

IMPERIAL

Brittle bones - tutorial



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Insendi: Genetics – Brittle Bones



Imperial College
School of Medicine

English (En-GB) ▾

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Principles of Medicine 1a 2023-24

MBBS Phase 1a 2023-24

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5: Genetics

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5.1 Modes of inheritance

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5.2 Chromosomal abnormalities

Campus ▾

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⋮

5.3 Complex disease & pharmacogenetics

Campus ▾

Unpublished ▾

⋮

5.4 Brittle bones

Campus ▾

Unpublished ▾

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Session Plan



Part 1

Case information - tutor

- Recap Insendi material
- How does it link to what I know?
- How does it link to future PoM material?

Part 2

Questions (Group work)

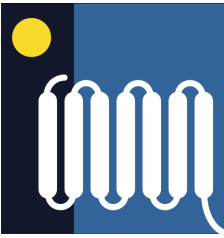
- Engage with content
- Discuss with each other/staff
- Formulate answers - active learning

Part 3

Consolidation

- Debrief with tutor
- Clarify understanding – ask questions?
- Take home messages

Susan M



- A female child, Susan M, had been born after a normal pregnancy.
- She had a history of frequent long-bone fractures resulting from minor injury and necessitating treatment in the Casualty Department of the local hospital.
- Child abuse was initially suspected.
- As Susan developed, skeletal abnormalities were detected which were confirmed on X-ray.
- At 3 years of age, she could still only crawl and had difficulty standing.
- Susan's type I collagen genes were investigated.

Brittle bones - 1



- Fibroblasts were cultured from a skin biopsy, and the collagen I they synthesised was analysed by gel electrophoresis in the presence of SDS.
- Type I collagen consists of two $\alpha_1(I)$ polypeptide chains and one $\alpha_2(I)$ chain. Some of the patient's $\alpha_1(I)$ polypeptide chains had unusually low electrophoretic mobility in the absence of 2-mercaptoethanol but co-migrated with normal $\alpha_1(I)$ chains in the presence of the reagent

