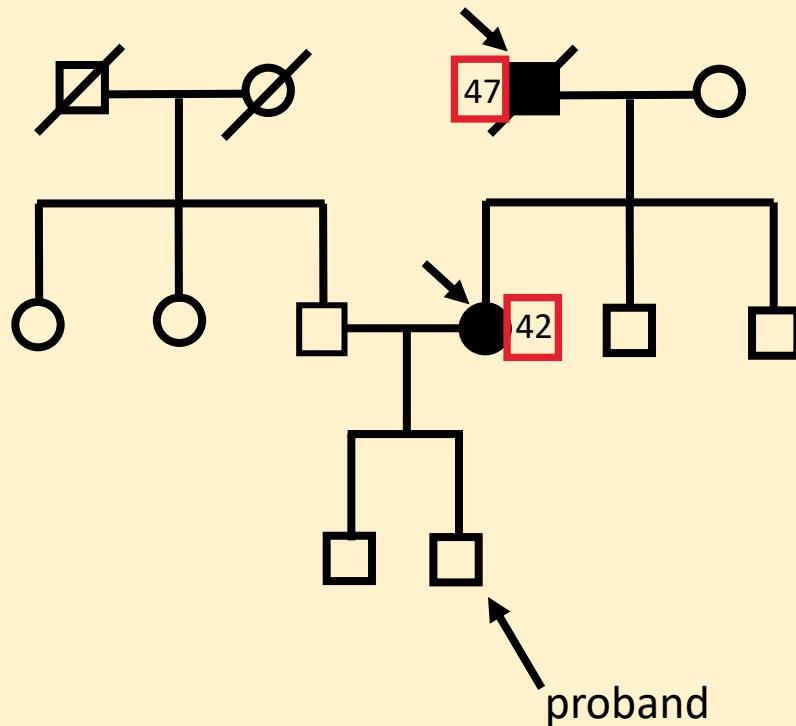


Question Time:4



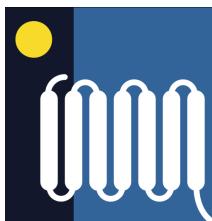
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Mr Jones is worried as he has recently read about an inherited form of AD
AD (~3% familial ~97% sporadic) overall prevalence ~1.6%
Age onset FAD 40-50
FAD autosomal dominant disease
His mother who is now 50 developed symptoms of dementia at 42
His maternal grandfather developed symptoms at 50
Neither of his two maternal uncles who are now in their late 60s are affected
No one on his fathers side is affected.
Worried he may have inherited gene and develop disease

Question Time:4

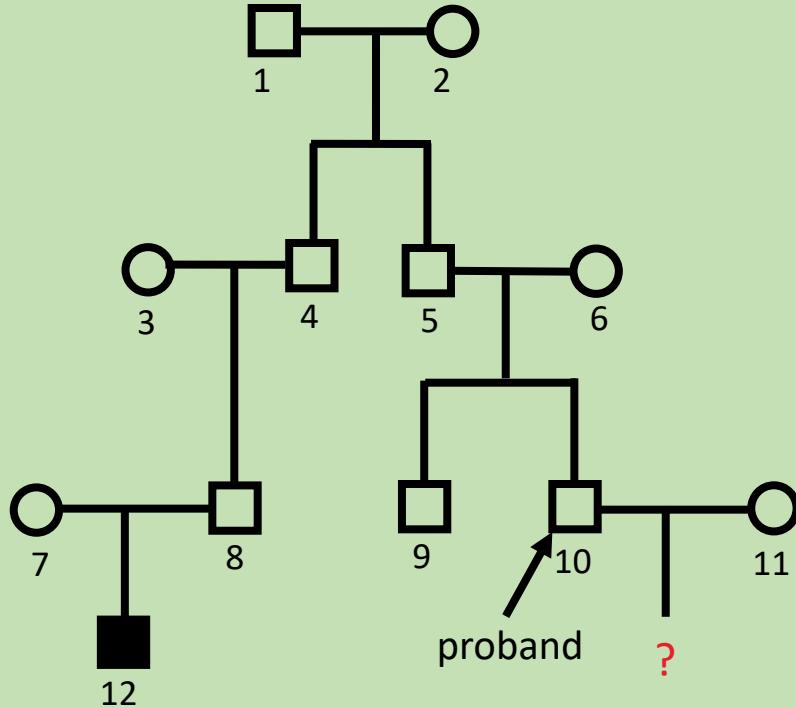


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 His mother who is now 50 developed symptoms of dementia at 42
 His maternal grandfather developed symptoms 47
 Worried he may have inherited gene develop disease

