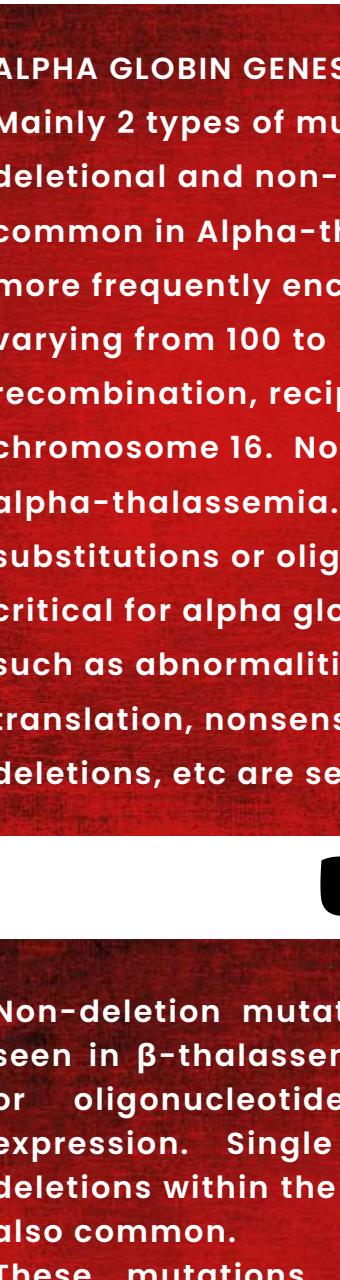
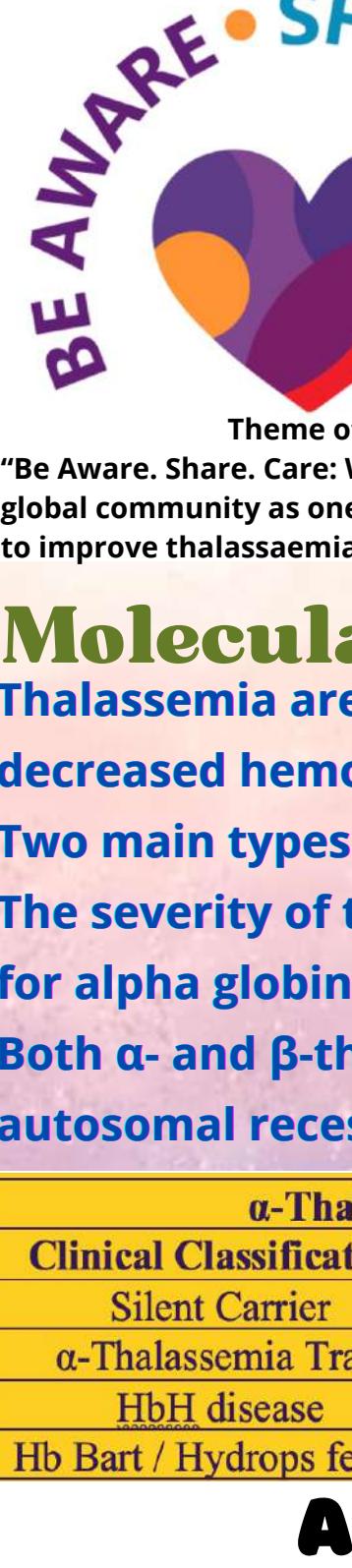


Newsletter



Theme of 2022
"Be Aware. Share. Care: Working with the global community as one to improve thalassaemia knowledge."

All India Institute of Medical Sciences, Rajkot

CLINICAL BIOCHEMISTRY & MOLECULAR BIOLOGY BULLETIN

Volume 1

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An estimated 30 million people have the Beta Thalassemia blood disorder in India*

Molecular Genetics of Thalassemia Thalassemia are inherited blood disorders characterized by decreased hemoglobin production.

Two main types are: Alpha thalassemia and Beta thalassemia.
The severity of the disease depends on how many of the four genes for alpha globin or two genes for beta globin are malfunctioning.
Both α- and β-thalassemia are commonly inherited in an autosomal recessive manner.

