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Single nucleotide polymorphisms in the promoter region of the interleukin-6 gene and the risk of recurrent pregnancy loss in Japanese women

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Running title

IL-6 gene and recurrent pregnancy loss

Title page

Single nucleotide polymorphisms in the promoter region of the interleukin-6 gene and the risk of recurrent pregnancy loss in the Japanese population

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Capsule

There was a statistically significant difference in the IL-6-634C→G genotype frequency between Japanese patients with recurrent pregnancy loss (RPL) and controls. Patients who carry the -634 G allele may have a decreased risk of RPL.

Structured Abstract

Objective: To investigate the relationships between recurrent pregnancy loss (RPL) and single nucleotide polymorphisms (-634 C→G and -174 G→C genotypes) in the promoter region of the IL-6 gene in the Japanese population.

Design: A case-control study.

Setting: Department of Obstetrics and Gynecology in a university hospital.

Patient(s): 76 cases with RPL. Controls were 93 fertile women.

Intervention(s): Frequency and distribution of the promoter region of the IL-6 gene allele.

Main Outcome Measure(s): Determination of IL-6 promoter gene polymorphisms was performed by polymerase chain reaction and gel electrophoresis

Results: There was a significant difference in the -634C→G genotype frequency (CC vs. CG/GG) between the women with RPL and the controls. The risk of RPL was lower in the carrier of the G allele than in women with the wild type (CC) , with OR = 0.46 (95% CI = 0.24-0.91). On the other hand, we did not detect any carrier of -174C in the 169 subjects.

Conclusion(s): The results suggest that women carrying the -634 G allele of the IL-6 gene may have a decreased risk of RPL in the Japanese population.

Key words: polymorphism; interleukin-6; recurrent pregnancy loss

Introduction

About 10-14% of clinically recognized pregnancies end in pregnancy loss among the Japanese population as well as in Caucasians. The etiology of recurrent pregnancy loss (RPL) remains largely unclear (1-3). The possible immunological etiologies of pregnancy failure have been intensively investigated (4).

A shift to type-2 T-helper (Th2) cytokine production with abundant interleukine (IL)-6 and IL-10 is considered essential for the maintenance of normal pregnancy. There is evidence of a diminished Th2 immune response to placental antigens in women with RLP (5). Plasma levels of IL-6, IL-8 and IL-11 have been found to be decreased in such women compared with normal pregnancies (6). Additionally, IL-6 levels in maternal serum (7), amniotic fluid (8), vaginal fluid (9), and placenta (10) have been found to increase during the process of normal labor compared to the nonlabor state.

One study demonstrated an increase in the frequencies of type-1 T-helper (Th1) cytokine IL-1 β gene promoter region variants IL-1B-511C and IL-1B-31T in RPL women with RLP (11). One of the Th2 cytokines, IL-10 promoter region variant (-1082G \rightarrow A) was not associated with RPL (12, 13). However, relationships between many diseases and IL-6 promoter gene polymorphisms, such as IL-6 -174G \rightarrow C (14, 15) and IL-6 -634C \rightarrow G (16), were recently demonstrated. It is also known that the former polymorphism is frequently found in Caucasians (14, 15) and the latter in Japanese (16). There have been no reports of other frequent polymorphisms of the IL-6 gene, and the relationship between any IL-6 gene polymorphism and RPL has not yet been investigated. In the present study, therefore, we assessed these two promoter polymorphisms and investigated whether either of these IL-6 gene polymorphisms

predisposes women to RPL.

Materials and Methods

This case-control study was performed in Sapporo, Japan, during the years 1999-2002. We studied 76 cases aged 20-42 years with a history of RPL and 93 controls aged 21-54 years who visited Hokkaido University Hospital. The characteristics of the study groups are shown in Table 1. RPL was defined as a history of two or more spontaneous consecutive abortions and stillbirths. The primary RPL group comprised 68 women with a history of two or more pregnancy losses but no live birth. The 8 secondary RPL women experienced three or more pregnancy losses after one live birth.

