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ORAL DISEASES

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TGF-β1 gene polymorphism in renal transplant patients with and without gingival overgrowth

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Short title: TGF-β1 gene polymorphism in gingival overgrowth

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BACKGROUND: The incidence of gingival overgrowth among renal transplant patients treated with cyclosporine A ranges from 13 to 84.6%, and the overgrowth is not only esthetic but also medical problem. The study was aimed at determination of association between TGF- β 1 gene (*TGFB1*) polymorphism and gingival overgrowth in kidney transplant patients medicated with cyclosporin A.

METHODS: 84 kidney transplant patients with gingival overgrowth and 140 control transplant patients without overgrowth were enrolled into the case control study. *TGFB1* polymorphism was determined using the PCR-RFLP assay for +869T>C in codon 10 and +915G>C in codon 25 as well as TaqMan real-time PCR assays for promoter -800G>A and -509C>T SNPs.

RESULTS: In kidney transplant patients suffering from gingival overgrowth mean score of gingival overgrowth was 1.38 ± 0.60 , whereas in control subjects was 0.0. The patients with gingival overgrowth were characterized by similar distribution of *TGFB1* genotypes and allele in comparison to subjects without gingival overgrowth. Among 16 potentially possible haplotypes of *TGFB1* gene, only four were observed in the studied sample of kidney transplant patients: G_C_T_G, G_T_C_G, G_C_C_C, and A_C_T_G, with similar frequency in patients with and without gingival overgrowth.

CONCLUSION: No association between the *TGFB1* gene polymorphism and gingival overgrowth was revealed in kidney transplant patients administered cyclosporine A.

Key words: TGF- β 1; gingival overgrowth; polymorphism

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Introduction

Gingival overgrowth frequently occurs in transplant patients receiving immunosuppressive drugs such as cyclosporine (INN ciclosporin), which has been widely used since 1970s. The incidence of gingival overgrowth among renal transplant patients treated with cyclosporine A ranges from 13 to 84.6% (Somacarrera *et al*, 1994; Margiotta *et al*, 1996; Pernu *et al*, 2001; Afonso *et al*, 2003). Gingival overgrowth induced by cyclosporine A is probably the consequence of alterations in the turnover of epithelial and connective tissues, resulting in an increase in structural proteins sometimes associated with epithelial thickening. Numerous studies have investigated potential risk factors in the development of the condition, including the extent of renal disease, interval since transplantation, duration of renal replacement therapy, dose of cyclosporine A, gingival inflammation, plaque indices and recipient HLA-phenotype. Whilst some studies have suggested some associations between the incidence and severity of gingival overgrowth and sex, pre-transplant diagnosis, age at transplantation, duration of therapy and cyclosporine dosage, others have failed to find any significant drug-related risk factors for the development of the disease as previously reported (Seymour *et al*. 1996; Thomas *et al*. 2000; Thomas *et al*. 2001; Afonso *et al*. 2003). However, it remains unclear why a proportion of patients are susceptible to gingival overgrowth while others remain unaffected.

It was hypothesized, that gingival overgrowth induced by cyclosporine results in a disturbance in the homeostatic balance which is characterized by an increase in both the number of fibroblasts and an increase in the volume of the extracellular matrix. This loss of growth control results in an accumulation of redundant tissue of relatively normal composition (McGaw *et al*, 1988). There is considerable information supporting the role for transforming growth factor- β 1 (TGF- β 1) in mediation the dysregulated fibroblast proliferation and extracellular matrix synthesis. Evidence from experimental studies demonstrate that gingival fibroblasts in culture exhibit a proliferative response to TGF- β 1 (Anderson *et al*, 1998; James *et al*, 1998). It was also reported that TGF- β 1 suppresses extracellular matrix degradation via downregulation of metalloproteinases (MMP) production and inducing tissue inhibitor of MMP synthesis (Edwards *et al*, 1987; Overall *et al*, 1989). Also, administration of TGF- β 1 antisense oligonucleotide increase the expression of collagen degrading enzymes in human gingival fibroblasts (Cotrim *et al*, 2002). Experimental data also revealed that proliferation of rat gingival cells under cyclosporine A depended on TGF- β 1 mRNA induction by the drug (Yoshida *et al*, 2005). Clinical studies demonstrated that circulating and gingival crevicular fluid levels of TGF- β 1 were increased in cyclosporin-medicated transplant patients (Buduneli *et al*, 2001), and TGF- β 1 was found to be an independent risk factor for gingival overgrowth in immunosuppressed patients, also medicated with cyclosporine A (Ellis *et al*, 2004;

