

The prevalence and outcomes of α - and β -thalassemia among pregnant women in Hubei Province, Central China

An observational study

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Abstract

There is no information concerning the prevalence of thalassemia among pregnant women in Hubei Province currently. This study is aimed to explore the prevalence of α - and β -thalassemia genotypes among pregnant women in Hubei Province, and to explore the clinically applicable screening approach, as well as to investigate the pregnancy outcomes of α - and β -thalassemia carriers.

Pregnant participants were recruited from 4 hospitals for the screening of α - and β -thalassemia mutations in Hubei Province. Polymerase Chain Reaction and flow cytometry methods were used to examine α - and β -thalassemia mutations. The hematological parameters and pregnancy outcomes of α - and β -thalassemia carriers were obtained from the hospital information system. The chi-square tests were used to evaluate the difference in hematological parameters between pregnant thalassemia carriers and the control group.

Among 11,875 participants, 414 (3.49%) were confirmed with α -thalassemia carriers, 228 (1.92%) were confirmed with β -thalassemia carriers, and 3 (0.03%) were confirmed with both α - and β -thalassemia carriers. The frequency of $-\alpha^{3.7}$ accounted for 2.05% and it was the most frequent genotype of α -thalassemia; the proportion of IVS-II-654 was 0.85% and it was the most frequent genotype of β -thalassemia in Hubei Province. Furthermore, the proportion of patients with low mean corpuscular volume (MCV) or mean cell hemoglobin (MCH) values was accounted for 36.64% and 93.97% among α -thalassemia and β -thalassemia carriers, respectively. And participants with normal MCV and MCH values were accounted for 95.07% among non-thalassemia participants. High prevalence of pregnancy-induced diabetes (16.97%), preterm birth (9.96%), pregnancy-induced hypertension (8.12%), and low birth weight (5.90%) were observed among pregnant thalassemia carriers.

MCV and MCH values were suggested to apply on the preliminary screening of pregnant β -thalassemia; however, it's unpractical on that of α -thalassemia. Furthermore, thalassemia carriers might have a high risk of negative pregnancy outcomes. These findings could be useful for the preliminary screening of thalassemia and perinatal care for the pregnant thalassemia carriers.

Abbreviations: MCH = mean cell hemoglobin, MCV = mean corpuscular volume, TI = thalassemia intermedia.

Keywords: Hubei, pregnant, prevalence, screening, thalassemia

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1. Introduction

Normal hemoglobin of hemoglobin A is a tetramer composed of a pair of α -globin chains and a pair of β -globin chains. Thalassemia is a group of diseases characterized by the reduced or deficient synthesis of one or more globin chains. As one of the most common hereditary disorders, there are great differences in the distribution of thalassemia in different areas. According to the Global Burden of Disease, the 3 countries with the highest prevalence of thalassemia mutations were Thailand (12.71%), Cambodia (12.17%), and Lao People's Democratic Republic (11.48%) in 2019.^[1] And the prevalence of thalassemia mutations in China ranked fifth in the world (9.79%).^[1] Moreover, thalassemia was reported to be prevalent in southern China.^[2,3] The Li people in Hainan Province were reported to have a high prevalence of thalassemia (65.27%).^[4] According to a meta-analysis based on 16 studies in mainland China, the prevalence of β-thalassemia was 0.01% to 1.59% in Hubei Province, but there is no corresponding supportive information on the prevalence of thalassemia among pregnant women in Hubei Province currently.^[5]