All women with RLP were subjected to blood analyses for syphilis, anti-nuclear antibody (ANA), anti-DNA antibody, lupus anticoagulant, anti-cardiolipin antibody, anti-cardiolipin beta 2-glycoprotein I complex antibody and activated partial thromboplastin time, d-dimer, protein S, protein C activity, and factor XII to look for patients with anti-phospholipid antibody syndrome and thrombophilia. If ANA or anti-DNA antibody was present, further serological tests, for lupus erythematosus (LE), rheumatoid factor, anti-SSA(B) antibody, anti-RNP antibody, and anti-Sm antibody were performed, and complements were measured for the diagnosis of definite autoimmune diseases. Genetic analysis for factor V Leiden mutation was not done in this study because we previously found no Japanese having factor V Leiden mutation (17, 18).

Subjects were examined by ultrasound and hysterosalpingography for detection of anatomical abnormalities of the genital tract. All couples were also

subjected to chromosome karyotypic analyses of peripheral blood. Couples with balanced type chromosomal translocation and women with RPL having uterine conformational abnormalities such as a septate uterus were excluded from this study, because these etiologies of RPL were known to have the close cause-effect relationship. Women tested positive for autoantibodies and women with antiphospholipid antibody syndrome were included in this study, because immunological abnormalities including IL-6 possibly underlain the pathophysiology of RPL in these women. Of the 76 women, 10 (13.2%) had autoimmune disease and/or antiphospholipid antibody syndrome and 8 (10.5%) had hematologic abnormalities, but none had a history of thromboembolism.

Control women consisted of 93 volunteers experiencing at least one live birth and no abortion who had no history of endometriosis or infertility. There were no significant differences in age between cases and controls. This study was conducted with all the subjects' informed consent and approved by the institutional ethical boards for human gene and genome studies at Hokkaido University Graduate School of Medicine.

Genomic DNA was extracted from lymphocytes of peripheral blood samples by the use of standard techniques. Sequence amplification was performed by using polymerase chain reaction (PCR). To analyze the -634 C→G genotype, PCR amplifications were carried out as described by Ota et al (16), using the primers 5'-GAG AGG CCT TGA AGT AAC TG-3' and 5'-AAC CAA AGA TGT TCT GAA CTG A-3'. After the PCR product was digested with BsrBI endonuclease, the restriction digest was separated in 3% agarose gel, generating a 120-bp fragment and a 60-bp fragment. The 120-bp and 60-bp fragments represented the "G" allele. The 180-bp fragment represented the "C" allele.

To analyze the -174G→C genotype, PCR amplifications were carried out as described by Fernandez-Real et al. (14), using the primers 5'-TGA CTT CAG CTT TAC TCT TTG T -3' and 5'-CTG ATT GGA AAC CTT ATT AAG-3'. After the PCR product was digested with Hsp92II endonuclease, the restriction digest was separated in 3% agarose gel, generating a 140-bp fragment and a 58-bp fragment. The 140-bp and 58-bp fragments represented the "C" allele. The 198-bp fragment represented the "G" allele.

Statistical analysis

We calculated age-adjusted odds ratios (OR) and 95% confidence intervals (CI) associated with the IL-6 genotypes by unconditional logistic regression analysis. All analyses were conducted using SPSS software for Windows (SPSS Inc., Chicago, U.S.A.).

Results

The characteristics of the study groups are shown in Table 1. The frequencies of the -634 C→G and -174 G→C genotypes in 76 cases with RPL were compared with those in 93 controls among a Japanese population (Table 2). There was a significant difference in the -634C→G genotype frequency (CC vs. CG/GG) between the women with RPL and the controls. The risk of RPL was lower in the carriers of the G allele than in women with wild type (CC), with an OR =0.46 (95% CI = 0.24-0.91). On the other hand, we detected no -174C/C or -174G/C allele in women with RPL or in the control women and all 169 subjects were found to have the -174G/G homozygote.