1-POM-1-5

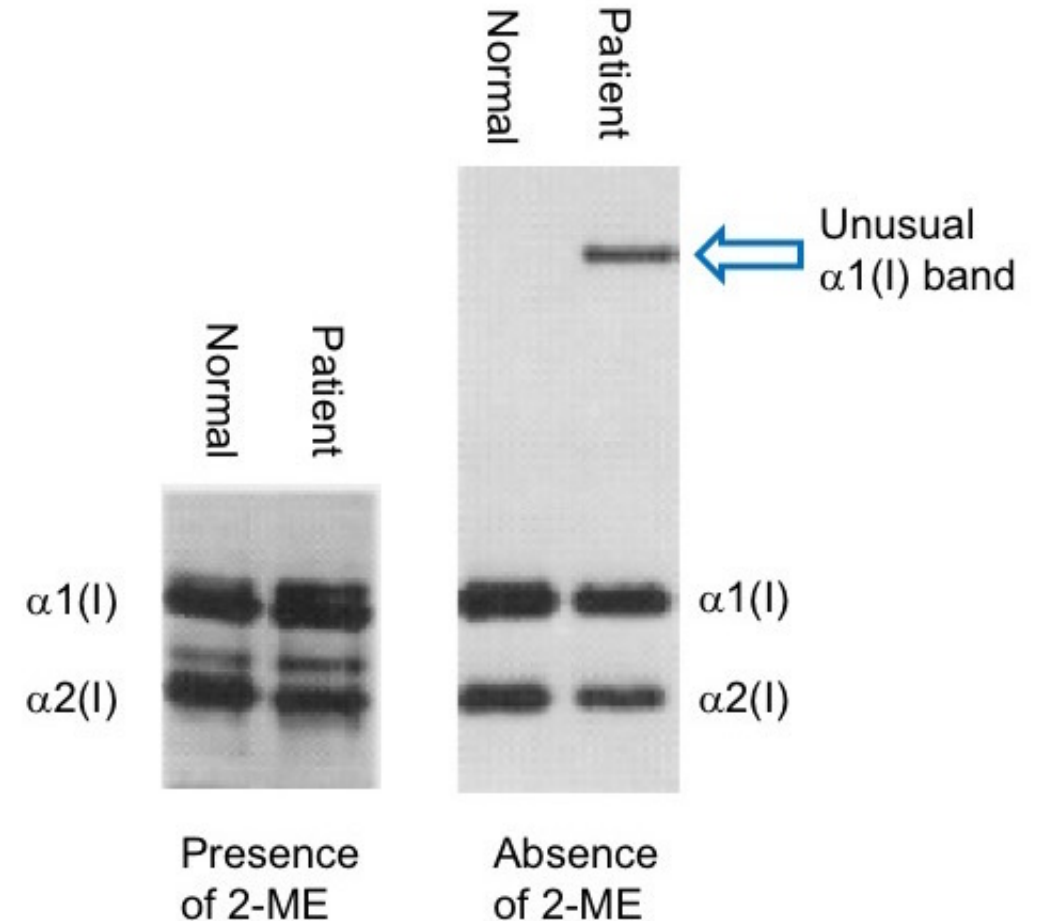


Figure: SDS-poly acrylamide gel electrophoresis of normal and patient samples

Brittle bones - 2



The child was shown to be heterozygous for a mutation in the COL1A1 gene which encodes the procollagen precursor of $\alpha_1(I)$ collagen. A comparison of part of the nucleotide sequences of the normal and mutant COL1A1 genes is shown below.

Normal COL1A1 sequence

5'...CCT CGG GGT GCC AAT GGT GCG CCT GGC AAC GAC GGT GCT AAG GGC GAT GCC GGT...3'

Patient's COL1A1 sequence

5'...CCT CGG GGT GCC AAT GGT GCG CCT TGC AAC GAC GGT GCT AAG GGC GAT GCC GGT...3'

Questions to be addressed (in groups)



- Use the genetic code to predict the effect of the mutation on the protein sequence.
- Explain why this change might cause the altered electrophoresis pattern.
- Why are only some of the α_1 (I) collagen chains affected?

Take home messages - 1



- Use the genetic code to predict the effect of the mutation on the protein sequence.

Normal COL1A1 sequence

5'...CCT CGG GGT GCC AAT GGT GCG CCT GGC AAC GAC GGT GCT AAG GGC GAT GCC GGT...3'
P R G A N G A P **G** N D G A K G D A G

Patient's COL1A1 sequence

5'...CCT CGG GGT GCC AAT GGT GCG CCT TGC AAC GAC GGT GCT AAG GGC GAT GCC GGT...3'
P R G A N G A P **C** N D G A K G D A G

Take home messages - 2

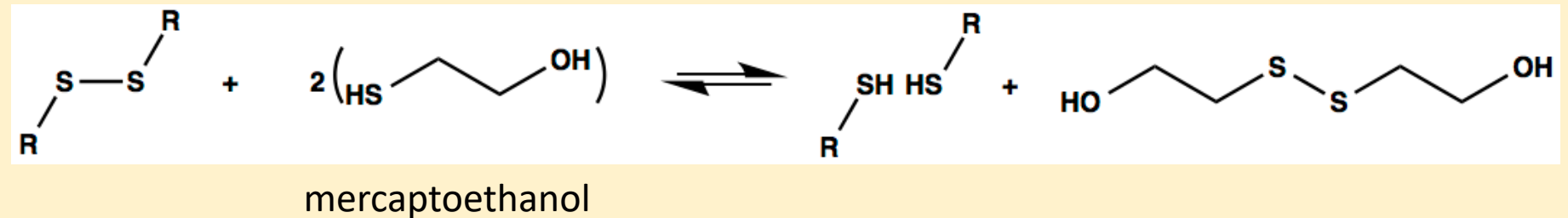
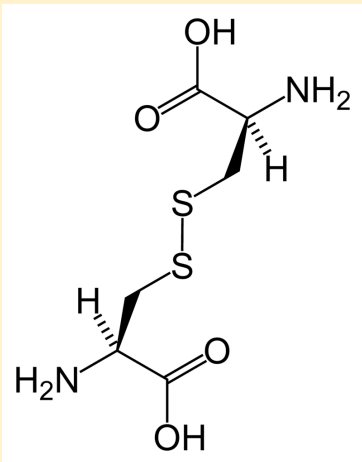


- Explain why this change might cause the altered electrophoresis pattern.

