# <u>Thalassaemias</u>

### **Definition:**

Family of inherited haemoglobinopathies resulting from decreased synthesis of  $\alpha$  or  $\beta$  globin chains.

### Clinically they r divided into :

1-Thalassamia major: (Transfusion dependent)

A- Hydrops foetalis

B-  $\beta$  Thalas. Major

### 2-Thalassamia intermedia:

Ch. by moderate an. usually e' splenomegaly & iron overload.

3-Thalassamia minor: symptomless carrier.

# <u>β Thalassamia</u>

#### Def:

Severe H.A in infancy & early childhood e' certain characteristic features:

- MHA
- Anisocytosis, poikilocytosis, target cs.
- Splenomegaly
- Mongoloid facies
- serum iron
- Patient usually dies from cardiac arrhythmias

#### **Clinical classification:**

Thalassamia major

- " intermedia
- ,, minor

#### Genetic classification:

 $\beta$  = normal gene  $\beta$ + = partial synthesis of  $\beta$  chain  $\beta$  o = complete absence of  $\beta$  chain

### Genetic expression may be:

<b>1-βοβο :</b> thalas. Major			
Hb: all r F & A2 (no Hb A)			
<b>2-</b> $\beta$ + $\beta$ + : thalas. major or intermedia			
variable amount of Hb A, F & A2			

- **3-β+βo :** thalas. major or intermedia variable amount of Hb A, F & A2
- **4-ββo**: thatas. minor or intermedia

Hb A, F & Hb A2

**5-**ββ+: thalas. minor or intermedia

just **†**Hb A2

### **Classification & Terminology of Beta Thalassemia**

•	Normal	β/β
•	Minor	$eta/eta^0$ $eta/eta^+$
•	Intermedia	$eta^0/eta^+$ $eta^+/eta^+$
•	Major	$eta^0/eta^0$ $eta^+/eta^+$
		$eta^0\!/eta^+$

# Molecular basis of β thalassamia

The defect mainly is Quantitative : due to amount of m-RNA. Most of  $\beta$  thal. syndromes r caused by mutations affecting gene regulation or expression rather than gene deletion (unlike  $\alpha$  thalas.)

#### N.B: causes:

- **1- Deletion:**
- 2- Non deletion:
- A- mutation of m-RNA transcription
- B- ", " processing
- C- ", ", translation



### I- Deletion: βo Rare

Deletion of a part or whole of 3' or 5' end of  $\beta$  gene (mainly in  $\alpha$  thalassamia not  $\beta$  ).

### **II-** Non deletion = Mutation:

### A- Mutations of transcription:

### **1-promotor region mutation:** β+

 $\downarrow \text{ m RNA transcription} \rightarrow \forall \text{ amount of globin synthesis} \rightarrow \\ \downarrow \text{ synthesis of } \beta \text{ chain } = \beta +$ 

### 2- chain terminator mutation: βo

This leads to : mRNA is incapable of being translated into full length globin chains resulting in  $\beta$ o phenotype.

### **B-** Mutations of processing:

Mutations affecting splicing, capping or polyadenylation  $\rightarrow$  unstable m-RNA $\rightarrow$   $\downarrow$ amount of globin synthesis  $\rightarrow$   $\downarrow$ synthesis of  $\beta$  chain

### **1-** Splice junction mutations : βο

- Point mutation involving splicing sites result in abnormal splicing.
- The m-RNA produced is useless as a messenger for  $\beta$  globin synthesis  $\longrightarrow \beta o$

### 2- Mutations of consensus sequences : $\beta$ +

- Mutations involving consensus seq. (boundaries surrounding splice junction) results in formation of cryptic donor site.
- These cryptic sites resemble N splice sites but r not recognized unless N sites r altered.
- These mutations leads to  $\oint$  splicing not abolish it  $\longrightarrow \beta +$

### 3-Mutations creating new splicing sites: $\beta$ + or $\beta$ o

- Nucleotide substitution e'in introns results in formation of new splice sites, despite of presence of functioning N splice sites.
- The new splice sites compete e' N splice site  $\rightarrow \beta + \text{ or } \beta 0$

# 4- Activation of cryptic donor site : β+

- Exons contain cryptic sites e' a nucleotide sequence resembling N seq.
- Mutations of these cryptic sites lead to their activation, competition between abnormal new splice seq. & N splice seq  $\rightarrow$  mild  $\beta$ +

# 5-Mutations of polyadenylation : $\beta$ +

Mutation of AATAAA seq. at 3' end  $\rightarrow$  transcription continue elongated m-RNA (unstable) $\rightarrow \beta$ +

