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Interleukin-4 promoter polymorphisms in patients with aggressive or chronic periodontitis

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Introduction

In periodontitis susceptibility individuals, disease progression involves a predominantly Th2-like immune response (1). IL-4 stimulates a Th2-type of immune response while inhibits Th1 cells (2,3). Moreover, IL-4 limits the persistence of macrophages in periodontitis lesions and down regulates CD14 receptor, one of the key receptors for lipopolysaccharides (4).

Objectives

Promoter polymorphisms of the interleukin-4 gene have been associated with altered IL-4 production (5-7). Thus, they could be indicative for periodontitis. Therefore, the aim of this study was to determine allele- genotype- and haplotype- frequencies of IL-4 SNPs at positions -1098T/G, -590C/T and -33C/T in patients with generalized aggressive (AP) or generalized chronic periodontitis (CP) in comparison to periodontitis free controls.Calculation of adjusted Odds ratios (OR) were carried out with respect to established cofactors for periodontitis such as age, gender, smoking, and plaque index.

Material and Methods

Study groups

121 patients with sever periodontitis (attachment loss >4mm in 80% of the teeth; CP: n=54, mean age 49,3 ±10.0 years; AP: n=67, mean age 41 ±9.9 years) and 81 individuals without periodontitis (Controls:mean age 46,9 ±10.7 years) were included.

Proof of IL-4 promoter polymorphisms

IL-4 SNPs were analyzed by PCR-SSP (CTS-Kit, Heidelberg, Germany). Distributions of single alleles, genotypes, and haplotypes were calculated by Chi²-Test with Yates correction or Fisher's exact test. Risk factor analyses were carried out by logistic regression under consideration of established cofactors for periodontitis such as age, gender, smoking status, and plaque index.

Results

Allele frequencies

The mutant allele G at position -1098 was significantly increased among both patient groups. The mutant alleles -590 T und -33 T were more frequently proven only among patients with AP (Fig. 1)

Genotype frequencies

The mutant genotype -1098 TG was significantly increased among both patient groups (Fig. 2). Moreover, the mutant genotypes -590 CT and -33 CT were likewise significantly increased (linkage disequilibrium!) among patients with AP (Fig. 3 and 4).

Haplotype frequencies

Among both, patients with AP and CP the mutant haplotype -1098, -590, -33 GCC was significantly increased. Moreover, the haplotype -1098, -590, -33 TTT was significantly more proven only among patients with AP (Fig. 5).

Risik analysis with binary logistical regression

The haplotype -1098,-590,-33 GCC increased the adjusted OR for AP and CP. Mutations (C>T) at positions -590 and -33 were indicative only for AP (Fig. 6).



Fig. 6



Conclusions

Genotypes and haplotypes who expressed the mutant allele G at position -1098 were indicative for both, CP and AP. Polymorphisms (C>T) at positions -590 and -33 (high interleukin-4 producer genotypes) increased the Odds ratio for an AP. In a further study it would be useful to investigate the IL-4 production in association to IL-4 polymorphisms.

Literature

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Abbreviations

AP = aggressive periodontitis CP = chronic periodontitis IL = interleukin Th = T helper cells PCR-SSP = polymerase chain reaction with sequence specific primers

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INTRODUCTION

IL-4 stimulates a Th2-type of immune response while inhibits Th1 cells. For example, IL-4 limits the persistence of macrophages in periodontitis lesions and down regulates CD14 receptor, one of the key receptors for lipopolysaccharides. Promoter poly-morphisms of the interleukin-4 gene have been associated with altered IL-4 production. Thus, they could be indicative for periodontitis.

AIMS

Determination of allele- genotype- and haplotype-frequencies of IL-4 SNPs at positions -1098T/G, -590C/T and -33C/T in patients with generalized aggressive (AP) or generalized chronic periodon

titis (CP) Calculation of adjusted Odds ratios (OR) with respect to the cofactors age, gender, smoking, and plaque index.

MATERIAL UND METHODS

Study groups

121 patients with sever periodontitis (attachment loss >4mm in 80% of the teeth; CP: n=54, age 49,3 years; AP: n=67, age 41 years) and 81 individuals without periodontitis (Controls: age 46,9 years) were included.

Proof of IL-4 promoter polymorphisms IL-4 SNPs were analyzed by PCR-SSP (CTS-Kit, Heidelberg, Germany). Distributions of single alleles, genotypes, and haplotypes were calculated by Chi²-Test with Yates correction or Fisher's exact test. Risk factor analyses were carried out by logistic regression under consideration of established cofactors for periodontitis such as age, gender, smoking status, and plaque index.

RESULTS

Allele frequencies

The mutant allele G at position -1098 was significantly increased among both patient groups. The mutant alleles -590 T und -33 T were more frequently proven only among patients with AP (Fig. 1)





CONCLUSIONS

Genotypes and haplotypes who expressed the mutant allele G at position -1098 were indicative for both, CP and AP. Polymorphisms at positions -590 and -33 (high interleukin-4 producer genotypes) increased the Odds ratio for an AP. In a further study it would be useful to investigate the IL-4 serum levels in association to IL-4 polymorphisms.

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Genotype frequencies The mutant genotype -1098 TG was significantly increased among both patient groups (Fig. 2). Moreover, the Abl. 2 Significant genotype trea at position -1098 T+G mutant genotypes -590 CT and -33 CT were likewise significantly increased (linkage disequilibrium!) among patients with AP (Fig. 3 and 4). OTT VS. TO



Haplotype frequencies

Among both, patients with AP and CP the mutant haplotype -1098, -590, -33 GCC was significantly increased. -1098, -590, -33 TTT was significantly more proven only among patients with AP (Fig. 5)

Fig. 5 Significant IL-4 haplotypes at p -1098,-590,-33



Risik analysis with binary logistical regression The haplotype -1098,-590,-33 GCC increased the adjusted OR for AP and CP. Mutations at positions -590 and -33 were indicative only for AP (Fig. 6).

