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ABSTRACT SHARING

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THE UNIQUE BETA ZERO SOUTH EAST ASIA DELETION (β°)-THAL SEA; β° -THALASSAEMIA OR HEREDITARY PERSISTENT FETAL HAEMOGLOBIN (HPFH)?

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Background

We report hematological parameters and molecular characterization of beta zero (β°) South East Asia (SEA) deletion with unusually high levels of Hb F as compared to a classical beta-thalassemia trait.

Materials and Methods

Retrospective analysis of 303 cases presented with high Hb F level (>5%) based on hemoglobin analysis were done to exclude Hereditary Persistent Fetal Haemoglobin (HPFH) and Delta-Beta Thalassemia ($\delta\beta$) by GAP-PCR. Seventeen cases were positive for the β° - SEA deletion. The mutation was analysed and the results were compared between another beta zero thalassaemia group (β° -FIL, β° -IVS1-1, β° - 3.5kb, Codon41/42), $\delta\beta$ (THAI) and HPFH group (HPFH-6 and Siriraj).

Results

Among 303 samples investigated, 228 (75.2%) were heterozygous, 22 (7.3%) were homozygous and 53 (17.5%) of them were found to be co-inherited with different types of β -thalassemia mutations. Of these 228 heterozygous cases, β^o -thalassemia mutations were identified in 148 (64.9%); 69 (46.6%) with Filipino deletion, 39 (26.4%) with Codon 41/42 mutation, 16 (10.8%) with IVS1-1 mutation, 15 (10.1%) with SEA deletion and 9 (6.1%) with 3.5Kb deletion. The demographic data including state and ethnicity of β^o -SEA deletion cases were analysed. The results showed that β^o -SEA deletion was frequently received from Sarawak with n= 5 (29.4%), followed by Sabah with 3 (17.6%) and Selangor with 3 (17.6%). Overall, Chinese had the highest number of β^o -SEA deletion cases with n=10 (58.8%), followed by Bidayuh with 4/17 (23.5%) and Sino with 1/17(5.9%). These data demonstrated that β^o -SEA deletion was not uncommonly reported in Malaysia especially in Sarawak, and commonly seen in the Chinese population.

The hematological parameters and RBC indices of heterozygous $\beta^o\text{-SEA}$ deletion were significant differences from other $\beta^o\text{-thalassemia}$ mutations except for RBC. The mean Hb for heterozygous $\beta^o\text{-SEA}$ deletion (13.43 \pm 1.40 g/dL) was significantly higher than heterozygous IVS 1-1 and Codon 41/42 (Post hoc test, p<0.05). As compared to other β^o mutations group, the mean Hb level for $\beta^o\text{-SEA}$ is within normal limit with hypochromic microcytic red cell similar to $\delta\beta$ and HPFH groups. The median for MCV and MCH of $\beta^o\text{-SEA}$ deletion was significantly higher than other heterozygotes $\beta^o\text{-thalassemia}$ traits (Man Whitney test, p<0.05).

The patient with β° -thal SEA deletion had Hb A2 level of classical β -thalassemia trait with Hb F level of classical HPFH or $\delta\beta$ carrier. The median for HbA2 was (4.00+1.00%) similarly observed in other β° -thalassaemia group except IVS 1-1 mutation (median of 5.30+ 0.45%) and it was statistically significant (Mann Whitney test, p<0.05). Interestingly, we found the HbF level for β° -thal SEA deletion was statistically higher as compared to other β° mutation with median of (19.00+ 5.50%, p<0.05) except for 3.5kb deletion group.

Conclusion

We conclude that β° -SEA deletion is a unique beta zero thalassaemia trait. This cases provides valuable information for accurate genetic counselling. Although the deletion region only involve the beta globin gene, it presented as a phenotype of HPFH. The possible mechanism of such variation will further be elucidated.

Key wards: β-thalassemia, HPFH, δβ-thalassemia