We next evaluated the -634C→G genotype in both subgroups of women with

three or more pregnancy losses (PLs) and women with two PLs (Table 3). There was a significant difference in the -634C→G genotype frequency (CC vs. CG/GG) between the women with three or more PLs and the controls. The risk of RPL was very much lower in the carriers of the G allele than in women with the wild type, with OR = 0.22 (95% CI = 0.09-0.57). However, no difference in the -634C→G genotype frequency (CC vs. CG/GG) was found between the women with two PLs and the controls.

When 10 women with autoimmune disease and/or antiphospholipid antibody syndrome were excluded in the risk evaluation, we also found a significant lower risk of RPL in the carriers of the G allele than in women with wild type, with an OR = 0.45 (95% CI = 0.22-0.93). The RPL risk in women with three or more PLs was also much lower in the carriers of the G allele than in women with the wild type, with OR = 0.25 (95% CI = 0.10-0.64).

Discussion

Many investigators have assessed possible associations between etiologies of RPL and gene polymorphisms including a family of enzymes responsible for metabolism of environmental toxins, glutathione S-transferase (GST) (19-22), and associations between etiologies of RPL and the GSTP1 (20) and GSTM1 polymorphisms (22) have been demonstrated. Others sought etiologies in gene polymorphisms susceptible to infection in women with RPL; these cytokine genes included tumor necrosis factor (TNF)- α (13, 22, 24), interferon (IFN)- γ (13), and IL-1B (23, 24), and anti-inflammatory cytokine IL-10 (12, 13). However, among these studies only one demonstrated an increase in the frequencies of IL-1B promoter region variants IL1B-511C and IL-1B-31T in women with RPL; a IL1B-511C variant was

found to be associated with Th1 overimmunity to trophoblast antigens (11).

In the present study, we for the first time demonstrated an association between RPL and the -634C→G genotype of the IL-6 gene, and found that RPL risk in the carriers of the G allele was lower than that in women with the wild type, with OR = 0.46. Ota et al. (16) reported frequencies of the IL-6-634C→G allele in 470 Japanese women as follows; gene frequencies of the C allele and G allele were 0.814 and 0.186, respectively. We obtained similar allelic frequencies from control women in our study, and these were as follows; gene frequencies of the C allele and G allele were 0.801 and 0.199, respectively. Thus, allelic frequencies of IL-6-634C→G among control women in the present study did not differ from the abovementioned allelic frequencies in a Japanese population. Therefore, population bias is not likely to affect the results in the present study.

Additionally, we found an extremely low RPL risk in the carriers of the G allele among women with three or more PLs with OR = 0.22 and no significant relation of the IL-6-634C→G genotype among women with two PLs. Women with three or more PLs have a more severe condition of RPL and poorer reproductive outcome in subsequent pregnancy than do women with two PLs. Thus, the causal association of the deviation of the IL-6-634C→G genotype with RPL was further suggested.

IL-6 is a Th2 cytokine, and abnormally decreased expression of IL-6 mRNA in the uterine endometrium in women with RPL was recently found (25). Additionally, low serum concentrations of IL-6 during early pregnancy in women with RPL were found to be associated with unsuccessful pregnancy outcome (26). The women with RPL had significantly decreased plasma levels of IL-6, IL-8 and IL-11 compared to those with normal pregnancies (6).

There have as yet been no reports concerning the relationship between the IL-6 -634 allele and IL-6 production in circulation. Ota et al. (16) speculated that a decrease in bone mineral density found in carriers of the G allele, but not in subjects with the wild type, was due to effects of transcriptional activation caused by the presence of the G allele, since IL-6 is known to stimulate the osteoclast development. Whether the IL-6-634G variant results in a lower risk of abortion through a shift toward Th2-dominance during early pregnancy and increased IL-6 production in maternal circulation or in the materno-fetal interface should be further studied.

In future studies, detailed mapping and molecular characterization of the IL-6 promoter are needed to demonstrate that the -634 polymorphism has direct biochemical significance. The -634 polymorphism might simply be a molecular marker for a larger chromosomal region that includes both IL-6 and a nearby gene that affects pregnancy survival; this is another hypothetical explanation for the low RPL risk in the carriers of the IL-6-634G allele found in the present study.

