

Received: 2015.12.03
Accepted: 2015.12.22
Published: 2016.07.08

Evaluation of ICAM-1 and VCAM-1 Gene Polymorphisms in Patients with Periodontal Disease

Authors' Contribution:
Study Design A
Data Collection B
Statistical Analysis C
Data Interpretation D
Manuscript Preparation E
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Source of support: This work was self-funded by the Department of Periodontology, Clinical School of Stomatology, First Affiliated Hospital of Harbin Medical University, Harbin, Heilongjiang, P.R. China

Background: We aimed to investigate the potential genetic relationships between the polymorphisms of gene rs5498 *ICAM-1* and rs1041163 *VCAM-1* and chronic periodontitis in a Chinese population within Heilongjiang.

Material/Methods: Genomic DNA was extracted from oral mucosa cells of 584 periodontal patients and 182 healthy individuals. Genotyping of the rs5498 *ICAM-1* and rs1041163 *VCAM-1* gene polymorphisms was performed with the Multiplex SNaPshot technique.

Results: Statistically significant associations were identified between the chronic periodontal patients and the controls in the gene polymorphisms of rs5498 *ICAM-1* ($P=0.007$) and rs1041163 *VCAM-1* ($P=0.029$). The distribution of rs5498 ($P=0.029$) and rs1041163 ($P=0.049$) differed significantly across the mild, moderate, and severe groups of periodontitis.

Conclusions: Our findings indicate that *ICAM-1* rs5498 and *VCAM-1* rs1041163 polymorphisms contribute to chronic periodontitis, and *ICAM-1* rs5498 and *VCAM-1* rs1041163 gene polymorphisms might be associated with periodontitis severity in the Heilongjiang Chinese population. Further studies should be conducted to determine whether these polymorphisms could be used as biomarkers of periodontitis.

MeSH Keywords: **Chronic Periodontitis • Inflammation • Polymorphism, Genetic**

Full-text PDF: <http://www.medscimonit.com/abstract/index/idArt/896979>



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Background

Periodontal disease is characterized by Gram-negative bacteria in the oral cavity, along with chronic inflammation [1,2], which could consequently lead to the destruction of the soft tissue and resorption of the periodontal bone. It was suggested that the alteration or progression of the disease may be associated with the host immune response to these bacteria, and that inflammatory cells in diseased periodontal tissues are related to the disease progression [3,4]. Recruitment and retention of these cells are determined by cell adhesion molecules, which are suggested to be involved in the pathogenesis of inflammatory diseases [5]. Evidence shows that cell adhesion molecules can be up-regulated as a result of stimulation from certain pro-inflammatory cytokines, and can act as co-stimulatory receptors in the activation of inflammatory cells [6].

Cellular adhesion molecule (CAM), an adhesion protein on the surface of the gums, epithelium of gingival sulcus, and junctional epithelium, is an important component of the adhesion between cells and their extracellular matrix [7]. CAM is involved in the occurrence and development of periodontitis, mainly through the participation of cell information transmission, inflammation, and immune response [8,9]. With the assistance of the specificity of CAM on the junctional epithelium, polymorphonuclear leukocyte (PMN) can enter the gingival and periodontal pocket, playing the role of sterilization. The level of CAM is positively correlated with the degree of periodontal inflammation [10].

Intercellular adhesion molecule 1 (ICAM-1) and vascular cell adhesion molecule 1 (VCAM-1) belong to the Ig superfamily [11,12]. ICAM-1 is widely expressed on leucocytes, endothelial cells, monocytes, synovial cells, fibroblasts, and epithelial cells [13,14]. In immunological and inflammatory reactions, ICAM-1 and VCAM-1 have the ability to promote cell adhesion and recruit leucocytes. These abilities of ICAM-1 and VCAM-1 were found to be involved in rheumatoid arthritis [15]. In confirmation of this information, a study also showed that the plasma and synovial fluid levels of ICAM-1 and VCAM-1 were significantly higher in rheumatoid arthritis patients in comparison with normal healthy controls [16]. The *ICAM1* gene is located on locus 19p13.3-p13.2 [17]. As an indispensable participant in the initiation of the immunological response, ICAM-1 mediates adhesion of leukocytes against the blood vessel wall, enabling them to enter the tissues by transendothelial migration [18]. The *VCAM1* gene is located on locus 1p31-32 [19]. As an endothelial receptor for VLA-4 and integrin $\alpha 4\beta 7$, *VCAM1* contributes to the initiation of the T cell response to alloantigens [20]. The polymorphisms of *ICAM1* and *VCAM1* genes may affect the function of the immune response [21]. Several studies have indicated that polymorphisms of rs5498 T>C in exon 6 of the *ICAM1* gene and rs1041163 T>C in the *VCAM1*

gene promoter are associated with various diseases, such as coronary artery disease, stenosis, myocardial infarction, and atherosclerosis [22–24]. Because the pathophysiology of periodontitis is similar to that of these diseases, which is inflammatory in nature, and no case-control study has been conducted on the associations between *ICAM-1* and *VCAM-1* gene polymorphisms and chronic periodontitis, we evaluated their roles in the susceptibility to periodontal diseases and investigated the potential associations between the polymorphisms and chronic periodontitis in the Heilongjiang Chinese population.