In developed areas of China, thalassemia genetic screening has been regarded as one of the routine prenatal examinations, but not in most developing areas in China. As other diseases, clinical practical approaches should be promoted to generalize the thalassemia screening.^[6,7] Furthermore, due to advances in treatment technology, the life expectancy of thalassemia patients has been greatly increased, and patients with thalassemia static, thalassemia minor, or thalassemia intermedia (TI) have no significant effect on the overall life expectancy.^[8] However, these thalassemia patients are prone to a variety of complications, including chronic hemolytic anemia, hypoparathyroidism, cirrhosis, heart failure, pulmonary hypertension, thrombosis, and so on.^[9-12] Pregnant thalassemia patients might have no lifethreatening symptoms, their pregnancy outcomes still warrant attention. A case-control study conducted by Vafaei et al^[13] reported that the frequency of neonates with low birth weight was significantly higher among women with B-thalassemia minor. Huang et al^[14] reported that the incidence of preterm birth and low birth weight were 6.5% and 7.3% in the thalassemia carriers. Whereas current studies focused on pregnancy outcomes among thalassemia carriers are insufficient.

For further understanding of thalassemia among pregnancies, this study is aimed to explore the prevalence of α - and β -thalassemia genotypes among pregnant women in Hubei Province, and to explore the clinically applicable screening approach, as well as to investigate the pregnancy outcomes of α - and β -thalassemia carriers.

2. Methods

2.1. Population

Hubei Province of Central China has 13 prefecture-level administrative regions and 4 county-level administrative regions. The Eastern region includes Huangshi City, Ezhou City, Huanggang City, Xianning City, and Xiaogan City; the Central region includes Wuhan City, Jingzhou City, Jingmen City, and the county-level administrative regions; the Western region includes Yichang City, Shiyan City, Xiangyang City, Suizhou City, and Enshi City. This study was approved by the Ethics Committee of Maternal and Child Health Hospital of Hubei Province on Sep 3, 2018. In this study, 4 hospitals providing medical services for patients mainly from the Eastern, Central, and Western regions of Hubei Province were selected. Pregnant women were recruited in these hospitals to participate in the screening of α - and β -thalassemia mutations from January 2019 to November 2020, and they were further classified by residence address. The least sample size in this study was calculated by the following determination formulas: $n=t^2 \times P(1-P)/d^2$, t=1.96, P=4%, d=0.4%, $n\approx9220$, and the current study recruited a total of 11,875 eligible participants.

2.2. Genetic analysis

In this study, thalassemia (α/β) Gene Diagnostic Kit (Polymerase Chain Reaction [PCR]- Flow cytometry fluorescence Hybridization Assay) was used to detect thalassemia genotypes. PCR primers and hybridization probes were designed for the hot spots of thalassemia mutation. A total of 20 point-mutations includes 3 α -mutations (WS122, QS125, and CS142) and 17 β -mutations (CD41-42, IVS-II-654, CD17, -28, CD26, CD71-72, CD43, -29, Int, CD14-15, CD27-28, -32, -30, IVS-I-1, IVS-I-5, CD31, and CAP). Moreover, a total of 3 deletion mutations (—sea, - α 3.7, and - α 4.2) and 1 normal controlling genotype were amplified. PCR and flow cytometry methods were used to examine α - and β -thalassemia mutations.

2.3. Data collection and analysis

The prevalence of different thalassemia alleles among pregnant women was described in Hubei Province stratified by East, Central, and West. The hematological parameters including red blood cell, hemoglobin, hematocrit, mean corpuscular volume (MCV), mean cell hemoglobin (MCH), and mean corpuscular hemoglobin concentration were collected. Thalassemia carriers with a singleton pregnancy were selected to follow up for pregnancy outcomes including preterm birth, pregnancy-induced hypertension, pregnancy-induced diabetes, eclampsia/preeclampsia, birth defect, miscarry, stillbirth, fetal weight, and Apgar scores. Data of the hematological parameters and pregnancy outcomes were obtained from the hospital information system. The chi-square tests were used to evaluate the difference in hematological parameters between α - and β -thalassemia carriers. Statistical analysis was conducted with SAS 9.4 (SAS Institute, Carv, NC, USA) for Windows. Two-sided P values <.05 were considered statistically significant.

3. Results

Among the 11,875 participants, 8150 (68.63%) provided the hematological parameters, 645 (5.43%) were diagnosed with thalassemia carriers, and 271 (42.02%) of the singleton thalassemia carriers were tracked with a pregnancy outcome in this study.