Gurkan *et al*, 2008). Recently, an accumulation of extracellular matrix related to TGF- β 1 overexpression in cultured fibroblasts from patients medicated with cyclosporine A has been reported by Dreyfuss *et al* (2010).

It was reported that expression of TGF- β 1 gene (*TGFB1*) may depend on genetic constitution, i.e. within the *TGFB1* some functional SNPs were defined. Clinical studies in kidney transplant patients medicated with cyclosporine A suggested an association between *TGFB1* gene polymorphism located in coding regions (codon 10 - +869T>C Pro10Leu and codon 25 +915G>C Arg25Pro) and gingival overgrowth (Linden *et al*, 2001; Radwan-Oczko *et al*, 2006). However, contradictory results in kidney transplant patients with respect of the same as aforementioned SNPs are also available (Radwan-Oczko *et al*, 2008).

Therefore, it was decided to investigate the role of four SNPs within *TGFB1* gene, previously associated with alterations in gene expression level, in gingival overgrowth in kidney transplant patients medicated with cyclosporine A. Two of the SNPs were previously evaluated, and lead to aminoacid substitution: +869T>C in codon 10 (rs1800470, Pro10Leu) and +915G>C in codon 25 (rs1800471, Arg25Pro). The other two SNPs are located in promoter region of *TGFB1*: -800G>A (rs1800468), and -509C>T (rs1800469), and thus can affect promoter function and gene expression (Grainger *et al*., 1999; Grainger *et al*., 2000).

Materials and Methods

The protocol of the study was approved by the ethics committee of the Pomeranian Medical University, Szczecin, Poland, i.e. the experiments were undertaken with the understanding and written consent of each subject and according to ethical principles, including the World Medical Association Declaration of Helsinki (version, 2002). Patients of Polish origin, Caucasians, hospitalized and than followed-up at the Department of Nephrology, County Hospital, Szczecin, Poland were recruited into the study from 2002 to 2010 year, after giving informed consent. A total of 84 unrelated kidney transplant patients suffering from gingival overgrowth (61 males, 23 females) aged from 16 to 70 years (mean 42.2 ± 13.1 years) were enrolled into the study. All patients were examined by two independent consultant periodontal specialists 6 months after kidney transplantation. Periodontists were blinded to SNP status. The patients were assessed using a clinical scoring method according to Pernu *et al* (1992). The patients were ascribed a general whole-mouth score of between 0 and 3: 0=no overgrowth seen; 1=mild gingival overgrowth (thickening of the marginal gingiva and/or lobular granulation of the gingival pocket as well as overgrowth covering the gingival third of the crown or less); 2=moderate gingival overgrowth (overgrowth

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2 extending to the middle of the crown); 3=severe gingival overgrowth (overgrowth covering two
3 thirds of the crown or affection of the whole attached gingiva).

5 Control samples were from 140 kidney transplant patients (91 males, 49 females), aged from 17 to
6 66 years (mean 39.7±11.4 years), who were free from gingival overgrowth signs at 6 months after
7 transplantation, as evaluated by consultant periodontal specialists. Patients' characteristics are
8 shown in table 1.

12 During the study period all subjects were administered cyclosporin A, azathioprine, prednisone and
13 one of two calcium channel blockers, i.e. diltiazem or verapamil; eight patients without gingival
14 overgrowth were given atenolol or prazosine. The medication regimens administered in both groups
15 of the study, including cyclosporin A dose and concentrations, dosages of verapamil, diltiazem and
16 prednisone evaluated at monthly intervals during 6 months were comparable, i.e. all calculated p-
17 values were >0.05 (Table 2). The serum concentrations of cyclosporin A were measured by
18 fluorescence polarization immunoassay using TDx analyzer (Abbott) in all patients.