Sequence changes from glycine to cysteine

Cysteines can form disulphide bridges linking two chains together

2-Mercaptoethanol breaks these disulphide bridges

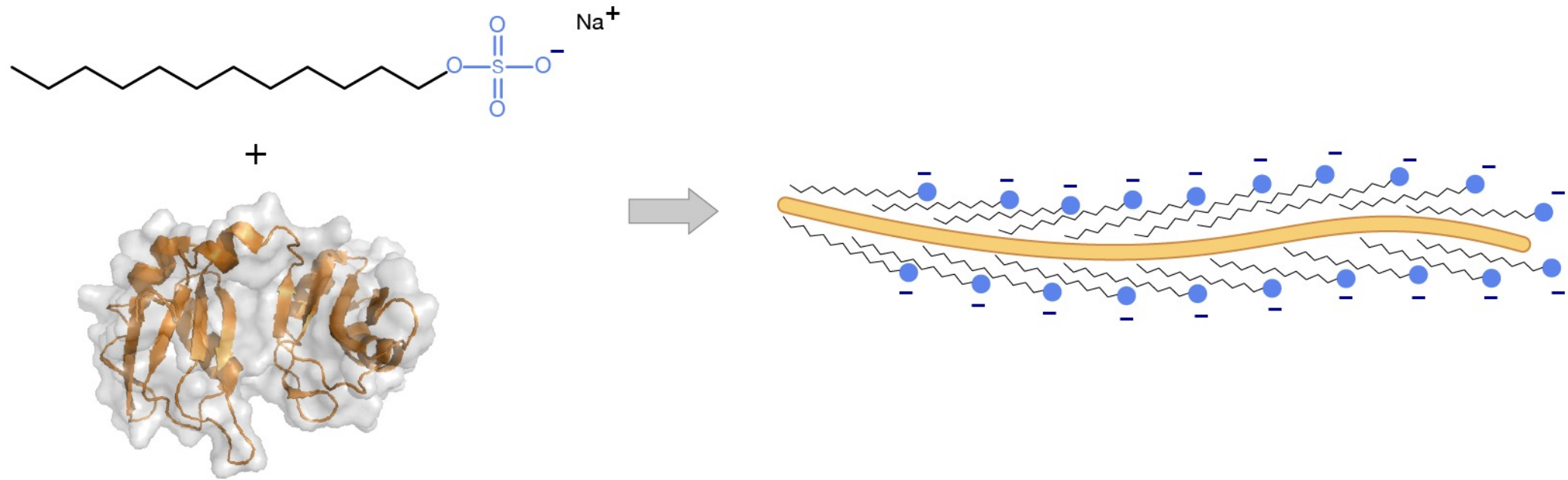


Take home messages - 2

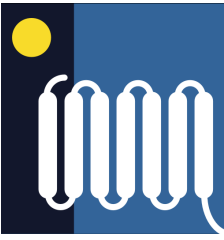


- Explain why this change might cause the altered electrophoresis pattern.

SDS-PAGE sodium dodecyl sulfate poly acrylamide gel electrophoresis (sodium lauryl sulfate)



Take home messages - 2



- Why are only some of the $\alpha_1(I)$ collagen chains affected?

Since only one of the two copies of ColA1 gene are mutated only some collagen molecules carry the mutation

To form complex need two copies of mutated protein to combine

Questions to be addressed (in groups)



- Predict possible biochemical consequences of the change on the assembly of type I collagen.
- Subsequent investigations showed the disorder had a dominant pattern of mutation in the patient's family. Explain this by reference to the structure of collagen.
- Why might the predicted change cause skeletal abnormalities and brittle bones?