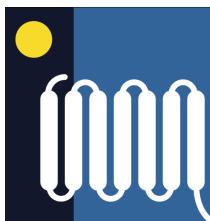
Autosomal dominant
 parent and grandparent affected
 Early age of onset
 Possibly FAD
 Mother may be affected therefore 1 in 2 (50%)



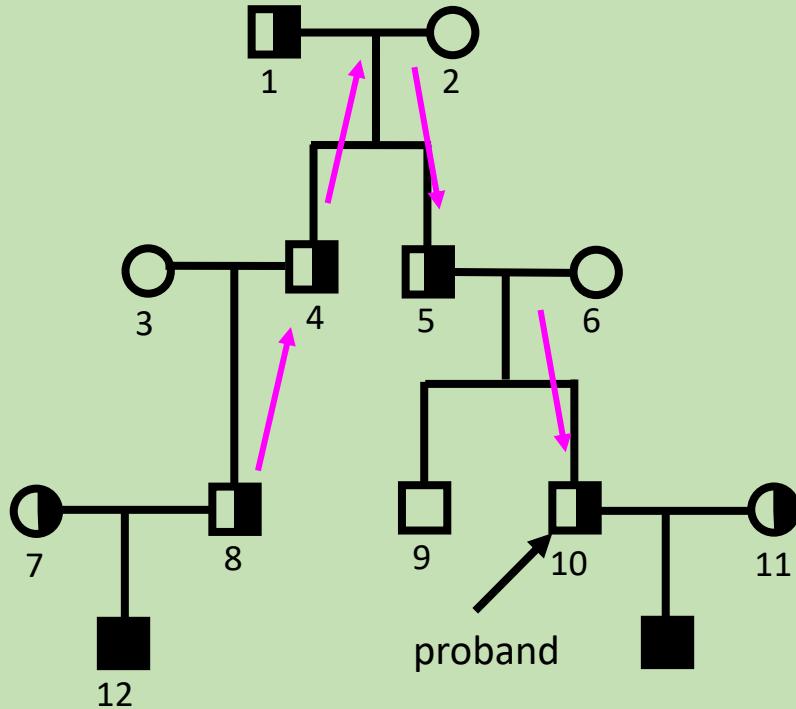
Question Time:5



Mr & Mrs Taylor consult you both of NW European descent
Mr Taylor's first cousin has a son diagnosed with CF
No one with disease in Mrs Taylors family
They are thinking of starting a family and want to know risk of having child with CF
CF autosomal recessive
1 in 22 NW European descent carriers



Question Time:5



Maximum risk of child with CF 1 in 704

7 and 8 must both be carriers

Either 3 or 4 must be carrier

Chance its 4 is 1 in 2

If 4 carrier 1 or 2 must be a carrier

Chance 5 inherited gene 1 in 2

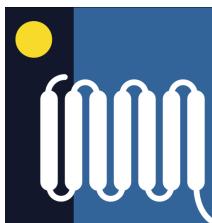
If 5 carrier chance 10 (Mr Taylor) inherited 1 in 2

Thus overall risk Mr Taylor carrier 1 in 8 ($1/2 \times 1/2 \times 1/2$)