A total of 23 genotypes and 17 gene mutations of α- and β-thalassemia were identified. Among the thalassemia carriers, 414 (3.49%) were confirmed with α-thalassemia carriers, 228 (1.92%) were confirmed with β-thalassemia carriers, and 3 (0.03%) were confirmed with both α- and β-thalassemia carriers (Table 1). Three types of α-thalassemia static including $-\alpha^{3.7}$, $-\alpha^{4.2}$, and $\alpha^{WS}\alpha$ accounted for 69.81% of α-thalassemia carriers, and more details were provided in Table S1, Supplemental Digital Content, http://links.lww.com/MD2/A885. The frequency of $-\alpha^{3.7}$ accounted for 2.05% and it was the most frequent genotype accounting for more than half of the α-thalassemia among Eastern, Central, and Western of Hubei Province, Table 1

The prevalence and proportions of pregnant α - and β -thalassemia carriers in Hubei Province.

Subtypes	East N (%)	Central N (%)	West N (%)	Total N (%)
α -thalassemia				
$-\alpha^{3.7}$ (static)	57 (2.07/56.43)	116 (2.00/61.38)	71 (2.13/57.26)	244 (2.05/58.94)
— ^{SEA} (TM)	31 (1.13/30.69)	55 (0.95/29.10)	26 (0.78/20.97)	112 (0.94/27.05)
$-\alpha^{4.2}$ (static)	10 (0.36/9.90)	13 (0.22/6.88)	14 (0.42/11.29)	37 (0.31/8.94)
$\alpha^{WS}\alpha$ (static)	2 (0.07/1.98)	2 (0.03/1.06)	4 (0.12/3.23)	8 (0.07/1.93)
α ^{QS} α (TM)	0	2 (0.03/1.06)	5 (0.15/4.03)	7 (0.06/1.69)
α ^{CS} α (TM)	1 (0.04/0.99)	0	2 (0.06/1.61)	3 (0.03/0.72)
$-\alpha^{3.7}/-\alpha^{4.2}$ (TM)	0	1 (0.02/0.53)	0	1 (0.01/0.24)
$-SEA/\alpha^{WS}\alpha$ (TI)	0	0	1 (0.03/0.81)	1 (0.01/0.24)
$\alpha^{CS}\alpha/-\alpha^{3.7}$ (TM)	0	0	1 (0.03/0.81)	1 (0.01/0.24)
Total	101 (3.68/100)	189 (3.26/100)	124 (3.72/100)	414 (3.49/100)
β-thalassemia				
IVS-II-654 (TM)	28 (1.02/50.91)	41 (0.71/50.00)	32 (0.96/35.16)	101 (0.85/44.30)
CD41-42 (TM)	12 (0.44/21.82)	11 (0.19/13.41)	21 (0.63/23.08)	44 (0.37/19.30)
CD17 (TM)	7 (0.25/12.73)	16 (0.28/19.51)	17 (0.51/18.68)	40 (0.34/17.54)
CD27-28 (TM)	5 (0.18/9.09)	1 (0.02/1.22)	8 (0.24/8.79)	14 (0.12/6.14)
CAP (TM)	0	6 (0.10/7.32)	2 (0.06/2.20)	8 (0.07/3.51)
-28 (TM)	1 (0.04/1.82)	2 (0.03/2.44)	5 (0.15/5.49)	8 (0.07/3.51)
CD71-72 (TM)	1 (0.04/1.82)	2 (0.03/2.44)	1 (0.03/1.10)	4 (0.03/1.75)
CD43 (TM)	1 (0.04/1.82)	1 (0.02/1.22)	1 (0.03/1.10)	3 (0.03/1.32)
CD26 (TM)	0	1 (0.02/1.22)	2 (0.06/2.20)	3 (0.03/1.32)
-29 (TM)	0	1 (0.02/1.22)	1 (0.03/1.10)	2 (0.02/0.88)
CD14-15 (TM)	0	0	1 (0.03/1.10)	1 (0.01/0.44)
Total	55 (2.00/100)	82 (1.42/100)	91 (2.73/100)	228 (1.92/100)
$\alpha + \beta$ -thalassemia				
$CD27-28 + -\alpha^{3.7}$	0	1 (0.02/50.00)	0	1 (0.01/33.33)
$VS-II-654 + -\alpha^{3.7}$	0	1 (0.02/50.00)	0	1 (0.01/33.33)
$CD71-72 + \alpha^{QS} \alpha$	0	0	1 (0.03/100)	1 (0.01/33.33)
Total	0	2 (0.03/100)	1 (0.03/100)	3 (0.03/100)