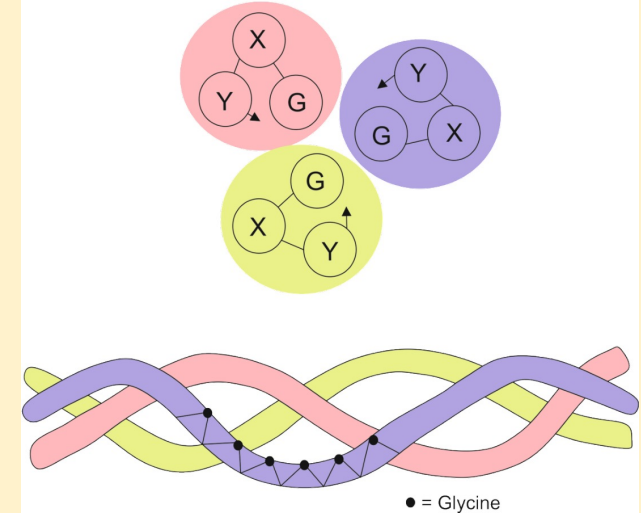
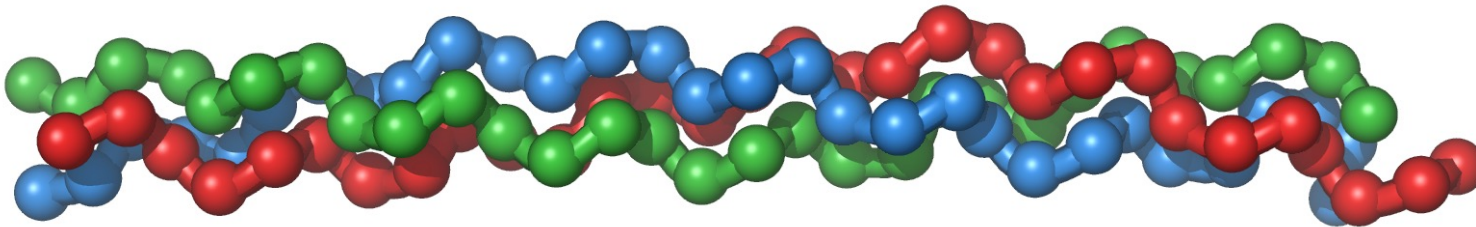
Take home messages - 4



- Predict possible biochemical consequences of the change on the assembly of type I collagen.

-Gly, X, Y-

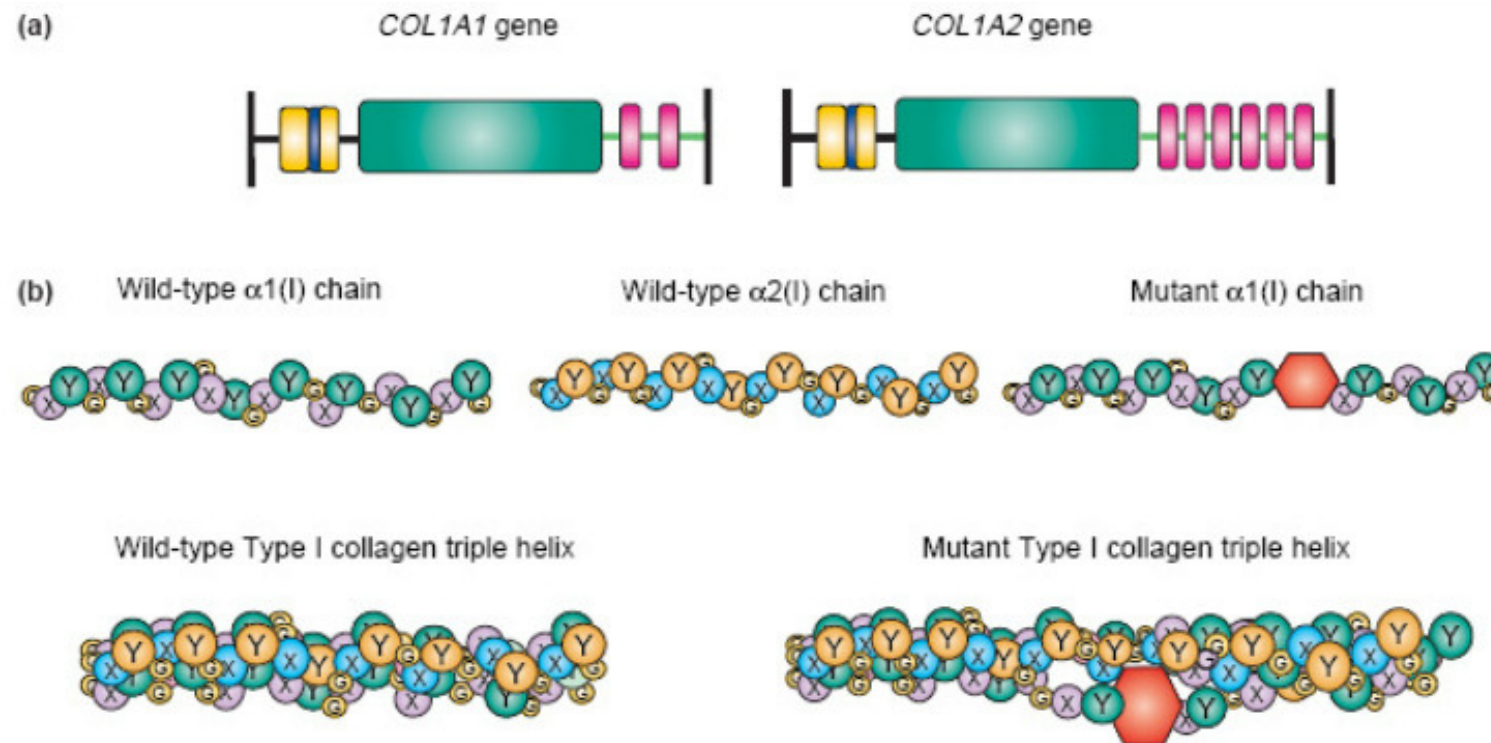
Glycine is in centre of triple helix. Other AA wont fit