27 *Genotyping*

29 Four SNPs within *TGFBI* gene, previously associated with alterations in gene expression level were
30 selected for the purpose of this study. Two of them are located in promoter region of *TGFBI*:
31 -800G>A (rs1800468), and -509C>T (rs1800469), the others lead to aminoacid substitution:
32 +869T>C (rs1800470, Pro10Leu) and +915G>C (rs1800471, Arg25Pro). Genomic DNA was
33 extracted from 200 µL of whole blood samples using GeneMATRIX Quick Blood DNA
34 Purification Kit (EURx, Poland). Each individual was genotyped for a presence of 4 SNPs in
35 *TGFBI* gene. The allelic discrimination TaqMan real-time PCR assays (Assay IDs: C_8708474_20,
36 C_8708473_10, and C_22272997_10, Applied Biosystems, USA) were used for detection of
37 -800G>A, -509C>T and +869T>C SNPs, respectively. Fluorescence data was captured using an
38 ABI PRISM 7500 FAST Real-Time PCR System (AppliedBiosystems), after 40 cycles of PCR. For
39 analysis of +915G>C polymorphism, a PCR-RFLP method was applied. Briefly, polymerase chain
40 reaction was performed in 15µl of total volume, with a pair of primers: 5'-cgc tgc tgt ggc tac tgg t-3',
41 and 5'-ctc cgg ttc tgc act ctc c-3', previously described by Pulleyn et al. [2001]. Subsequently, a 254
42 bp PCR product was digested with the *Cfr13I* endonuclease (Fermentas, Lithuania), yielding two
43 DNA fragments (171+83 bp) in case of wild-type G allele. The +915G>C substitution creates
44 additional restriction site recognized by *Cfr13I*, yielding 3 fragments (141+83+30 bp), which was
45 visualized after electrophoresis in 3.5% agarose gels stained with ethidium bromide (Pulleyn *et al*,
46 2001).

Statistical analysis

The data were tested for Hardy–Weinberg equilibrium by calculating expected frequencies of genotypes and comparing them to the observed values using the χ^2 test (Statistica 8.0, Statsoft Software, Warsaw, Poland). Associations between categorical variables were assessed by the Fisher exact test. The EH program (Jurg Ott, Rockefeller University, New York) was used to estimate haplotype frequencies. Linkage disequilibrium (LD) was measured as followed - the D' was calculated using 2LD software and squared correlation coefficient (r^2) was evaluated. Odds ratios (OR) and 95% confidence interval (95% CI) were calculated using the Newcombe-Wilson method without the continuity correction. A p level of less than 0.05 was considered statistically significant.

Results

Out of 84 patients with gingival overgrowth, 57 subjects were classified as score 1 of gingival overgrowth, 22 patients were ascribed score 2 and 5 subjects score 3. Mean score of gingival overgrowth was 1.38 ± 0.60 according to Pernu's scoring system (Pernu *et al*, 1992). Control transplant patients were characterized by healthy gingiva, i.e. were scored 0.

The genotype frequency distribution for all analyzed SNPs did not show a significant deviation from Hardy-Weinberg equilibrium in any of the study groups ($p > 0.1$). The distribution of *TGFB1* gene genotypes and alleles in kidney transplant patients is shown in Table 3. The patients with gingival overgrowth induced by immunosuppressive medication were characterized by similar distribution of *TGFB1* genotypes and allele to subjects without overgrowth, i.e. SNPs within the two coding regions: +869T>C in codon 10 (rs1800470, Pro10Leu) and +915G>C in codon 25 (rs1800471, Arg25Pro) as well as two promoter region SNPs: -800G>A (rs1800468), and -509C>T (rs1800469). In the next step of the study we subjected for analysis haplotypes of *TGFB1* gene (Table 4). The results of linkage analysis showed that all analysed SNPs are in strong linkage in a Polish population (Figure 1). The EH program (Jurg Ott, Rockefeller University, New York) was used to estimate haplotype frequencies based on results of typing for the following studied SNPs: -800G>A, -509T>C, +869T>C, +915G>C. Among 16 potentially possible haplotypes derived from 4 SNPs, only four were observed in the studied sample of kidney transplant patients, with respective frequencies: G_C_T_G (0.545), G_T_C_G (0.324), G_C_C_C (0.071), and A_C_T_G (0.060). Similarly to distribution of the genotypes and allele, frequency of *TGFB1* haplotypes were comparable in kidney transplant patients with and without gingival overgrowth.