α-Thalassemia	β-Thalassemia		
Clinical Classification	Genotype	Clinical Classification	Genotype
Silent Carrier	αα/α	β-Thalassemia Minor	β ⁺ /β or β ⁰ /β
α-Thalassemia Trait	αα/α or αα/αα	β-Thalassemia Intermedia	β ⁺ /β ⁺ or β ⁰ /β ⁺
HbH disease	αα/αα	β-Thalassemia Major	β ⁰ /β ⁰
Hb Bart / Hydrops fetalis	αα/αα		

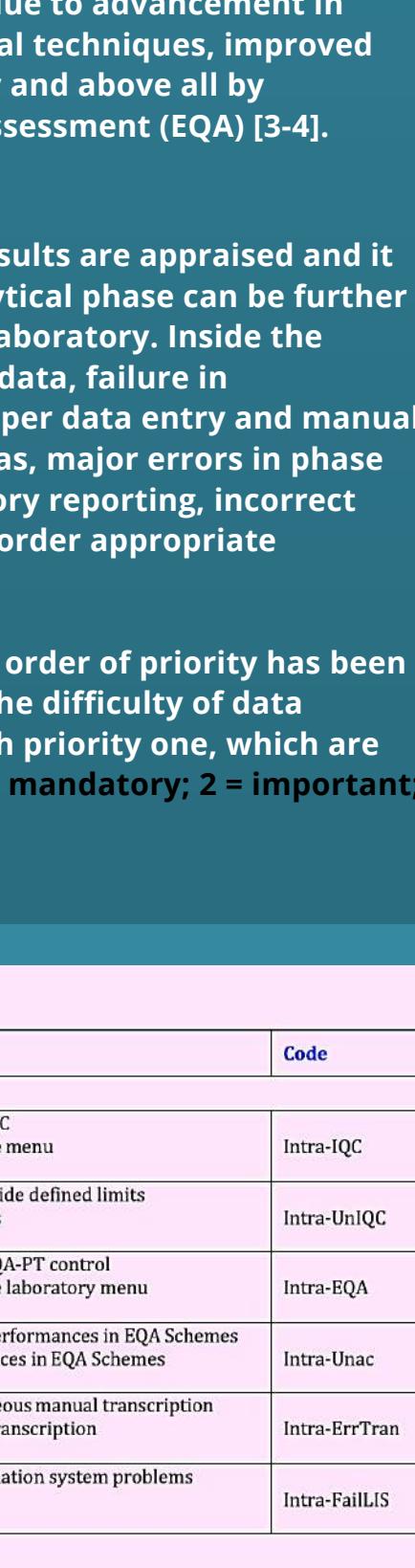
ALPHA- THALASSEMIA

ALPHA GLOBIN GENES ARE LOCATED AT CHROMOSOME 16 (16P13.3). Mainly 2 types of mutation detected in alpha thalassemia: deletional and non-deletional. Deletional mutations are more common in Alpha-thalassemia. -alpha3.7 and -alpha2.4 are more frequently encountered deletions. Extended deletions, varying from 100 to >250 kb may result in illegitimate recombination, reciprocal translocation, and truncation of chromosome 16. Nondeletion defects are less frequently found in alpha-thalassemia. These defects include single nucleotide substitutions or oligonucleotide deletions/insertions in regions critical for alpha globin gene expression. Several mechanisms such as abnormalities of RNA splicing and of initiation of mRNA translation, nonsense and frameshift mutations, in-frame deletions, etc are seen in alpha-thalassemia.