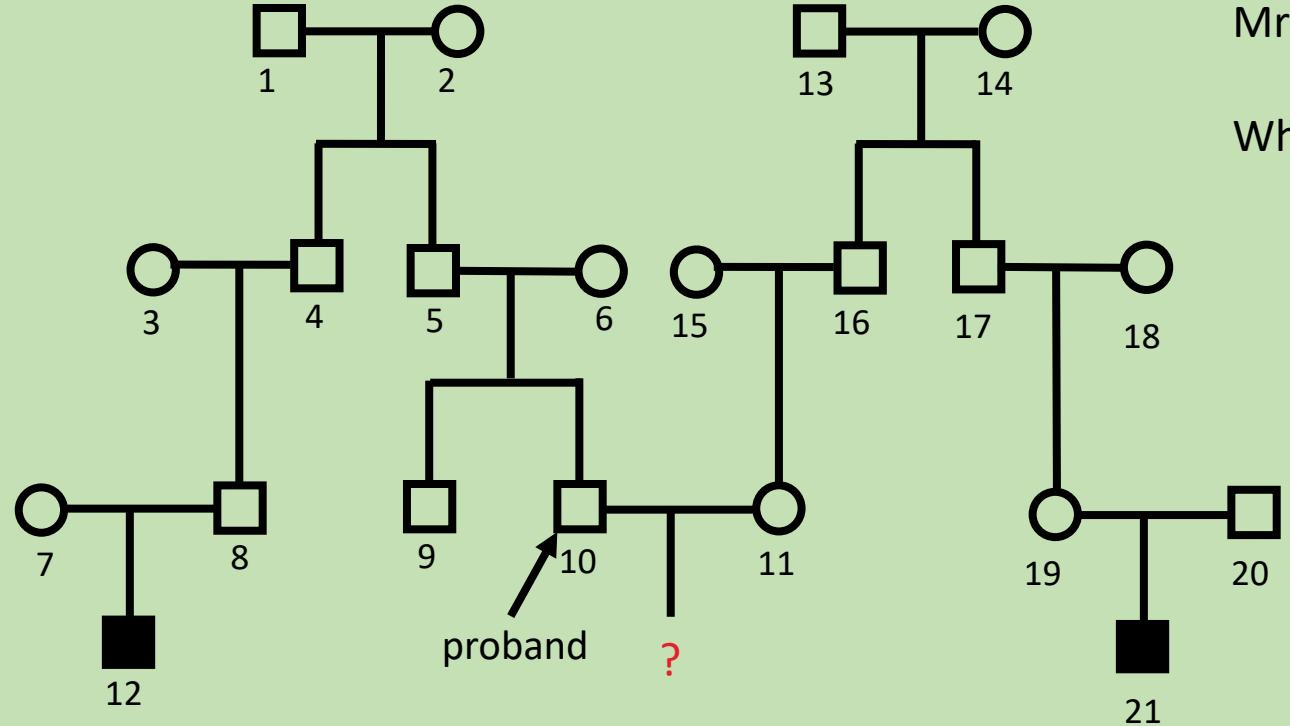
Mrs Taylor population risk of carrier 1 in 22