Scientific rationales

Previous studies indicated that *ICAM-1* rs5498 and *VCAM-1* rs1041163 gene polymorphisms are associated with various diseases, including coronary artery disease, stenosis, and atherosclerosis. Since the pathophysiology of these diseases is inflammation, similar to that of periodontitis, and because ICAM-1 and VCAM-1 are known to be involved in the occurrence and development of periodontitis, we hypothesized that the polymorphisms were also associated with periodontitis.

Main findings

ICAM-1 rs5498 and VCAM-1 rs1041163 polymorphisms contribute to chronic periodontitis, and ICAM-1 rs5498 and VCAM-1 rs1041163 gene polymorphisms might be associated with periodontitis severity in the Heilongjiang Chinese population.

Practical implications

Our study is the first to investigate the relationships between ICAM-1 and VCAM-1 polymorphisms and periodontitis.

Material and Methods

Ethics statement

This study was approved by the Ethics Committee of The First Affiliated Hospital of Harbin Medical University and each subject signed informed consent prior to enrollment in the current study. All procedures in this study were in accordance with the Declaration of Helsinki [25].

Subjects

In this study, 766 nonsmoking subjects, who were admitted to the First Affiliated Hospital of Harbin Medical University between September 2012 and December 2014, were recruited. There were 584 patients with periodontitis (284 females and 300 males; age range: 20–50 years) among whom 228 had mild, 212 had moderate, and 144 had severe chronic periodontitis.

Table 1. Genotype and allele frequency distribution of ICAM-1 rs5498.

	Controls (n=182)		Cases (n=584)		χ^2	P	OR (95%CI)
	N (frequency)		N (frequency)				
T/T	94	(51.6%)	252	(43.2%)	9.893	0.007	1.000
T/C	60	(33.0%)	268	(45.9%)			0.600 (0.416–0.866)
C/C	28	(15.4%)	64	(11.0%)			1.952 (1.156–3.303)
Allele							
T	248	(68.1%)	772	(66.1%)	0.517	0.472	1.031 (0.950–1.118)
C	116	(31.9%)	396	(33.9%)			0.940 (0.793–1.114)

ICAM-1 – intercellular adhesion molecule 1; OR – odds ratios; 95%CI – 95% confidence intervals.

Table 2. Genotype and allele frequency distribution of VCAM-1 rs1041163.

	Controls (n=182)		Cases (n=584)		χ^2	P	OR (95%CI)
	N (frequency)		N (frequency)				
C/C	7	(3.8%)	8	(1.4%)	7.069	0.029	1.000
C/T	40	(22.0%)	100	(17.1%)			1.143 (0.156–1.344)
T/T	135	(74.2%)	476	(81.5%)			0.709 (0.468–1.072)
Allele							
C	54	(14.8%)	116	(9.9%)	6.764	0.009	1.494 (1.106–2.018)
T	310	(85.2%)	1052	(90.1%)			0.946 (0.902–0.991)

VCAM-1 – vascular cell adhesion molecule 1; OR – odds ratios; 95%CI – 95% confidence intervals.

Table 3. Genotype and allele frequency distribution of ICAM-1 rs5498 among mild, moderate and severe cases.

	Mild cases (n=228)		Moderate cases (n=212)		Severe cases (n=144)		χ^2	P
	N (frequency)		N (frequency)		N (frequency)			
Genotype								
C/C	16	(7.0%)	32	(15.1%)	16	(11.1%)	11.519	0.021
C/T	104	(45.6%)	88	(41.5%)	76	(52.8%)		
T/T	108	(47.4%)	92	(43.4%)	52	(36.1%)		
C/C	16	(7.1%)	32	(15.1%)	16	(11.1%)	7.099	0.029
C/T+ T/T	209	(92.9%)	180	(84.9%)	128	(88.9%)		

ICAM-1 – intercellular adhesion molecule; OR – odds ratios; 95%CI – 95% confidence intervals.