TI = thalassemia intermedia, TM = thalassemia minor.

followed by —^{SEA} (0.94%) and $-\alpha^{4.2}$ (0.31%) genotypes. Three cases of 2 α -globin mutations including $-\alpha^{3.7}/-\alpha^{4.2}$, $-^{SEA}/\alpha^{WS}\alpha$, and $\alpha^{CS}\alpha/-\alpha^{3.7}$ were identified.

The prevalence of IVS-II-654 accounted for 0.85% and it was the most frequent genotype of β -thalassemia in the Eastern, Central, and Western of Hubei Province. Other high prevalence β -thalassemia genotypes were CD41-42 (0.37%), CD17 (0.34%), and CD27-28 (0.12%). Different from Eastern Hubei and Western Hubei, CD-17 accounted for 19.51% and it ranked second among all of the β -thalassemia genotypes in Central Hubei. Moreover, 1 TI case of —^{SEA}/ α ^{WS} α genotype and 3 cases carried both α - and β -globin mutations were identified, which were CD27-28 combined with $-\alpha^{3.7}$, IVS-II-654 combined with $-\alpha^{3.7}$, and CD71-72 combined with $\alpha^{QS}\alpha$.

Table 2 shows that there were 11.83% and 12.07% of α -thalassemia and β -thalassemia carriers suffered from low red blood cell (<3.810⁹/L), respectively. The majority of β -thalassemia carriers showed low hemoglobin (<110 g/L) (84.48%) and hematocrit (<0.35 L/L) (81.90%). Only 34.73% of α -thalassemia carriers were detected with low MCV (<82 fL), while corresponding proportion was 93.97% among β -thalassemia carriers. The β -thalassemia carriers also showed a higher proportion of low MCH (<27 pg) than that of α -thalassemia carriers (93.97% vs 35.88%). There were 93.97% and 36.64% of β -thalassemia carriers and α -thalassemia carriers presented low MCV or MCH, respectively. Meanwhile, 97.47% of non-thalassemia carriers presented normal MCV and MCH.

Table 3 shows that the most prevalent complication of pregnant thalassemia carriers was pregnancy-induced diabetes

(16.97%), followed by preterm birth (9.96%). Pregnancyinduced hypertension was also prevalent among α -thalassemia carriers (10.16%). The prevalence of eclampsia or preeclampsia was 2.95% among all of the thalassemia carriers. For neonatal outcomes, neonates with 1-minute Apgar scores <10 points comprised a proportion of 15.50%. Low birth weight neonates accounted for 5.88% and 5.95% among both α - and β -thalassemia carriers, respectively. The prevalence of birth defects among α - and β -thalassemia carriers accounted for 2.14% and 4.76%, respectively.

4. Discussion

This study firstly reported the frequency of α - and β -thalassemia carriers among pregnant women in Hubei Province, which was lower than the prevalence of thalassemia carriers among the general population of Chongqing (7.76%) and Shenzhen (6.49%),^[15,16] and this gap could be explained by the difference of targeted populations.