Discussion

TGF- β 1 pathophysiological data as well as reports on functional polymorphism of its gene – *TGFB1*

1 suggest potential role of the cytokine and polymorphism in the pathology of drug-induced including
2 cyclosporin A, gingival overgrowth (Edwards *et al*, 1987; Anderson *et al*, 1998; James *et al*, 1998;
3 Overall *et al*, 1989; Grainger *et al*, 1999; Grainger *et al*, 2000 Buduneli *et al*, 2001; Cotrim *et al*,
4 2002; Ellis *et al*, 2004; Yoshida *et al*, 2005; Gurkan A *et al*, 2008). However, available data on the
5 effects of *TGFB1* gene polymorphism on gingival overgrowth in kidney transplant patients
6 medicated with cyclosporin A are controversial. Linden *et al* (2001) and Radwan-Oczko *et al*
7 (2006) revealed that polymorphisms in *TGFB1* gene located in coding regions of the gene, i.e.
8 codon 10 - +869T>C Pro10Leu and codon 25 +915G>C Arg25Pro are genetic factors associated
9 with risk of gingival overgrowth. Subsequent report from the latter group (Radwan-Oczko *et al*,
10 2008) showed that the aforementioned polymorphisms are not associated with gingival overgrowth
11 observed in kidney transplant patients medicated with cyclosporin A.

12 Therefore, it was decided to evaluate *TGFB1* gene polymorphism in kidney transplant patients
13 medicated with cyclosporine A. The former studies included polymorphisms in the coding regions
14 of the gene resulting in aminoacid substitutions, namely in codon 10 - +869T>C Pro10Leu and
15 codon 25 +915G>C Arg25Pro (Linden *et al*, 2001; Radwan-Oczko *et al*, 2006; Radwan-Oczko *et al*
16 2008). However, it is well known that polymorphisms located in promoter region of the gene can
17 also be functional, affecting expression level of genes, hence in the present study polymorphisms in
18 the promoter of *TGFB1* gene were evaluated at positions -800G>A and -509C>T (Grainger *et al*,
19 1999; Grainger *et al*, 2000). The promoter polymorphisms, although functional, have not been
20 studied in posttransplant patients with gingival overgrowth yet. Furthermore, we evaluated
21 haplotypes of *TGFB1* gene as potential risk factors associated with gingival overgrowth in patients
22 administered cyclosporine A.

23 Previous reports, similarly to our report included patients from Caucasian populations, but number
24 of cases analysed was smaller than in the present study, study by Linden *et al* (2001) involved 164
25 kidney transplant patients medicated with cyclosporin A (of them 46 with gingival overgrowth), by
26 Radwan-Oczko *et al* from 2006 comprised 92 cases (of them 50 with gingival overgrowth) and
27 Radwan-Oczko *et al* from 2008 included 98 subjects (of them 54 with gingival overgrowth). The
28 present study recruited 224 patients, including 84 with gingival overgrowth. Having in mind power
29 of the study the present report seems to be most reliable in comparison to the further studies. The
30 results of the present study are in line with the report of Radwan-Oczko *et al* (2008) demonstrating
31 no association of the coding region polymorphisms of *TGFB1* gene and gingival overgrowth,
32 contrary to studies of Linden *et al*, 2001 and Radwan-Oczko *et al* 2006. The present study was
33 extended to functional polymorphisms located in promoter region of the *TGFB1* gene, and similarly
34 to results from coding regions any associations of genetic factors with gingival overgrowth in

kidney transplant patients were revealed. Finally, haplotype frequencies based on results of typing for the studied SNPs: -800G>A, -509T>C, +869T>C, +915G>C were analysed. Similarly to genotypes, the analysed haplotypes did not discriminate the studied groups, i.e. kidney transplant patients medicated with cyclosporin A with and without gingival overgrowth.

