

In the name of God



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Prevalence and molecular characterization of alpha-thalassemia and Detection of Hb Bart's and Hb H diseases using capillary electrophoresis among newborns in Ardabil Province

Introduction

- **Apha-thalassemia** is one of the **most** recessively **congenital** hemoglobin disorders in the world, which is characterized by decreased or absence of alpha globin chains production . Worldwide, up to around 5% of the general population are affected . **Although** the frequency of **α -thalassemia** in Iran is **greater** than it is **worldwide**, it is **not** exactly determined .

- Alpha-thalassemia syndrome results from deletion or mutation of one or both α -globin genes that are located on 16p13.3 . The phenotype of the disease is related to the number of α -globin genes deficiency as well as the type of mutation, ranging from asymptomatic phenotype to a fatal in utero disease . The loss of **one ($-\alpha$) and two ($-$) genes** of α -globin as the most **common cause** of α -thalassemia are not associated with clinical symptoms. While, the loss of three genes of α -globin ($--/-\alpha$) leads to hemoglobin H disease, which sometimes results in moderate anemia and the need for transfusion, the loss of four genes of α -globin ($--/--$) leads to Hb Bart's Hydrops Fetalis Syndrome, which can be fatal .

- Worldwide, **more than 770 mutations** have been recognized in α -globin gene clusters, and of those, **more than 19 mutations** documented **in Iran** . Gene sequencing and Multiplex Ligation-dependent Probe Amplification (**MLPA**) techniques can be useful for better detection of novel suspected α -thalassemia mutations . the aim of the present study was to determine the prevalence and molecular characteristic of α -thalassemia among newborns in **Ardabil Province**.

- The diagnosis of α -thalassemia is based on the diagnosis of **Hb Bart's** in infants, which indicates one or more defective or missing α -globin genes . Bart's hemoglobin level was found to be correlated with the number of defective globin α genes and is used to screen alpha thalassemia in infants . Hb Bart's has been reported to have a **greater affinity for oxygen** and therefore is unable to deliver effective oxygen to tissues . Hb Bart's levels in carriers of α genes **range from zero** to a small amount up to **1%** and may even be found in people with normal α genes . Because **Hb H** is **fast moving**, and also Hb H is **unstable** and may **not** be **detected** by Hb electrophoresis.

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- In general, previous research has shown that Hb Bart's and H in cord blood can be used as a suitable marker of alpha thalassemia, so that Bart hemoglobin levels increase in proportion to defective α genes .
 - However, the effectiveness of this method has never been tested in our population. In this study, we used **molecular analysis** an **acetate cellulose** electrophoresis evaluations to detect alpha thalassemia from cord blood. Finally, our group determined the probability of α -thalassemia in Bart's and H hemoglobin screening program.

1. Material and Methods

- In this cross-sectional study, **one thousand** newborns were screened for α -thalassemia in the pediatric unit (the only research center for thalassemia cases) of Bu-Ali hospital in Ardabil city between April 2016 and March 2018.
1. Using EDTA-containing tubes, a total of 2 ml venous blood was taken from each infant for analysis during routine hematological tests (**CBC**, Htc, Hb, MCH and MCHC as well as **ferritin**) and the remainder was kept at 4°C and sent for laboratory analysis in the first three days of life. There is a general agreement that the α -thalassemia trait is considered typical and could be present when **MCV** (Mean Corpuscular Volume) is **below 94FL** and **MCH** (mean corpuscular hemoglobin) is **below 30** pg, respectively, except in neonates with iron deficiency

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- . But in the current study, cases with **MCV** and **MCH** below **100 fL** and **33** pg respectively, were referred to laboratory for measurement of serum ferritin level, **acetate cellulose** electrophoresis and **molecular analysis**. Molecular techniques have been applied to identify genetic mutations and genomic DNA was extracted from whole blood.
 - Collected data were analyzed by statistical methods in IBM© SPSS© Statistics version 21 (IBM© Corp., Armonk, NY, USA).