It was found that the serum IL-6 concentration of the carriers of the -174C/C allele was lower than that of the carriers of the -174G/G allele in Caucasian subjects (15), and that the frequency of the -174C allele was 0.40-0.45 in Caucasians (14, 15). Although we carried out analyses for the IL-6-174G→C genotype, we detected no -174C/C or -174G/C allele in women with RPL or in the control women. Recently, no -174C allele was found among 388 Japanese men (27), and only one person with the -174C allele was found in 259 Chinese men (28). Another report found that -174G/C polymorphism was rare in Koreans (29). Thus, our results concerning the IL-6-174G→C genotype in Japanese were compatible with these previous results. Carriers of the -174C allele seem to be extremely rare among East Asians.

Numerous investigations have been performed focusing on the possible roles of immunological abnormalities in RPL. Murine studies have provided evidence that Th1 cytokines are harmful to pregnancy, causing fetal death, whereas Th2 cytokines produced at the materno-fetal interface are beneficial to the maintenance of pregnancy by suppressing cellular cytotoxicity (30, 31). Abnormal Th1/Th2 balance with Th1 dominance of peripheral mononuclear cells in response to trophoblast antigens has been found to be associated with the etiology of RPL (32).

As regards another immunological abnormality found in women with RPL, high peripheral blood natural-killer (NK) cell activities before conception (33) and early in gestation (34) have been associated with subsequent miscarriage. Additionally, decreased serum levels of macrophage migration inhibition factor during early gestation were recently found in women with RPL with subsequent miscarriage of a normal fetal karyotype (35). This evidence suggested that immunological abnormalities were causally associated with RPL.

Based on the results of the present and previous studies (11, 20, 22), we suggest that not only one but many genetic factors might be associated with RPL as predisposing factors. Whether genetic factors predisposing to RPL are reflected by the abovementioned immunological abnormalities should be further investigated. If the relation between the genetic factors and the immunological abnormalities is elucidated, genetic polymorphisms such as IL-6 -174G→C could be a marker for selecting therapeutic options and for counseling of women with RPL.

Some couples may be predisposed to produce embryos with high rates of somatic mosaicism, leading to abnormal development and RPL (38, 39). However, there are no investigations of the relationships between cytokine gene polymorphisms

and somatic mosaicism of zygotes, and this point also should be further clarified. In conclusion, it is suggested that women with the -634 G allele of the IL-6 gene have a decreased risk of RPL.

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TABLE 1. Characteristics of 76 cases with recurrent pregnancy loss (RPL) and 93 controls in a Japanese population

		Cases		Controls	
		Number	%	Number	%
Age					
	20-29	28	36.8	36	38.7
	30-39	41	53.9	48	51.6
	≥ 40	7	9.2	9	9.7
Number of previous pregnancy losses					
	2	29	38.2	—	—
	3	32	42.1	—	—
	≥ 4	15	19.7	—	—

TABLE 2. Distribution of Il-6 genotypes among 76 cases with RPL and 93 controls

	Cases		Controls		OR* (95% CI)	P value
	Number	%	Number	%		
IL-6						
-634 C→G						
CC	58	76.3	56	60.2	1.00	0.026
CG/GG	18	23.7	37	39.8	0.46 (0.24-0.91)	
-174 G→C						
GG	76	100	93	100	-	-
CG/CC	0	0	0	0	-	

*Age-adjusted logistic regression analysis.

TABLE 3. Distribution of IL-6 -634 genotypes among the cases with three or more pregnancy losses (PLs) and with two PLs and controls

	Cases		Controls		OR* (95% CI)	P value
	Number	%	Number	%		
Three or more PLs						
IL-6-634 C→G						
CC	41	87.2	56	60.2	1.00	
CG/GG	6	12.8	37	39.8	0.22 (0.09-0.57)	0.002
Two PLs						
IL-6-634 C→G						
CC	17	58.6	56	60.2	1.00	
CG/GG	12	41.4	37	39.8	1.05(0.45-2.46)	NS

*Age-adjusted logistic regression analysis.