

Hemoglobinopathies

Types of Hereditary Hemoglobin Disorders

Type	Description
1. Quantitative abnormalities	• Abnormalities in the relative amounts of α and β subunits. → Thalassemias .
2. Qualitative abnormalities	• Structural variants caused by mutations altering amino acid sequence of globin chain. → >800 variants identified.
3. Hereditary persistence of fetal hemoglobin (HPFH)	• Impairment of the switch from γ -globin → β -globin after birth. • Fetal Hb ($\alpha_2\gamma_2$) has higher O_2 affinity than adult Hb ($\alpha_2\beta_2$). • Persistence of HbF → mild or no symptoms.

Thalassemias Overview

| Definition |

Most common single-gene disorder (either α or β). Caused by reduced synthesis of α or β chains → imbalanced $\alpha:\beta$ ratio.

| Main consequences |

• Imbalance in chain production → precipitation of excess unpaired chains → RBC destruction → anemia.

| Types |

α -Thalassemia: reduced α -chain synthesis.

β -Thalassemia: reduced β -chain synthesis.

Alpha-Thalassemia

| Definition |

Reduced or absent synthesis of α -globin chains. Mainly due to deletion mutations that inactivate α -globin genes.

| Genetic basis |

- 4 α -globin genes total (2 per chromosome 16).
- Severity depends on number of genes deleted (1–4).

| Mechanism |

↓ α -chain synthesis → excess β or γ chains → formation of unstable tetramers (β_4 or γ_4).

| Pathophysiology |

Unstable Hb (HbH or Hb Bart's) → high O_2 affinity → poor O_2 release → tissue hypoxia → hemolysis.

Subtypes of α -Thalassemia

Subtype	Genes Deleted	Main Hb Formed	Clinical Features
α^0 -Thalassemia (Hydrops fetalis / Hb Bart's)	4/4	Hb Bart's (γ_4)	• Very high O_2 affinity (cannot release O_2). • Severe anemia, edema (hydrops fetalis). • Stillbirth or death soon after birth.
α^{+++} -Thalassemia (HbH disease)	3/4	HbH (β_4)	• β-chain tetramers with high O_2 affinity, unstable $\rightarrow$ Heinz bodies. • Mild–moderate (sometimes severe) hemolytic anemia. • May need transfusions; not fatal.
α^{++} -Thalassemia trait	2/4	—	• Mild microcytic anemia. • Clinically mild.
α^+ -Thalassemia (Silent carrier)	1/4	—	• No symptoms, clinically silent.

Summary of α -Thalassemia Severity

Deleted genes	Type	Main Hb formed	Severity
4	α^0 (Hydrops fetalis)	Hb Bart's (γ_4)	Lethal
3	α^{+++} (HbH disease)	HbH (β_4)	Mild–moderate anemia
2	α^{++} (Trait)	—	Mild microcytic anemia
1	α^+ (Silent carrier)	—	Asymptomatic

Beta-Thalassemia

| Definition |

Reduced (β^+) or absent (β^0) synthesis of β -globin chains. Caused mainly by point mutations in β -globin gene on chromosome 11.

| Mechanism |

↓ β -chain → excess α -chain → precipitation → RBC membrane damage → hemolysis.

| Main mutation type |

Point mutations (not deletions).

| Result |

Imbalanced α : β chain ratio → ineffective erythropoiesis and anemia.

Types of β -Thalassemia

Type	Genetic basis	Clinical severity
β^0 -Thalassemia	No β -chain synthesis	Severe
β^+ -Thalassemia	Reduced β -chain synthesis	Mild–moderate
β -Thalassemia major (Cooley's anemia)	Both β genes defective	Severe anemia; requires lifelong transfusion
β -Thalassemia intermedia	Partial β production	Moderate anemia; occasional transfusion
β -Thalassemia minor (trait)	One gene defective	Mild microcytic anemia; often asymptomatic

Pathophysiology of β -Thalassemia

Defect	Effect
↓ β synthesis	→ Excess α -chains precipitate in RBCs.
Precipitated α -chains	→ Damage RBC membrane → intramedullary destruction.
Ineffective erythropoiesis	→ Bone marrow expansion (especially in skull/face).
Compensatory mechanisms	→ Extramedullary hematopoiesis → splenomegaly.
Iron overload	→ From transfusions + ↑ absorption → hemosiderosis.
Clinical findings	Anemia, bone deformities, hepatosplenomegaly, growth retardation.