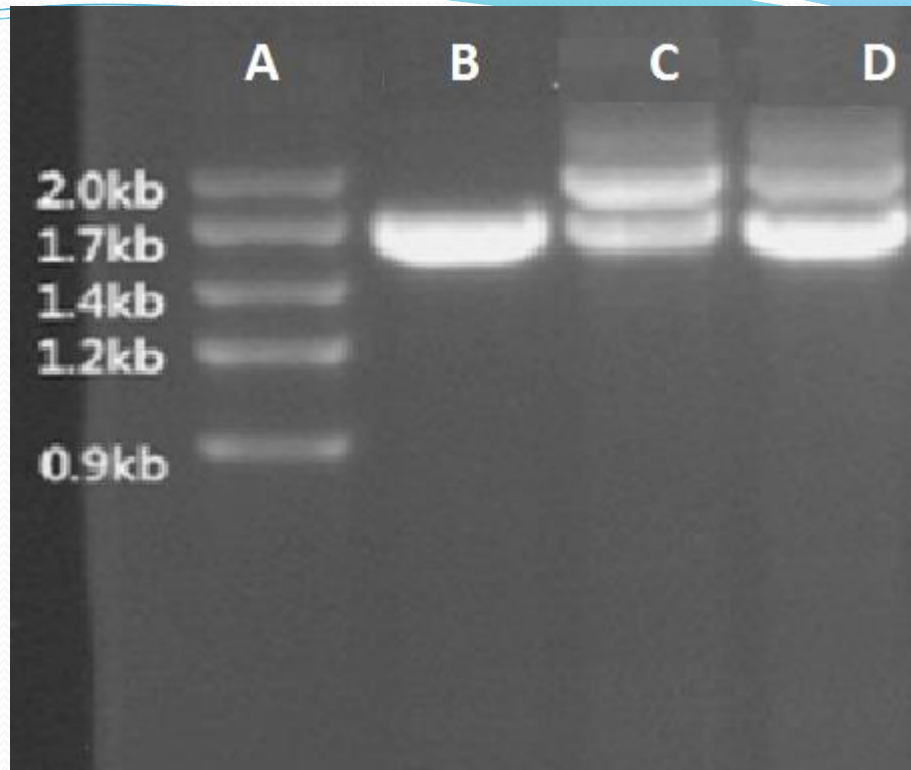


Figure 1. Multiplex PCR for screening of the most common α -thal alleles. A- Ladder marker; B- Natural genotype control ($\alpha\alpha / \alpha\alpha$); C- patient ($\alpha -3.7 / \alpha\alpha$); D-positive control to remove $\alpha -3.7$.

Results

- In total, 97 newborns had $MCV < 100$ fL and $MCH < 33$ pg. Hb electrophoresis, serum ferritin level was carried out on all cases. Based on these results, each suspected newborn was referred for genetic analysis. In summary, we found 33 newborns with α -thalassemia based on genetic analysis. The prevalence of α -thalassemia in studied newborns was 3.3 % in Ardabil province. The most common mutation was the 3.7 single gene deletions that were found in 42.4% (14 cases) of newborn with α -thalassemia. HbH was only detected in two cases (14.7% and 15.6%) in total (6.06%) and Hb Bart's was seen only in three neonates (2.5%, 4.5% and 5.5%) in total (9.09%) (Table 1,2).

Number	RBC	Hb	Htc	MCV	MCH	MCHC	HbA	HbA ₂	HbF	Ferritin
33	4.50 ± 0.52	13.68±1.82	41.04 ± 5.47	92.618 2± 3.70	31.20±1.63	33.78±0.82	22.36 ± 4.84	0.97 ± 0.41	73.53 ± 12.2	187.5 ± 75.5

Table 1. The mean of hematological parameters in studied samples

HbA: Hemoglobin A; HbA₂: Hemoglobin A₂; HbF: Fetal hemoglobin; MCH: Mean Corpuscular Hemoglobin; MCHC: Mean Corpuscular Hemoglobin Concentration; MCV: Mean Corpuscular Volume

Genotype	N	%	HbA2 (%)
- $\alpha^{3.7} / \alpha \alpha$	14	42.4	0.8
- $\alpha^{3.7} / - \alpha^{3.7}$	6	18.2	0.72
aaa anti $\alpha^{3.7} / aa$	2	6.1	1
- $\alpha^{PA2} / \alpha \alpha$	1	3.0	0.9
- $\alpha^{4.2} / \alpha \alpha$	4	12.2	1.75
$\alpha 2$ IVS1(-5nt)	1	3.0	0.8
$\alpha 2$ cd19 [-G] het	1	3.0	1.5
het deletion med1	1	3.0	0.8
HET.C.427T>C at hbA2	1	3.0	2.9
del G at codon 126/Wt	1	3.0	1.8
het -a 20.5	1	3.0	1.5

Table 2. The frequency of found mutations in studied samples

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Discussion

- Although no compressive study has been performed about the prevalence of α -thalassemia in Iran, the rate of α -thalassemia carriers was estimated to be about 7.8% in the general population. Since the Iranian population is a mixture of different ethnic group, the rate of α -thalassemia can be different in different regions. Alpha-thalassemia is more prevalent around the Caspian Sea and Persian Gulf (North, and South of Iran). The rate of α -thalassemia differs from 2.7% (in East Azerbaijan; north-west of Iran) to 15.3% (in Mazandaran; northern Iran). Alpha-thalassemia is prevalent in populations in South East Asia, the Mediterranean and the Middle East. It has been reported that the prevalence of α -thalassemia in Thailand, Morocco and turkey is 30-40%, 0.96% and 0.25%-4.1%, respectively.



