



β-Thalassemia: Prevention in Jordan



Daggag Hinda, Qubbaj Waffa', Abu Rubeiha Zaid, Shboul Mohammed, Natasha Saleh and El-Khateeb Mohammed
Molecular Genetics Laboratory, National Center for Diabetes, Endocrinology and Genetics,
Amman, Jordan.hinda.daggag@ncd.org.jo

Abstract

β-thalassemia is an autosomal recessive genetic disorder, characterized by defective beta-globin chain synthesis. At a carrier rate of 24% in a Jordanian population of about 5 million, with no effective treatment available, β-thalassemia represents a major financial and social burden on the patients and their families as well as on the national health system. This is only worsened by the high rate of consanguineous marriages in Jordan, estimated to be between 2030% (this includes only first-cousin marriages) for marriages contracted after 1980. To try and address this issue and reduce the number of infants born with β-thalassemia, a prenatal diagnosis program was initiated in Jordan in 1996, in collaboration between the National Center for Diabetes, Endocrinology and Genetics and the Gynecology Department at the King Hussein Medical Centre. In total, 324 chorionic villus samples (CVS) were screened for one of the seven most common β-thalassemia mutations in the Mediterranean region, between 1996-2009. Although most of the couples requesting the screen already had at least one affected child, in majority of cases, where CVS screen returned positive (i.e. the fetus was affected), the couples opted for selective termination of pregnancy. The initiation of prenatal screening (first of its kind in Jordan) together with the introduction of obligatory premarital screening for β-thalassemia in 2007, are both seen as positive steps towards the reduction and eventual prevention of β-thalassemia in Jordan.

Introduction

Thalassemia occurs when one or more of the disease-related genes fails to produce functional protein, leading to a shortage in one of the four hemoglobin subunits (hemoglobin normally consists of two alpha and two beta subunits) (Figure 1). β-thalassemia occurs when an β-globin gene fails (HBA1 and/or HBA2). β-thalassemia, characterised by reduced synthesis of the hemoglobin subunit beta, results from different mutations in the hemoglobin, beta gene HBB (Panigrahi and Agarwal, 2008; Weatherall and Clegg, 1981). So far, more than 200 mutations of the HBB gene have been characterised (Figure 2), most of them being single point mutations

β-Thalassemia In Jordan

β-thalassemia is one of the major inherited disorders in Jordan, with the carrier rate for the disease estimated at 24% (Hammamy et al., 2007). High incidence of β-thalassemia and other genetically determined recessive disorders in Jordan is largely due to high consanguinity marriages, with studies indicating the rate to be between 2030% for first cousin marriages (Hammamy et al., 2007). As no effective treatment exists for β-thalassemia and affected individuals require regular blood transfusion, this can represent a major financial and psychological load on the patients and their families, in addition to the challenges introduced onto the national healthcare system (Sadiq et al., 2001).

Premarital Screening

In order to try and address the above mentioned issues and reduce the rate of affected individuals, Jordanian Ministry of Health has introduced a national obligatory premarital screening for β-thalassemia, in 2007. Once the couple undergo the screening, they are offered genetic counseling, explaining the results of the screen and the available options (especially in cases where both individuals might be carriers for β-thalassemia). Nevertheless, in order to observe the true benefits of premarital screening on the rate of β-thalassemia in general population, thorough statistical analysis needs to be conducted on the national level, including figures pre and post 2007.

Prenatal Diagnosis

As part of prevention of β-thalassemia in Jordan, a prenatal diagnosis program was initiated in collaboration with the Gynecology Department at the King Hussein Medical Centre. In total, 324 chorionic villus samples (CVS) were screened for one of the seven most common β-thalassemia mutations in the Mediterranean region, between 1996-2009. Those mutations included IVSII-1 (G-A), IVSI-110 (G-A), IVSII-745 (C-G), IVSI-6 (T-C), IVSI-1 (G-A), codon 39 (C-T) and codon 37 (G-A). Amplification Refractory Mutation System by Polymerase Chain Reaction (ARMS-PCR) was used in order to test the CVS samples for different mutations. Restriction fragment length polymorphism (RFLP) haplotype analysis was also performed to enable exclusion of maternal contamination in the CVS.

Out of 300 couples tested, 71% were consanguineous (1st and 2nd degree relatives). In 25% of CVS cases, the foetus was unaffected whereas 53.5% of foetuses were found to be heterozygous and 13.4% were homozygous affected.