Interaction between simultaneously administered drugs affecting enlargement have been also reported. Cyclosporin A-treated patients are often comedicated with azathioprine and prednisolone, which can modify the severity of gingival overgrowth (Wilson *et al*, 1998). In contrast patients on cyclosporin A who are also receiving calcium channel blockers present with a greater risk of the gingival lesions than patients treated with cyclosporine alone (Thomasson *et al*, 1997). However, in the present study, both groups, i.e. patients with and without gingival overgrowth were medicated similarly during observation period. So, the effect of treatment modality of gingival pathology could be neglected in the data analysis.

Another factor which could potentially influence the study conclusions is a number of evaluated cases. As in the present study analysed groups comprised of 84 and 140 cases the conclusions drawn should be considered as preliminary data, and the results of the study should be confirmed by observations from other populations involving larger groups of patients.

Based on the results from the present study it can be concluded that there is no significant association between the *TGFB1* gene polymorphism and gingival overgrowth in kidney transplant patients administered cyclosporine A as a principal immunosuppressive agent.

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Conflict of interest

There is no conflict of interest.

Author contributions

Dr Kozak and Dr Dziwanowski were responsible for patient evaluation and data analysis, Dr Wajda, Dr Lapczuk and Dr Kurzawski were responsible for experiments, study design and manuscript preparation, Dr Lipski and Dr Drozdziak M were responsible for research design and manuscript revisions.

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References

Afonso M, de Oliveira Bello V, Shibli JA, Sposto MR (2003) Cyclosporine A-induced gingival overgrowth in renal transplant recipients. *J Periodontol* **74**: 51-56.

Anderson TJ, Lapp CA, Billman MA, Schuster GS (1998) Effects of transforming growth factor-beta and platelet derived growth factor on human gingival fibroblasts grown in serum-containing and serum-free medium. *J Clin Periodontol* **25**: 48-55.

Buduneli N, Kutukculer N, Aksa G, Atilla G (2001) Evaluation of transforming growth factor-beta1 level in crevicular fluid of cyclosporin A-treated patients. *J Periodontol* **72**: 526-531.

Cotrim P, de Andrade CR, Martelli-Junior H, Graner E, Sauk JJ, Coletta RD (2002) Expression of matrix metalloproteinases in cyclosporine-related gingival fibroblasts is regulated by transforming growth factor (TGF)- β 1 autocrine stimulation. *J Periodontol* **73**: 1313-1322.

Dreyfuss JL, Veiga SS, Coulson-Thomas VJ *et al* (2010) Differences in the expression of glycosaminoglycans in human fibroblasts derived from gingival overgrowths is related to TGF-beta up-regulation. *Growth Factors* **28**: 24-33.

Edwards DR, Murphy G., Reynolds JJ *et al* (1987) Transforming growth factor beta modulates the expression of collagenase nad metalloproteinase inhibitor. *EMBO J* **6**: 1899-1904.

Ellis JS, Morgan CI, Kirby JA, Taylor JJ, Thomason JM (2004) Plasma TGF- β 1 as a risk factor for gingival overgrowth. *J Clin Periodontol* **31**: 863-868.

Grainger DJ, Mosedale DE, Metcalfe JC (2000) TGF- β in blood: a complex problem. *Cytokine Growth Factor Rev* **11**: 133-145.

Grigner DJ, Heathcote K, Chiano M *et al* (1999) Genetic control of the circulating concentration of transforming growth factor type β 1. *Hum Mol Genet* **8**: 93-97.

Gurkan A, Afacan B, Emingil G, Toz H, Baskesen A, Atilla G (2008) Gingival crevicular fluid transforming growth factor- β 1 in cyclosporine and tacrolimus treated renal transplant patients without gingival overgrowth. *Arch Oral Biol* **53**: 723-728.

James JA, Invin CR, Linden GJ (1998) Gingival fibroblast response to cyclosporine A and transforming factor beta 1. *J Periodontal Res* **33**: 40-48.