Moreover, this study presented the proportion of $-\alpha^{3.7}$ genotype ranked the first among all the α -thalassemia genotypes, in line with Chongqing China, Northern Thailand, and Southwest Iran.^[17–19] Diejomaoh et al^[20] also confirmed that $-\alpha^{3.7}$ was the commonest α -thalassemia allele in the pregnant Kuwaiti population. Additionally, $-\sum^{SEA}$ was another common genotype that accounted for 27.05% of all α -thalassemia mutations in this study. However, $-\sum^{SEA}$ was previously reported to be the most frequent α -thalassemia genotype in China.^[5,21,22] Additionally, this study only included 1 case of TI who was

Table 2

	α -thalassemia N (%)	β-thalassemia N (%)	None-thalassemia N (%)	χ ²	Р
RBC, 10 ⁹ /L				54.15	<.001
<3.8	31 (11.83)	14 (12.07)	2310 (29.72)		
≥3.8	231 (88.17)	102 (87.93)	5462 (70.28)		
HGB, g/L				281.57	<.001
<110	55 (20.99)	98 (84.48)	654 (8.41)		
≥110	207 (79.01)	18 (15.52)	7118 (91.59)		
HCT, L/L				95.14	<.001
<0.35	65 (24.81)	95 (81.90)	1348 (17.34)		
≥0.35	197 (75.19)	21 (18.10)	6424 (82.66)		
MCV, fL				1938.71	<.001
<82	91 (34.73)	109 (93.97)	129 (1.66)		
≥82	171 (65.27)	7 (6.03)	7643 (98.34)		
MCH, pg				1688.67	<.001
<27	94 (35.88)	109 (93.97)	181 (2.33)		
≥27	168 (64.12)	7 (6.03)	7591 (97.67)		
MCHC, g/L				522.73	<.001
<316	50 (19.08)	55 (47.41)	193 (2.48)		
≥316	212 (80.92)	61 (52.59)	7579 (97.52)		
MCV $<$ 82 and/or MCH $<$ 27				1643.75	<.001
Yes	96 (36.64)	109 (93.97)	197 (2.53)		
No	166 (63.36)	7 (6.03)	7575 (97.47)		

HCT=hematocrit, HGB=hemoglobin, MCH=mean cell hemoglobin, MCHC=mean corpuscular hemoglobin concentration, MCV=mean corpuscular volume, RBC=red blood cell.

diagnosed with Hemoglobin H disease. In prenatal diagnosis genetic counseling, mothers with $-\frac{SEA}{\alpha^{WS}\alpha}$ genotype can be exempted from prenatal diagnosis similarly to $-\frac{SEA}{\alpha\alpha}$ genotype.^[23]

In line with our result, Cai et $al^{[21]}$ focused on the prevalence of β-thalassemia genotypes and confirmed that the 3 most common β-thalassemia genotypes were IVS-II-654, CD41-42, and CD17 among neonates in Wuhan, Hubei Province. The proportion of different B-thalassemia genotypes in Eastern, Central, and Western areas were inconsistent, which might be explained by the unbalanced distribution of the floating population in Hubei Province. Central Hubei has a large number of floating populations than Eastern and Western areas, and the migrant population was confirmed to be one of the important factors in the prevalence of thalassemia.^[24]

The sensitivity of low MCV or MCH values on the β-thalassemia screening was 93.97% in this study, which further confirmed that MCV and MCH values are reliable on the first

step screening of B-thalassemia.^[25] Phanmany et al^[26] recommended that MCV and MCH values are regarded as important indicators in the first step of both α - and β -thalassemia screening. Whereas the proportion of pregnant α -thalassemia carriers with decreased MCV or MCH in this study were only 34.73% and 35.88%, respectively. As reported previously, most α-thalassemia static carriers have normal hematological characters.^[27] In this study, pregnant α -thalassemia static carriers comprised a high proportion of 69.81%, and little α -thalassemia static carriers were detected with MCV <82 (9.09%) or MCH <27 (10.70%) (Table S1, Supplemental Digital Content, http://links. lww.com/MD2/A885). Thus, it's not appropriate to use MCV and MCH values on the preliminary screening of pregnant α -thalassemia carriers. These results could be contributed to generalizing the thalassemia screening in perinatal care.