Take home messages - 4



- Predict possible biochemical consequences of the change on the assembly of type I collagen.



TRENDS in Molecular Medicine

1-POM-4-1: Modes of inheritance: Differentiate between recessive and dominant autosomal and X-linked disorders, mitochondrial disorders, and explain their segregation patterns.

1-POM-1-8: Tissues: Summarise the main components of tissues, and outline the organisation of different types of epithelial tissues.

Take home messages - 5



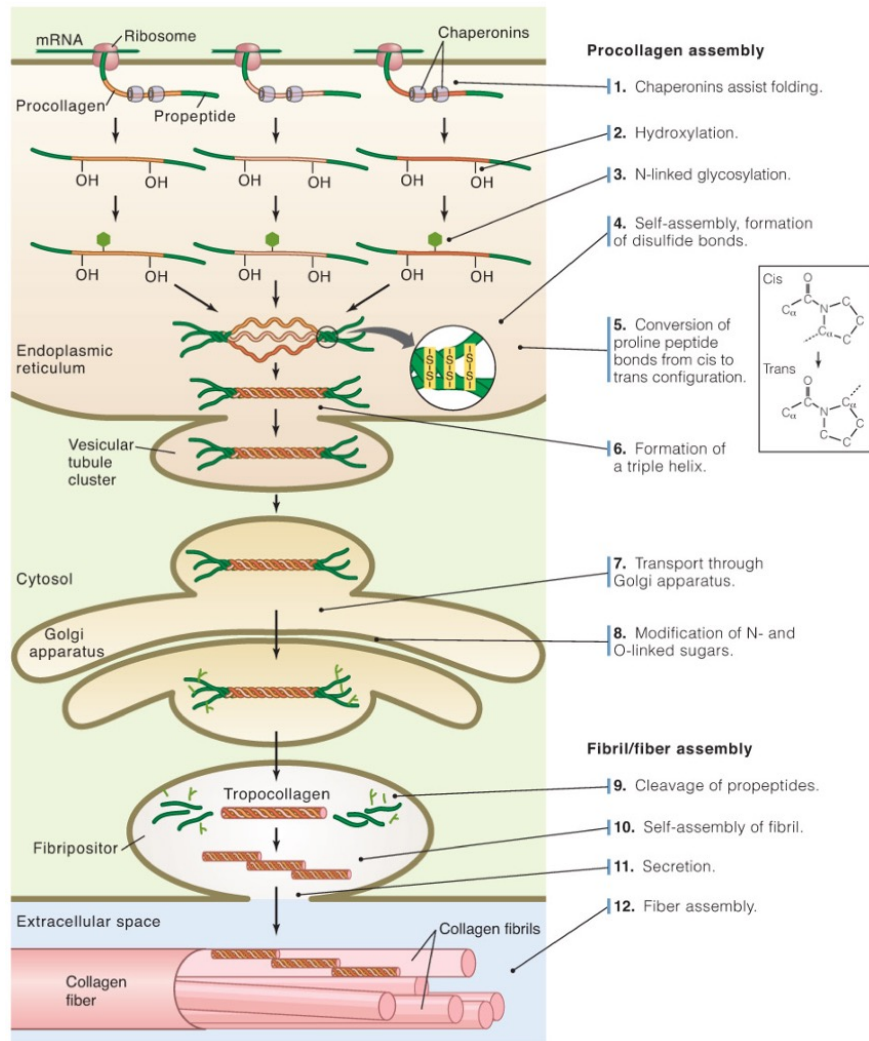
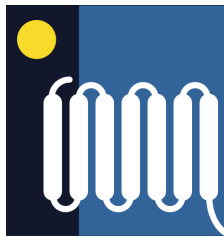
- Subsequent investigations showed the disorder had a dominant pattern of mutation in the patient's family. Explain this by reference to the structure of collagen.