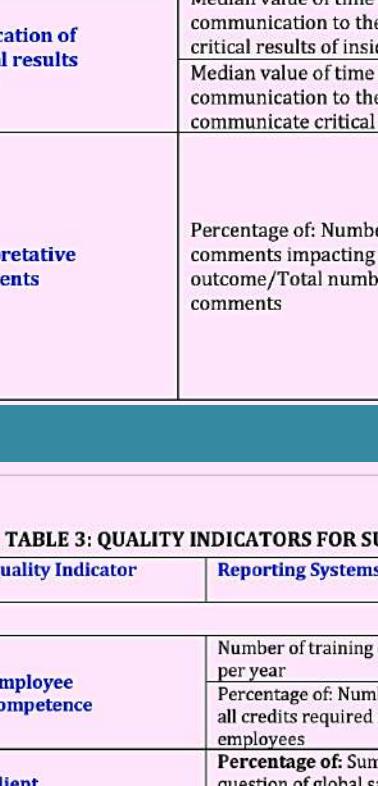
BETA- THALASSEMIA

Non-deletion mutations, especially, point mutations commonly or oligonucleotide insertions/deletions affecting β gene expression. Single base substitutions, small insertions, or deletions within the gene or its immediate flanking sequences are also common. These mutations affect β-globin expression in 3 different mechanisms: mutations leading to defective β-gene transcription (promoter and 5' untranslated region [UTR] mutations), mutations affecting messenger RNA (mRNA) processing (splice-junction and consensus sequence mutations, polyadenylation, and other 3' UTR mutations), and mutations resulting in abnormal mRNA translation (nonsense, frameshift, and initiation codon mutations).



Deletional mutations are less common in β-thalassemia. Deletion of portion of structural gene, enhancer region or promoter region can affect functionality of the beta globin gene significantly.

Experts Talk



Dr. Pinky Meena
Asst. Prof Pediatrics

Thalassemia is one of the most common preventable single gene disorders. In severe form it is associated with chronic, life-imposing and life-threatening diseases with inherent serious health sequelae that can lead to disability or death. Unfortunately, a terrible series of children in our country continue to be born and suffer from such disorders mainly due to lack of strategies and lack of comprehensive programme and systematic approaches to prevent it. Data on the incidence of thalassemia ranges from 2.0-4.8%. It is estimated that about 1000-1500 babies with the Thalassemia Major (α0/α0) are born every year. The only cure available for these children with thalassemia is hematopoietic stem cell transplant (HSCT). However, this percolates to only a few hematologists because of cost, limited BMT centres, or unavailability of a suitable HLA matched donor. Therefore, the mainstay of treatment is a regimen of regular blood transfusions followed by adequately monitored iron chelation therapy. Considering the magnitude of the problem and the cost implications of this disease, prevention includes identifying the carriers and avoidance of marriage of carrier couples and secondary by preventing the birth of affected child through prenatal diagnosis. In countries like Cyprus, Italy, and Canada where successful screening programme in high schools and young adults for Thalassemia has led to decrease in the incidence of thalassemia.

MOLECULAR BIOLOGY & CLINICAL CHEMISTRY (MCC) UPDATES

THALASSEMIA CAN BE DIAGNOSED BY SEPARATION AND QUANTIFICATION OF HEMOGLOBIN FRACTIONS BY ELECTROPHORESIS OR HPLC.

MOLECULAR DIAGNOSIS OF THALASSEMIA CAN BE DONE WITH PCR AND NEXT GENERATION SEQUENCING TECHNIQUES.

THESE TECHNIQUES CAN SPECIFICALLY IDENTIFY CAUSATIVE MUTATIONS SUCH AS SINGLE NUCLEOTIDE SUBSTITUTIONS, INSERTIONS, SHORT OR LONG DELETIONS. THESE MOLECULAR BIOLOGY TECHNIQUES CAN BE USED FOR PRENATAL DIAGNOSIS OF FOETUS OF THALASSAEMIC COUPLE.

Recently, the Clinical Biochemistry service laboratory has got HPLC analyser installed in AIIMS, Khanderi. Our lab is now equipped to perform thalassemia diagnosis



Inappropriate turnaround times

Turnaround time (minutes) from sample reception in laboratory to release of result, of International Normalized Ratio (INR) at 90th percentile (STAT).

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Turnaround time (minutes) from sample reception in laboratory to release of result, of White Blood Cell (WBC) count at 90th percentile (STAT).

Turnaround time (minutes) from sample reception in laboratory to release of result, of Cardiac Troponin I (TnI) or Troponin T (TnT) at 90th percentile (STAT).

Turnaround time (minutes) from sample reception in laboratory to release of result, of Total Potassium (K) at 90th percentile (STAT).

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