- The prevalence of α -thalassemia in the current study was 3.3%. Our results are similar to the study of East Azerbaijan province, in the north-west of Iran. Data showed that $-\alpha^{3.7}$ and $-\alpha^{4.2}$ are the most common mutation, causing α -thalassemia among 10,849 cases of α -thalassemia carriers in a comprehensive study of Iran. Like other regions, $-\alpha^{3.7}/\alpha\alpha$ and $-\alpha^{3.7}/-\alpha^{3.7}$ was the most common mutation in this study, followed by $-\alpha^{4.2}/\alpha\alpha$. The second most common mutation in other studies was $aa/-\alpha^{4.2}$, poly A2, deletion $-\text{MED}$, $\alpha\text{-5 nt}$. The most common mutation in the world is $-\alpha^{3.7}$ which was similar to our study results.



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- The mean of **MCV** and **MCH** was 92.62 ± 3.7 and 31.20 ± 1.63 , respectively. So, by using **these cut-off** points (**MCV < 94** and **MCH < 30**), a large number of cases are **not detected**.

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- The **values** of MCV and MCH are strongly affected by the **number** of α -genes, cases with one α gene deletion have slightly decreased of these parameters and can be **overlapped** with **normal** values .
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- The mean of HbA2 and HbF were 0.97 ± 0.4 and 73.53 ± 12.3 , respectively. HbH was only detected in two cases (14.7% and 15.6%) and Hb Bart's was seen only in 3 neonates (2.5%, 4.5% and 5.5%). The level of HbA2 in newborns with α -thalassemia carrier is normal or lower. Especially in HbH disease, it can drop to less than 2% (94% of cases in our study had HbA2 below 2%).

- Remarkably, the mean HbA2 level in this study was 0.97 ± 0.41 . HbA2 levels in infants with thalassemia carriers are normal or slightly lower, which can distinguish α -thalassemia from thalassemia in particular. In Hb H disease, HbA2 levels can be reduced to less than 1. Normal or low HbA2 levels combined with decreased mean body hemoglobin (MCH) ($<33\text{pg}$) and mean cell volume (MCV) ($<100\text{fl}$) may be a good chance to detect α -thalassemia carriers.



. Alpha-thalassemia carriers can be identified by detection of Hb Bart's in newborns. The level of Hb Bart's is found in a large number of newborns with α -thalassemia and it is correlated with the number of defective α -globin genes. In this study, Hb Bart's was detected only in three cases (9.09%) and the majority of cases were negative. It is important to note that the absence of Hb Bart's does not exclude the presence of α -thalassemia in newborns (especially in mild α -3.7/ $\alpha\alpha$ interactions) and diagnosis can only be confirmed by the molecular analysis.

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- Hb Bart's disease is not accurately diagnosed in infancy because it disappears quickly after birth. Therefore, determining the amount of Hb Bart's after infancy is not reliable . On the other hand, Hb Bart's levels have been shown to vary between different ethnic groups .

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- Hb H and Hb Bart's Hb are also fast moving, appearing on electrophoresis. As a result, some reports indicate that these two parameters are unstable and may not be detectable by conventional methods .Therefore, it seems that molecular analysis of is necessary to confirm α -thalassemia.

- In this study, only five cases (15.15%) were detected by Hb Bart's and also Hb H by electrophoresis. Therefore, most infants with α -thalassemia are missed when electrophoresis alone is used. In some normal infants, Hb Bart's can be detected in about 0.5-1.5% of cases. Various factors may play a role in the variability of Hb Bart levels, and among them, the amount of γ - β globin change may play an important role.

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- Our findings show that all infants with two alpha gene defects had elevated Hb Bart , while a large proportion of single-defect alpha gene carriers do not show traceable Hb Bart.

- Previous studies have shown that the $-\alpha 4.2$ allele causes a considerable synthesis imbalance of α -non- α -globin chains and the considerable production of the γ chain in patients than the $-\alpha 3.7$ allele. According to such results, the level of Hb Bart in newborns with $-\alpha 3.7$ was 0.2 ± 0.5 was, while in newborns with $-\alpha 4.2$, the level of Hb Bart was represented in 0.3 ± 0.7 . Therefore, our group guessed that a small amount of Hb Bart related to the $-\alpha 3.7$ allele is not technically reliable, while the $\alpha 4.2$ allele is reliably detectable due to the high level of hemoglobin Bart's.

Conclusions

- Results of this study showed that, the prevalence of α -thalassemia was 3.3% in Ardabil province. The most common mutation causing α -thalassemia in this study was $-\alpha^{3.7}/\alpha\alpha$ (42.4%). Although the evaluation of Hb Bart's and Hb H are direct methods for the diagnosis of α thalassemia , only a few cases of Alpha- thalassemia in this study were detected by Hb Bart's and Hb H (15.15%) in acetate cellulose electrophoresis and most of them were missed. Therefore molecular analysis in suspected infants is necessary to confirm α -thalassemia.

Thanks For your Attention