Chance of child being affected if both carriers 1 in 4

Thus overall chance $1/22 \times 1/8 \times 1/4 = 1 \text{ in } 704$

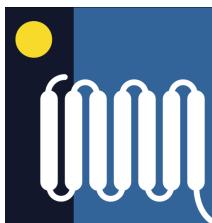


Add information from other side of family

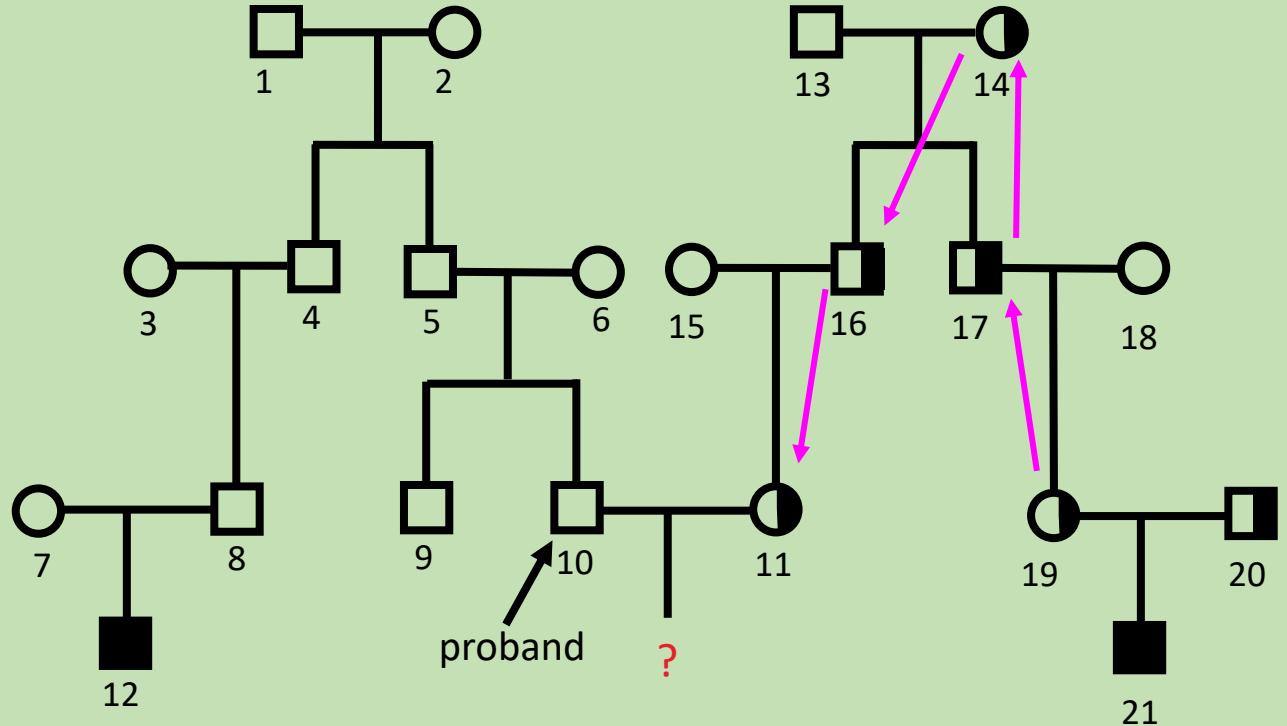


Mrs Taylors first cousin has now had a child with CF

What is the risk of Mrs and Mr Taylor having an affected child

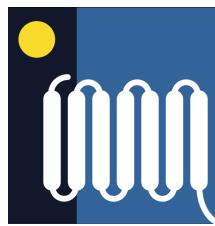


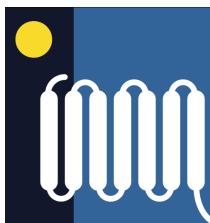
Add information from other side of family



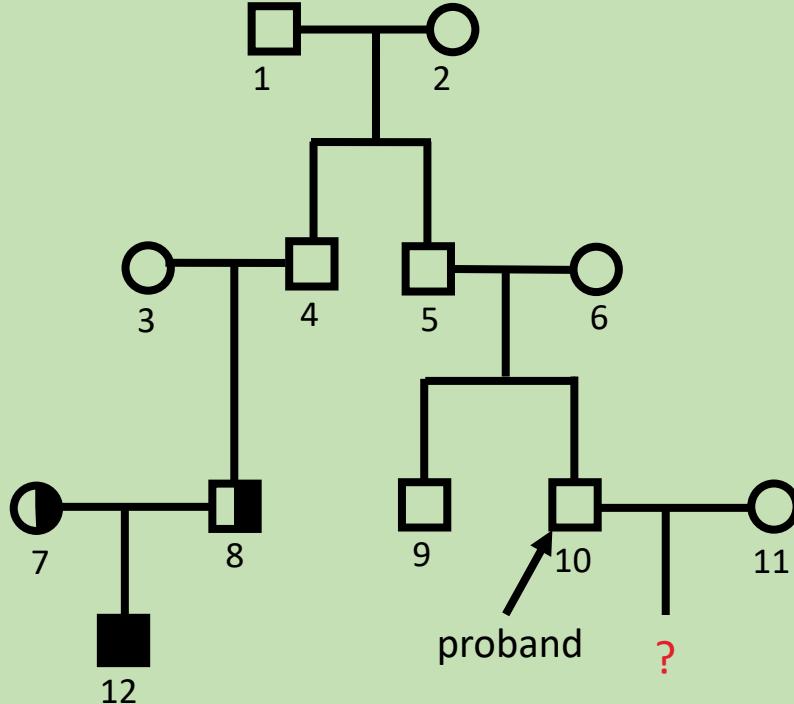
20 and 21 must both be carriers
Either 17 or 18 must be carrier
Chance its 18 is 1 in 2
If 17 carrier 13 or 14 must be a carrier
Chance 16 inherited gene 1 in 2
If 16 carrier chance 11 (Mrs Taylor) inherited 1 in
Overall risk Mrs Taylor carrier 1 in 8 ($1/2 \times 1/2 \times 1/2$)
Mr Taylors risk 1 in 8
Therefore risk both carriers $1/8 \times 1/8$ so 1 in 64
Risk of having affected child is $1/64 \times 1/4$ so 1 in 256