# 6- Mutations at cap site : $\beta$ +

Substitution of C for A in 1<sup>st</sup> position may  $\downarrow$  transcription or slow the 5' capping  $\rightarrow \downarrow$  m-RNA stability $\rightarrow \beta$ +

# **<u>C- Mutations causing abnormal translation of m-RNA:</u>**

# **1-** Nonsense mutation : βο

Point mutation (single aa substitution) creation of a stop codon  $\rightarrow$  prevent translation of m-RNA  $\rightarrow$  premature stop codon  $\rightarrow \beta o$ 

# 2- Frame shift mutation:

1,2,4 base insertion or deletion  $\rightarrow$  disturbance of N reading frame  $\rightarrow$  creation of termination codon



# **Pathophysiology**



# Thalassamia major (Cooley's anaemia)

Severe anaemia manifested early in life e' splenomegaly & bony deformaties

### Genetic expression:

βo/βo β+/β+ βo/β+ δβ lepore / δβ lepore  $βo/E \rightarrow$  (Hb E → thalassamic Hb as it is

abnormal Hb &  $\downarrow$  amount

C/P:

of chronic H.A

# Lab findings:

1- Evidences : 3

# **2-CBC:**

- MHA
- Target cells, macrocytes ( **†**etics)
- Anisocytosis, poikilocytosis
- Normos
- basophilic stippling, capot rings
- retics put doesn't correlate e' degree of anemia (ineff.eryth)
- WBCs & platelets: N

# **3-BM:**

# Hypercellular, erythroid hyperplasia

Iron stain:

↑ iron stores, ↑ sideroblasts.

# **4- Evidence of ineffective erythropoiesis**

**5-Special tests:** 

A- Hb electrophoresis:

**†** Hb F & A2

Hb A (variable acc. to genetic variant & molecular basis)

B- Osmotic fragility: ↓ due to ↑ retics w' resist lysis C- Ferrokinetics:

Ineffective erythropoiesis

# **D-** Gene study by PCR:

For prenatal diagnosis.

# Thalassemia major





# HB ELECTROPHORESIS

# Thalassemia Intermedia

Milder than thal. major but more severe than asymptomatic thal. Trait

**Genetic expression:** 

 $\begin{array}{c} \beta + /\beta + \\ \beta /\beta o \\ \beta o / (\delta \beta) o \\ \beta + / (\delta \beta) o \\ (\delta \beta) o / (\delta \beta) o \\ double heterozygous lepore: \\ \beta + / \delta \beta lepore \\ \beta o / \delta \beta lepore \\ Coinheritance of a thalas. ( \downarrow a chain) \\ Hb H (ao/a+) \end{array}$ 

### **C/P:**

### Varies from:

### 1- severe:

- Patient presents e' anaemia later than thal. Major
- Hb 6 g/dl e'out transfusion
- Growth retardation
- Skeletal deformaties
- Splenomegaly
- Leg ulcers

# 2- completely asymptomatic until adult life & transfusion independent

- e' Hb level 10-12 g/dl
- 3- varieties of intermediate severity

### Lab diagnosis:

- same as thal.minor
- Hb A : 20-40%

# <u>Thalassaemia minor</u>

# **Pathogenesis:**

 $\downarrow \beta$  or  $\downarrow \delta\beta$  synthesis

# Genetic expression:

- $\beta/\beta+$
- $\beta/\beta o$

 $\beta/(\delta\beta)o$ 

 $\beta/\delta\beta$  lepore

Hereditary persistence of fetal Hb (HPFH)

 $\alpha$ o thalas. trait

 $\alpha_{\scriptscriptstyle +}$  thalas. Trait

# C/P:

Asymptomatic, discovered accidently

### Lab :

- Anaemia is mild or absent, but  $\downarrow$  MCV,  $\downarrow$  MCH
- RBCs:
- Hypochromia, target cs, basophilic stippling
- WBCs & plat: N
- chemistry:
- firon or ferritin
- Hb E/P:
- † Hb A2 & F

# D.D:

Iron def. an. (MHA)

	Thalassamia minor	Iron def. anaemia
Serum iron	<b>↑</b>	
Serum ferritin		↓ ↓
Iron stores		Absent
sideroblasts		
Hb A2		+

# **Treatment for Beta Thalassemia**

- **Trait** no treatment required
- Intermedia
- Major (Cooley anemia)
  - Regular folate supplementation
  - **RBC transfusion** (Splenectomy may decrease need

for transfusions)

- to maintain [Hgb] ~9-10g/dL
- Blood transfusions → iron accumulation → iron overload
- Iron chelators (disferroxamin)
- Bone marrow transplantation (BMT)

-BMT has been attempted from donors with matching alleles.

• Gene therapy—the future