Dominant negative effect or Gain of function mutation

The mutation disrupts the activity of the normal version of ColA1

1-POM-4-1

Take home messages - 5



only half of col α 1 protein mutated
all fibrils will be affected due to packing

Take home messages - 6



- Why might the predicted change cause skeletal abnormalities and brittle bones?

Initially skeleton laid down as collagen

Later stage mineralisation

If collagen defective then bone defective.

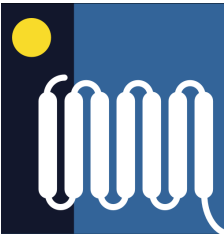
Skeleton is an active tissue whole skeleton turns over every 5-10 years

Questions to be addressed (in groups)



- Suggest a suitable prenatal diagnostic test to identify foetus who will suffer from OI

Take home messages - 7



- Suggest a suitable prenatal diagnostic test to identify foetus who will suffer from OI

1-POM-4-8

Need a sample derived from foetus eg amniocentesis or chorionic villus sampling
Then can use one of several approaches

Take home messages - 7



- Suggest a suitable prenatal diagnostic test to identify foetus who will suffer from OI

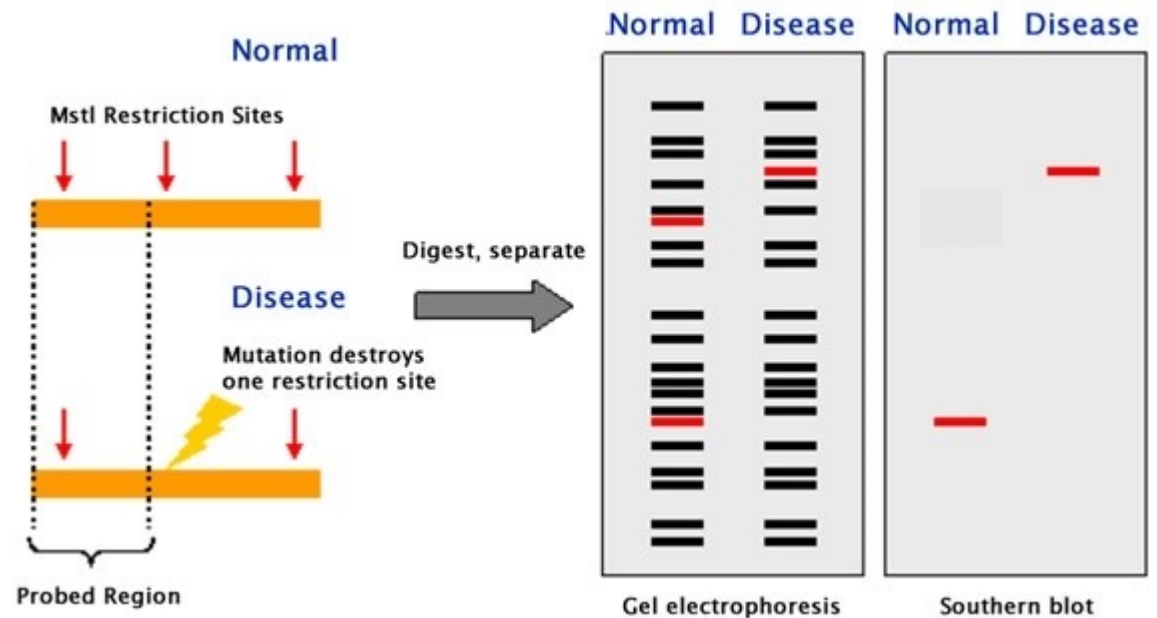
RFLP –restriction fragment length polymorphism

Mutation either removes or creates restriction site

Digest DNA

Gel electrophoresis

Use a probe to region of DNA near mutation



Take home messages - 7



- Suggest a suitable prenatal diagnostic test to identify foetus who will suffer from OI

PCR –amplify region with mutation
Gel electrophoresis
Use a probe specific for mutation
Sequence PCR product.

Polymerase chain reaction - PCR