Discussion

It has been proven that periodontitis, a complex multifactorial disease, is determined by both environmental and genetic factors. In addition to the pathogenic bacteria and other environmental factors (e.g., smoking and stress) [26,27] involved in the pathogenesis of periodontitis, there is also evidence suggesting that genetic factors might participate in the etiology of

periodontitis [28]. Recently, growing interest has been focused on identifying allelic variants of genes that can be used to assess the risk of periodontal diseases [29]. *ICAM-1* and *VCAM-1* polymorphisms are currently being investigated and have been found to be associated with several diseases, including stenosis, coronary artery disease, and atherosclerosis [30,31]. To the best of our knowledge, however, our study is the first to investigate the association between these polymorphisms and periodontitis.

Table 4. Genotype and allele frequency distribution of VCAM-1 rs1041163 among mild, moderate and severe cases.

Genotype	Mild cases (n=228)		Moderate cases (n=212)		Severe cases (n=144)		χ^2	P
	N (frequency)		N (frequency)		N (frequency)			
C/C	4	(1.8%)	0	(0%)	4	(2.8%)	9.559	0.049
C/T	48	(21.1%)	32	(15.1%)	20	(13.9%)		
T/T	176	(77.2%)	180	(84.9%)	120	(83.3%)		

VCAM-1 – vascular cell adhesion molecule 1; OR – odds ratios; 95%CI – 95% confidence intervals.

There was a significant difference in the rs5498 polymorphism of the ICAM-1 gene between the chronic periodontal patients and the controls. The genotype distribution of VCAM-1 rs1041163 was also significantly different between the patients and the controls. Moreover, a significantly higher prevalence of 'T' allele was observed in the patients than in the controls. These results suggest that rs5498 and rs1041163 polymorphisms may contribute to chronic periodontitis. Among the mild, moderate, and severe cases of periodontitis, the genotypes of rs5498 were found to have different distributions; similar results were obtained for the genotypes of rs1041163, suggesting that both rs5498 and rs1041163 may affect periodontitis severity.

The exact biological function of intronic SNPs is currently unclear. However, the probable mechanism underlying the function is as follows. Both ICAM-1 and VCAM-1 mRNAs are expressed by non-stimulated hepatocyte growth factor (HGF) [32]. In addition, ICAM-1 is induced by pro-inflammatory cytokines, such as IL-1 β , TNF- α , IFN- γ , and IL-2 [33,34]. *Escherichia coli* LPS [35] and VCAM-1 are induced by IL-1 β . ICAM-1 mediates the middle screw rotary body into the process of human gingival epithelium as an adhesion receptor on leukocytes and endothelial cells [36,37]. A previous study suggested the ICAM-1 plays an important role in the early stage and progression of chronic periodontitis [38]. VCAM-1, combined with monocytes and lymphocytes, can activate vascular endothelial cells to express vascular adhesion molecules [39]. The release of cytokines and enzymes promotes the progress of inflammation. Animal studies have shown that suppressing the expression of VCAM-1 can significantly inhibit the inflammatory responses [40]. In order to control the inflammatory responses, it is critical to effectively suppress the expression of VCAM-1. These results indicate that SNPs of ICAM-1 and VCAM-1 may cause susceptibility to periodontitis, and that different distributions may result in different severities of periodontitis. It is thus possible to intervene in and control the progression of periodontitis and to treat periodontitis on the SNP level by identifying the polymorphisms of ICAM-1 and VCAM-1.

The pathophysiology of periodontitis, similar to that of some other complex diseases, is characterized by various biological pathways [41,42], which ultimately lead to the same clinical manifestation. It is important to note that the number and type of genes involved in the same disease may not be the same in different ethnic populations. Accordingly, a functional SNP may be in linkage disequilibrium with distinct markers in different ethnic groups. Therefore, we cannot deny the possibility that functional rs5498 and rs1041163, together with the other uncovered functional variants in linkage disequilibrium, might synergistically influence diseases.

Taken together, our study shows that *ICAM-1* rs5498 and *VCAM-1* rs1041163 gene regions may affect the susceptibility to chronic periodontitis in the Heilongjiang Chinese population. These 2 polymorphisms might also affect the severity of periodontitis. Our work provides the first evidence of the involvement of human *ICAM-1* and *VCAM-1* genes in chronic periodontitis. However, the basic functions of *ICAM-1* and *VCAM-1* gene mutations remain to be further elucidated.

Conclusions

Our findings indicate that *ICAM-1* rs5498 and *VCAM-1* rs1041163 polymorphisms participate in the pathogenesis of chronic periodontitis, and that the gene polymorphisms at both *ICAM-1* rs5498 and *VCAM-1* rs1041163 might be associated with periodontitis severity in the Heilongjiang Chinese population. Future studies should focus on the application of these polymorphisms in the diagnosis and treatment of periodontitis as biomarkers.

Conflict of interest

We declare that we have no conflicts of interest.

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