This study showed a high rate of negative pregnancy outcomes among thalassemia carriers, compared with general pregnant women.^[28] Thame et al^[29] announced that infants of mothers

Table 3							
Pregnancy outcomes of α - and β -thalassemia carriers.							
Variables	α -thalassemia N (%)	β -thalassemia N (%)	Total N (%)				
Maternal outcomes							
Pregnancy-induced diabetes	31 (16.58)	15 (17.86)	46 (16.97)				
Preterm birth	19 (10.16)	8 (9.52)	27 (9.96)				
Pregnancy-induced hypertension	19 (10.16)	3 (3.57)	22 (8.12)				
Eclampsia or preeclampsia	7 (3.74)	1 (1.19)	8 (2.95)				
Neonatal outcomes							
1-minute Apgar scores < 10	29 (15.51)	13 (15.48)	42 (15.50)				
5-minute Apgar scores < 10	4 (2.14)	4 (4.76)	8 (2.95)				
Fetal weight < 2500	11 (5.88)	5 (5.95)	16 (5.90)				
Fetal weight > 4000	9 (4.81)	1 (1.19)	10 (3.69)				
Birth defect	4 (2.14)	4 (4.76)	8 (2.95)				
Miscarry	1 (0.53)	0	1 (0.37)				
Stillbirth	0	1 (1.19)	1 (0.37)				

with homozygous sickle cell disease might have a higher risk of poor pregnancy outcomes. It is noticeable that the proportion of pregnancy-induced diabetes was as high as 16.58% and 17.86% among pregnant α - and β -thalassemia carriers, respectively, which was largely higher than what was reported in thalassemia major patients (6.54–9.0%).^[30,31] Another study focused on Chinese populations reported that the prevalence of diabetes was 1.90% (4/211) among patients with non-transfusion-dependent thalassemia.^[32] Attention should be paid to pregnancy-induced diabetes in thalassemia patients.

It should also be noted that α -thalassemia carriers might have worse pregnancy outcomes compared with β -thalassemia, especially for pregnancy-induced hypertension and eclampsia/ preeclampsia. The frequency of pregnancy-induced hypertension among pregnant α -thalassemia was close to what was reported in China (8.60–11.95%).^[33] However, it was much lower among pregnant β -thalassemia carriers. Serum cholesterol levels, blood viscosity, or arterial blood pressure, which were positively related with pregnancy-induced hypertension, were lower in β -thalassemia carriers compared with α -thalassemia carriers.^[34,35] Moreover, this study presented significantly decreased levels of hematological indicators including MCV and MCH among β -thalassemia carriers as mentioned above. The mechanism of the associations between α - and β -thalassemia and pregnancy outcomes needs further study.

This study has several limitations. First, this study only examined the thalassemia gene mutations and focused on the prevalence, screening, and pregnancy outcomes among pregnant women, but the hereditism of thalassemia was not examined. Second, this study only detected 17 common gene mutations; other rare mutations were not examined. Third, the prevalence of thalassemia carriers among different nations in Hubei Province was not examined in this study. Fourth, due to the high rate of lost follow-up, this study only collected the pregnancy outcomes of thalassemia carriers instead of all the participants including the control group.

5. Conclusion

This study filled the gap in the prevalence of α - and β -thalassemia carriers among pregnant women in Hubei Province. MCV and MCH values were suggested to apply on the first step of β -thalassemia screening; however, it might be unpractical on the preliminary screening of pregnant α -thalassemia carriers, among whom genetic screening is recommended as a priority. Furthermore, thalassemia carriers might have a high risk of negative pregnancy outcomes, and further study focused on the impact of thalassemia on the pregnancy outcomes is promoted in the future. These findings could be useful for the preliminary screening of thalassemia and perinatal care for the pregnant thalassemia carriers.

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