

Molecular and clinical characteristics of Hemoglobin Ottawa detected in a Chinese population

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Abstract. Hemoglobin (Hb) Ottawa [α 15 (A13) Gly→Arg], also known as Hb Siam, was first described in an 82-year-old Canadian in 1974. The same year, a second case was reported in a 28-year-old Chinese male living in Thailand. A third case was found in 1986 in a Chinese female living in the Hubei province of China. Since then, there have been no reports of Hb Ottawa in mainland China in the English literature. Hb Ottawa results from a GGT→CGT mutation in codon 15 of the α_1 or α_2 -globin gene. Hb Ottawa carriers do not present any clinical symptoms or hematological changes, and are often diagnosed during a health examination and thalassemia screening. In a hemoglobin survey of 9745 students in Chaozhou, Guangdong, China, we identified four cases of Hb Ottawa in a thalassemia screening by Hb electrophoresis, and confirmed it to be the result of a GGT→CGT mutation in codon 15 of the α_2 -globin gene by DNA sequence analysis.

Introduction

Hemoglobin (Hb) Ottawa [α 15 (A13) Gly→Arg], also known as Hb Siam, was first identified in an 82-year-old Canadian (1). The same year, a second case was reported in a 28-year-old Chinese male living in Thailand (2). According to the human hemoglobin variant database (HbVar) (<http://globin.cse.psu.edu>), Hb Ottawa has only been found in individuals of Polish-Canadian and Chinese ethnicity. Hb Ottawa results from a GGT→CGT mutation in codon 15 of the α_2 -globin gene (1-3), although some studies have reported that it is a result of a GGT→CGT mutation in the α_1 -globin gene (4,5). Hb Ottawa carriers do not present any clinical symptoms or hematological changes, and are often diagnosed in thalassemia screening by Hb electrophoresis, confirmed by DNA sequence analysis.

In 1986, a case of Hb Ottawa was found in a Chinese female in the Hubei province of China. This was the first

report of Hb Ottawa in China (3). Since then, there have been no reports of Hb Ottawa in mainland China in the English literature. Recently, during a hemoglobin survey of 9745 students in Chaozhou, Guangdong, China, we identified four cases of Hb Ottawa in a thalassemia screening by Hb electrophoresis, and confirmed it to be the result of a GGT→CGT mutation in codon 15 of the α_2 -globin gene by DNA sequence analysis.

Materials and methods

Materials. During a hemoglobin survey of 9745 students aged 7-19 years in Chaozhou, Guangdong, China, four cases of abnormal hemoglobin were found by Hb electrophoresis.

Hematological analysis. The specimens were analyzed by cellulose acetate electrophoresis (pH 8.6), and the percentage of Hb variants was measured by densitometry on alkaline electrophoresis (6). Hb stability was examined by the isopropanol and heat methods (7,8). Hb variants were dissociated by parachloro-mercuri-benzoate (PCMB), which is capable of differentiating α - or β -globin mutations (9).

DNA analysis. Genomic DNA was extracted from the peripheral blood leukocytes of the specimens using the genomic DNA mini-preparation kit (Decipher Bioscience Shenzhen Ltd.) according to previously described methods (10). The α_1 and α_2 globin genes were separately amplified by polymerase chain reaction (PCR) in the DNA thermal cycler MJ mini (Bio-Rad, Hercules, CA, USA). The primers and products lengths are shown in Table I, and were determined as described previously (11,12).

The α_1 -globin gene was amplified in 50 μ l PCR reaction mixtures containing 0.1 μ g DNA, 15 pmol primers, 200 μ mol dNTPs and 2.5 units Taq DNA polymerase (Shanghai Sangon Biological Engineering Technology and Services Co., Ltd., P.R. China) in 10 mmol/l Tris-HCl (pH 8.0), 50 mmol/l KCl and 3 mmol/l $MgCl_2$. PCR conditions were as follows: after an initial denaturation at 95°C for 3 min to activate the DNA polymerase, 35 cycles of PCR at 98°C for 40 sec, 60°C for 5 sec and 72°C for 1 min were carried out, followed by a final extension at 72°C for 5 min. The amplification process was similar for the α_2 -globin and α_1 -globin genes. The amplified products were separated on 2% agarose gels and detected under UV light after staining with ethidium bromide. The

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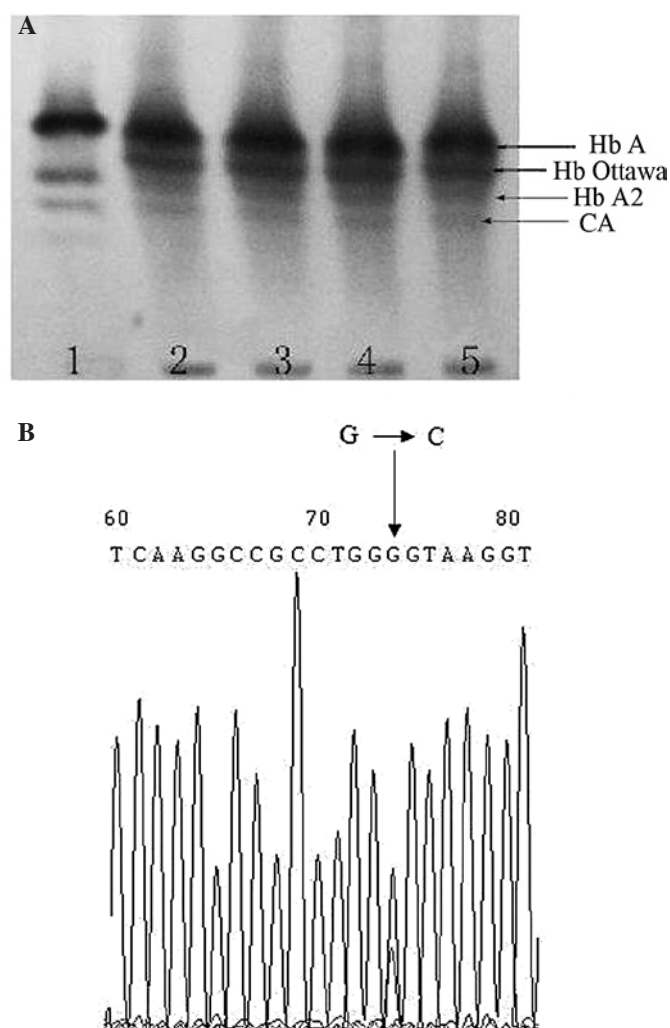