How more information modifies risk

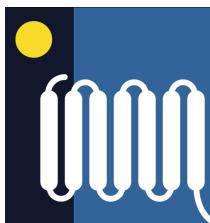




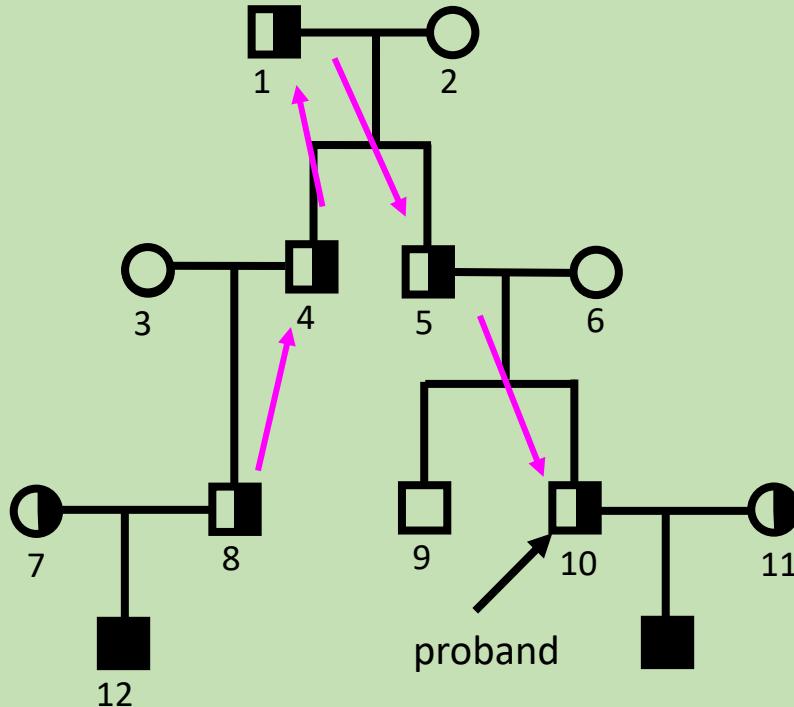
More information and risk



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Mr Taylor's first cousin has a son diagnosed with CF
No one with disease in Mrs Taylors family
They are thinking of starting a family and want to know risk of having child with CF
CF autosomal recessive
1 in 22 NW European descent carriers
What is risk if
3 carrier
4 carrier
5 carrier
10 carrier
10 and 11 carrier



How could risk of affected child risk be modified



If we know 3 is carrier risk becomes effectively zero

If we know 4 is carrier risk becomes 1 in 352 (1 or 2 must be carrier)

If we know 5 is carrier risk becomes 1 in 176

If we know 10 is carrier risk becomes 1 in 88

If we know 10 and 11 are carriers risk becomes 1 in 4

Thus the more information the better



Question Time:6

Couple who are both known to be carriers of recessive disease (Aa)
Have unaffected child want to know risk of them being a carrier (Aa)
A- normal allele a- disease allele

Parent Allele	A	a
A	AA	Aa
a	Aa	aa

- a. 1 in 4
- b. 1 in 3
- c. 1 in 2
- d. 2 in 3
- e. 3 in 4



Question Time:6

Couple who are both known to be carriers of recessive disease (Aa)

Have unaffected child want to know risk of the child being a carrier (Aa)

Parent Allele	A	a
A	AA	XX
a	XX	XX

2/3 Since you know child not affected remove one possibility

If you know child not affected nor carrier it is AA

But also because not affected (aa) nor carrier (aA)

Bayesian statistics

Hereditary haemochromatosis



“So THAT'S why you're feeling tired all the time! The thousands of patients told they're healthy when they really have genetic condition which can damage your liver and even cause kidney failure”

Daily Mail

Killer haemochromatosis is ‘most common genetic disease’ – often mistaken for extreme tiredness

The Sun

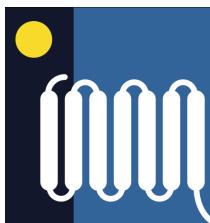
Haemochromatosis ‘bigger threat than we thought’
BBC

How a simple blood test got me thinking about our behaviour and choices
Guardian

‘Stealth disease’ written off as old age may be behind thousands of cancers and joint problems, study shows

Genetic disorder dubbed ‘Celtic curse’ afflicts one in five men who carry mutation with liver disease or joint issues
Independent

Hereditary haemochromatosis



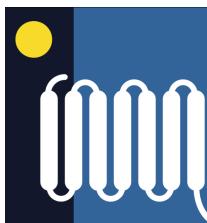
Hereditary (Familial) Haemochromatosis autosomal recessive disease

Mutation in Human homeostatic iron regulator protein (HFE)

Result excess iron absorption. Humans cannot excrete excess iron so builds up in tissues

Wide range of symptom including tiredness

Hereditary haemochromatosis – a common cause of tiredness ?



Prevalence

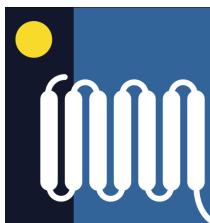
HH approximately 1 in 200 northern Europeans homozygous 0.5%

Of these only 10% have clinically relevant iron accumulation 0.05%

Chronic tiredness/fatigue main symptom of 7% all primary care patients

So even if all HH in tiredness group 1 in 140 have HH

In UK men cause 1.6% hip fractures, 5.8% liver cancer.



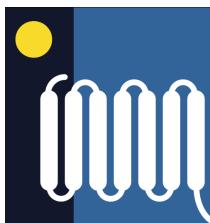
Progress check

Part 1

- Pedigree allows identification of inheritance type if any
- Needed to calculate risk

Part 2

- Risk helps informed decision making
- Can reassure some patients if not affected
- Identify patients who require further support
 - Genetic counselling



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