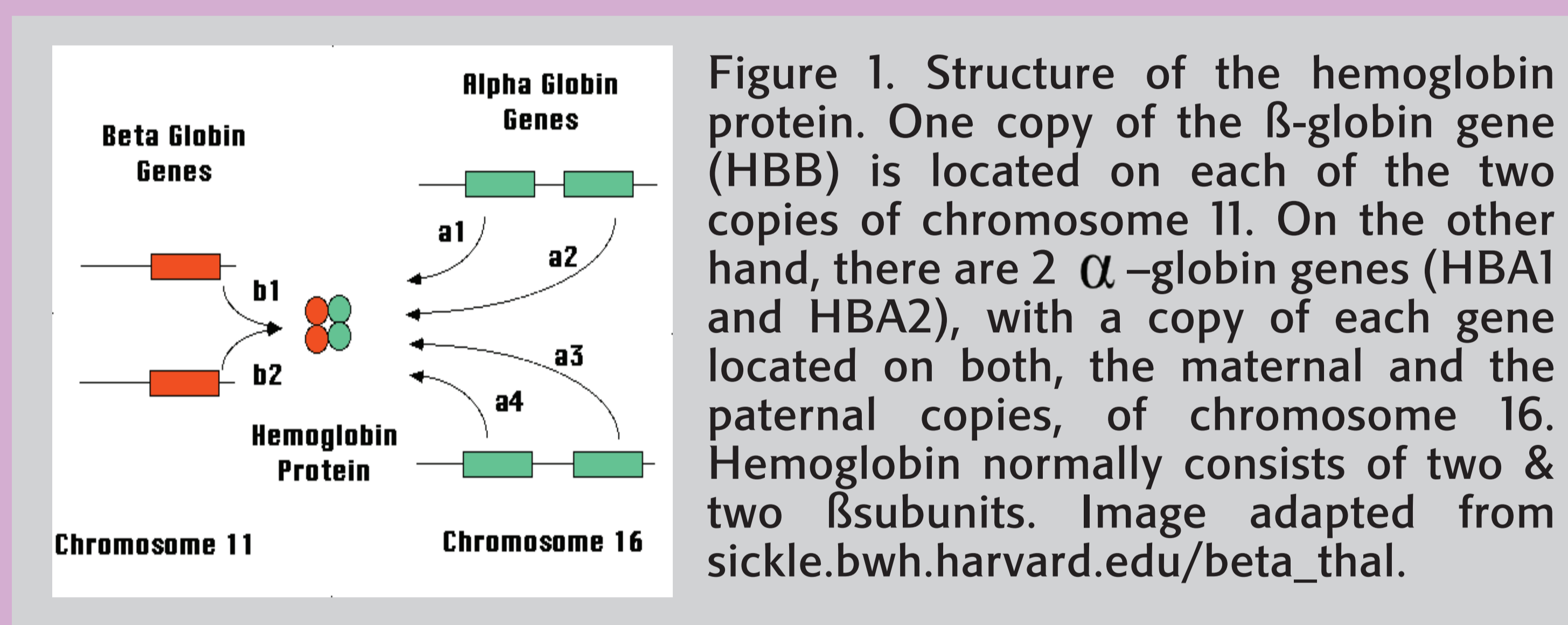


Figure 1. Structure of the hemoglobin protein. One copy of the β-globin gene (HBB) is located on each of the two copies of chromosome 11. On the other hand, there are 2 α-globin genes (HBA1 and HBA2), with a copy of each gene located on both, the maternal and the paternal copies, of chromosome 16. Hemoglobin normally consists of two & two β subunits. Image adapted from sickle.bwh.harvard.edu/beta_thal.