Figure 1. Hemoglobin electrophoresis and DNA sequencing analysis. (A) Hemoglobin analysis with cellulose acetate electrophoresis at pH 8.6. 1, Normal control; 2, Case 1; 3, Case 2; 4, Case 3; 5, Case 4; (B) DNA sequence analysis of the amplified α_2 -globin gene. The downward arrow indicates the G-C substitution in codon 15 of the α_2 -globin gene.

Table I. Primers and length of globin gene amplification.

Primers	Sequences (5'-3')	PCR product length (bp)
α_1 -B	CCATGCCTGGCA CGCTTTGCTGAG	880
α_1 -L	TCCCCACAGACT CAGAGAGAACC	
α_2 -D	AACACCTCCATTC GTTGGCACATTC	880
α_2 -L	TCCCCACAGACT CAGAGAGAACC	

PCR products were analyzed by DNA sequencing with the ABI 3700 automated sequencer.

Results

During a hemoglobin survey of 9745 students aged 7-19 years in Chaozhou, Guangdong, China, four cases of abnormal hemoglobin were identified by Hb electrophoresis. The subjects had no clinical symptoms or hematological changes. The hematological data are summarized in Table II. The Hb variant patterns were revealed by Hb cellulose acetate electrophoresis at pH 8.6 (Fig. 1A). This Hb variant (about 15% of total Hb) migrated toward the anode faster than Hb A2 and slower than Hb A, and the distance between it and Hb A2 was equal to that of the distance between it and Hb A. The results of Hb stability tests were normal. After the α_1 and α_2 -globin genes were amplified and their PCR products were sequenced, a transition from G→C in codon 15 of the α_2 -globin gene was observed (Fig. 1B). These results revealed that the identified subjects were Hb Ottawa carriers.

Table II. Summary of hematological findings.

Parameter	Case 1	Case 2	Case 3	Case 4
RBC ($10^{12}/l$)	5.30	5.38	4.9	4.99
Hb (g/l)	145.00	160.00	136.0	132.00
MCV (fl)	87.30	87.40	84.4	89.30
HCT (%)	46.80	47.00	41.3	44.50
MCH (pg)	27.30	29.70	27.7	26.40
MCHC (g/l)	313.00	340.00	329.0	296.00
RDW (%)	13.30	13.60	15.7	13.50
Hb A (%)	83.40	81.90	79.8	82.70
Hb A2 (%)	2.50	2.60	2.8	2.50
Hb Ottawa (%)	14.10	15.50	17.4	14.80
Genotype	$\alpha\alpha/\alpha\alpha^{\text{Ottawa}}$ β/β	$\alpha\alpha/\alpha\alpha^{\text{Ottawa}}$ β/β	$\alpha\alpha/\alpha\alpha^{\text{Ottawa}}$ β/β	$\alpha\alpha/\alpha\alpha^{\text{Ottawa}}$ β/β

RBC, red blood cell count; Hb, hemoglobin; MCV, mean corpuscular volume; HCT, hematocrit; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; RDW, red blood cell distribution width.

Discussion

Guangdong province is a high-incidence area of thalassemia and abnormal hemoglobin. During this hemoglobin survey of 9745 students in the Chaozhou area of Guangdong, the incidence of Hb Ottawa was found to be 0.41% (4/9745). Hb G-Chinese [α 30 (B11) Glu→Gln], Hb G-Waimanalo [α 64 Asp→Asn] and Hb Ottawa have the same electrophoresis phenomenon (11,13). All are stable hemoglobin variants and their carriers usually do not present any clinical symptoms or hematological changes. Therefore, the differential diagnosis of Hb Ottawa from Hb G-Chinese and Hb G-Waimanalo mainly depends on the DNA sequence analysis of the amplified α_2 -globin gene. High performance liquid chromatography (HPLC), multiplex allele specific PCR, high-resolution melting (HRM) or polymerase chain reaction and restriction fragment length polymorphism (PCR-RFLP) can also be applied to the differential diagnosis of Hb Ottawa (3,14,15). In addition, we confirmed that the cases of Hb Ottawa identified in our population were the result of a GGT→CGT mutation in codon 15 of the α_2 -globin gene, not the α_1 -globin gene as previously described (4,5). Diagnosis of these relevant abnormal hemoglobins is difficult to achieve by routine screening methods. The development of novel methods would be useful for providing an accurate and fast diagnosis.

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