Comparison: α -Thalassemia vs β -Thalassemia

Feature	α -Thalassemia	β -Thalassemia
Defective chain	α -chain	β -chain
Mutation type	Deletion	Point mutation
Excess unpaired chain	β or $\gamma \rightarrow$ form β_4, γ_4	α -chains precipitate
Major Hb abnormality	Hb Bart's or HbH	$\downarrow$ HbA, $\uparrow$ HbA ₂ , $\uparrow$ HbF
Severity depends on	Number of genes deleted	β^0 vs β^+ mutation
Main mutation location	Chromosome 16 (α gene cluster)	Chromosome 11 (β gene)
Pathophysiology	Unstable tetramers	Ineffective erythropoiesis
Mutation effect	Quantitative defect	Quantitative defect

🧩 Qualitative Abnormalities (Structural Variants)

| Definition |

Structural variants of Hb caused by point mutations that alter amino acid sequence
 $\rightarrow$ change in solubility, stability, or O₂ affinity.

| Categories |

- A. Mutations in surface residues
- B. Mutations in internal residues
- C. Mutations at $\alpha 1$ - $\beta 2$ contacts
- D. Mutations stabilizing methemoglobin

| Total variants |

> 800 hemoglobin variants known.

A.

Mutations in Surface Residues – Hemoglobin C (HbC)

| Mutation |

$\beta 6$ Glutamic acid $\rightarrow$ Lysine

| Effect |

Replacement of negatively charged residue by positive $\rightarrow$ promotes Hb crystallization.

| Clinical features |

Mild chronic hemolytic anemia (less severe than sickle cell).

| Heterozygotes (HbA/HbC) |

Usually asymptomatic.

B.

Mutations in Internal Residues – Hemoglobin E (HbE)

| Mutation |

β 26 Glutamic acid $\rightarrow$ Lysine

| Effect |

Occurs near splice site $\rightarrow$ abnormal splicing $\rightarrow$ $\downarrow$ β -chain synthesis.

| Nature of defect |

Both qualitative and quantitative (like mild thalassemia).

| Clinical |

Mild anemia, common in Southeast Asia.

C.

Mutations at α 1- β 2 Contacts – Hemoglobin S (HbS)

| Mutation |

β 6 Glutamic acid $\rightarrow$ Valine

| Effect |

Valine (nonpolar) interacts hydrophobically with adjacent Hb $\rightarrow$ polymerization in deoxy state.

| Result |

RBCs deform (sickle shape) $\rightarrow$ vaso-occlusion, hemolysis, ischemia.

D.

Mutations Stabilizing Methemoglobin – Hemoglobin M

| Mutation site |

Near heme pocket (e.g., His $\rightarrow$ Tyr).

| Effect |

Stabilizes Fe^{3+} (ferric state) $\rightarrow$ cannot bind O_2 .

| Clinical |

Methemoglobinemia $\rightarrow$ cyanosis, usually mild symptoms.

🧬 Sickle Cell Disease (HbS)

| Genetic mutation |

Point mutation in β -globin gene: Glu6Val (Valine replaces Glutamic acid).

| Mechanism |

- Valine forms hydrophobic patch $\rightarrow$ polymerization of deoxy-HbS $\rightarrow$ sickled RBCs.

| Factors promoting sickling |

$\downarrow$ O₂, dehydration, high Hb concentration, low pH.

| Reversibility |

Sickling is initially reversible with oxygenation, but repeated cycles cause membrane damage $\rightarrow$ irreversible sickle cells.

| Clinical features |

- Hemolytic anemia
- Painful vaso-occlusive crises
- Splenic infarction $\rightarrow$ autosplenectomy
- Increased risk of infection by encapsulated bacteria
- Bone and organ infarctions

| Sickle cell trait (HbA/HbS) |

Heterozygous; mostly asymptomatic; protection against Plasmodium falciparum malaria.

| Sickle cell disease (HbS/HbS) |

Homozygous; severe anemia, chronic crises, organ damage.

| Treatment |

- Hydroxyurea ($\uparrow$ HbF production)
- Transfusions, folic acid, infection control
- Bone marrow transplantation (curative in some).

🧐 Hereditary Persistence of Fetal Hemoglobin (HPFH)

| Definition |

Genetic condition with continued synthesis of γ -globin after birth.

| Mechanism |

Mutation in β -globin gene cluster $\rightarrow$ failed switch from $\gamma \rightarrow \beta$ production.

| Effect |

Elevated HbF ($\alpha_2\gamma_2$) levels in adults.

| Clinical significance |

- Benign; usually asymptomatic.
- In β -thalassemia or sickle cell $\rightarrow$ HbF presence lessens symptoms (because HbF inhibits polymerization of HbS).