

Molecular Genetics of β -thalassemia in the Eastern Province of Saudi Arabia

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Summary

Hemoglobinopathies are the most commonly inherited genetic disorders encountered among the population of the Eastern Province of Saudi Arabia. The thalassemia disease has two major forms known as α and β -thalassemia, and are characterized by quantitative deficiency of either α -globin or β -globin respectively. The high frequency and heterogenicity of β -thalassemia constitutes a major health problem. The objective of this project was to establish the molecular genetics of β -thalassemias in the Eastern Province. Genome DNA was isolated from the blood of 25 patients suspected of being carriers of β -thalassemia. Multiplex PCR for amplification of 4 regions in β -globin gene was undertaken.

The data show that IVS-2 nt 1 and IVS-1 nt 110 mutations are the most common in the population. Codon 39 mutation was also found with relatively high incidence. One of the subjects had a compound mutation which included IVS-2 nt 1 and IVS-1 nt 110 mutations. Six of the subjects with high Hbg A₂ showed none of the five common mutations. The results demonstrate the feasibility of using this cost effective technique to screen for the mutation as a pre-marriage requirement.