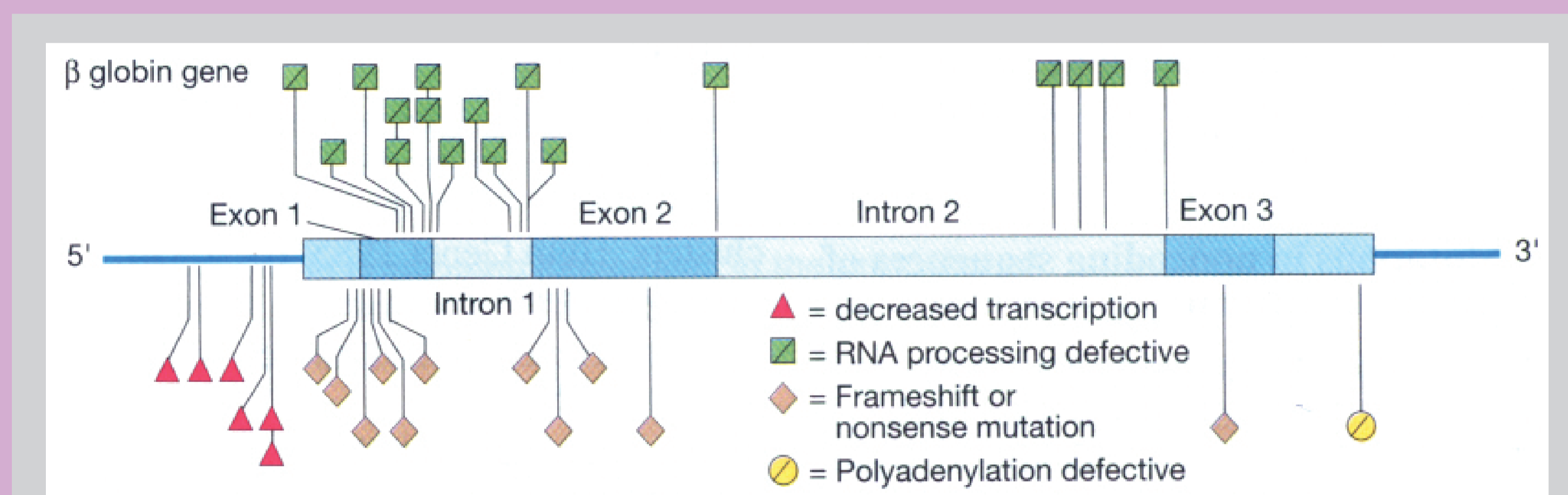


Figure 2. Mutations of the β-globin gene (HBB). HBB consists of 3 exons, with mutations occurring in both the intronic and exonic structures. Decreased transcription of the gene results from mutations occurring upstream of the HBB open reading frame (ORF) region.

Mode of inheritance

β-thalassemia is inherited in an autosomal recessive manner. Provided that both of the parents are carriers (carrying one normal and one mutated copy of the HBB gene), each of their offspring has a 25% chance of being affected (inheriting two mutated copies of the HBB gene, one from each parent), 50% chance of being an asymptomatic carrier (inheriting a normal copy from one parent and the mutated copy of the HBB gene from the other parent) and 25% chance of being unaffected and not a carrier.

Conclusion

The development of safe prenatal testing methods for the thalassemia has evolved in Jordan despite an atmosphere of profound ethical, legal and social concerns. Prenatal diagnosis for β-thalassemia was the first preventative genetic service to be set up in Jordan. The PCR-based methods utilized have been found to have a low error rate (0.1%) in comparison with other fetal diagnostic methods such as globin chain synthesis and southern blot analysis. Gene sequencing has also been successfully utilized at NCDEG, in cases where mutations could not be identified using the conventional PCR method. In our service, no misdiagnoses were reported. Implementing the obligatory pre-marital screening together with prenatal diagnosis for β-thalassemia in Jordan are aimed at reducing the rate of occurrence and eventually prevention of β-thalassemia. Continued progress depends on enhanced awareness of prenatal diagnosis and improved access to reproductive genetics services.

References

- Hammamy, H., Al-Hait, S., Alwan, A. and Ajlouni, K. (2007). Jordan: Communities and Community Genetics. *Community Genetics* 10, 5260-.
- Panigrahi, I. and Agarwal, S. (2008). Genetic determinants of phenotype in beta-thalassemia. *Hematology* 13, 24752-.
- Sadiq, M. F., Eigel, A., and Horst, J. (2001). Spectrum of β-Thalassemia in Jordan: Identification of Two Novel Mutations. *American Journal of Hematology* 68, 1622-.
- Weatherall J., Pressley L., Wood G., Higgs R. and Clegg B. (1981). Molecular basis for mild forms of homozygous beta-thalassemia. *Lancet* 1, 5279-.

Table 1. The summary of the results obtained from the molecular analysis of 324 CVS.

β	Number of CVS samples/324	Percentage %
β	83	25
β	175	53.5
β	22	6.4
β	44	13.4