Linden GJ, Haworth SE, Maxwell P *et al* (2001) The influence of transforming growth factor- β 1 gene polymorphisms on the severity of gingival overgrowth associated with concomitant use of cyclosporine A and a calsium channel blocker. *J Periodontol* **72**: 808-814.

Margiotta V, Pizzo I, Pizzo G, Barbaro A (1996) Cyclosporine- and nifedipine-induced gingival overgrowth in renal transplant patients: correlations with periodontal and pharmacological parameters, and HLA-antigens. *J Oral Pathol Med* **25**: 128-134.

McGaw WT, Porter H (1988) Cyclosporin-induced gingival overgrowth: an ultrastructural

- stereologic study. *Oral Surg* **65**: 186-190.
- Overall CM, Warana JL, Sodek J (1989) Independent regulation of collagenase 72-kDa progelatinase and metalloproteinase inhibitor expression in human fibroblasts by transforming growth factor β . *J Biol Chem* **264**: 1860-1869.
- Pernu HE, Pernu HM, Huttunen KR, Nieminen PA, Knuuttila ML (1992) Gingival overgrowth among renal transplant recipients related to immunosuppressive medication and possible local background factors. *J Periodontol* **63**: 548-553.
- Pernu HE, Pernu LM, Huttunen KR, Nieminen PA, Knuuttila ML (2001) Gingival overgrowth among renal transplant recipients related to immunosuppressive medication and possible local background factors. *J Periodontol* **62**: 548-553.
- Pulley LJ, Newton R, Adcock IM, Barnes PJ (2001) TGF β 1 allele association with asthma severity. *Hum Genet* 2001, **109**: 623-629.
- Radwan-Oczko M, Boratynska M, Zietek M, Dobosz T (2008) Transforming growth factor- β 1 gene expression and cyclosporine A-induced gingival overgrowth: a pilot study. *J Clin Periodontol* **35**: 371-378.
- Radwan-Oczko M, Boratynska M, Zietek M, Zoledziewska M, Jonkisz A (2006) The relationship of transforming growth factor- β 1 gene polymorphism, its plasma level, and gingival overgrowth in renal transplant recipients receiving different immunosuppressive regimens. *J Periodontol* **77**: 865-873.
- Seymour RA, Thomason JM, Ellis JS (1996) The pathogenesis of drug-induced gingival overgrowth. *J Clin Periodontol* **23**: 165-175.
- Somacarrera ML, Hernandez G, Acero J, Moskow BS (1994) Factors relating to the incidence and severity of cyclosporine-induced gingival overgrowth in transplant patients. A longitudinal study. *J Periodontol Res* **65**: 671-675.
- Thomas DW, Baboolal K, Subtamanian N, Newcombe RG (2001) Cyclosporine A-induced gingival overgrowth is unrelated to allograft function in renal transplant recipients. *J Clin Periodontol* **28**: 706-709.
- Thomas DW, Newcombe RG, Osborne G (2000) Risk factors in the development of cyclosporine-induced gingival overgrowth. *Transplantation* **70**: 552-556.
- Thomasson JM, Ellis JS, Kelly PJ, Seymour RA (1997) Nifedipine pharmacological variables as risk factors for gingival overgrowth in organ-transplant patients. *Clin Oral Invest* **1**: 35-39.
- Wilson RF, Morel A, Smith D *et al* (1998) Contribution of individual drugs to gingival overgrowth in adult and juvenile renal transplant patients treated with multiple therapy. *J Clin Periodontol* **25**: 457-464.

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Yoshida T, Nagata J, Yamane A (2005) Growth factors and proliferation of cultured rat gingival cells In response to cyclosporin A. *J Periodontal Res* **40**: 11-19.

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Table 1 Patients' characteristics

Parameter	Patients with gingival overgrowth (n=84)	Patients without gingival overgrowth (n=140)	p
Gender distribution (F/M)	23/61	49/91	0.301
Age (years)*	42.2±13.1	39.7±11.4	0.233
Place of residence: city/country	44/40	83/57	0.332
Smoking	16	31	0.615
Diabetes	39	47	0.065
Acute rejection	3	20	0.011

All calculations performed by means of Fisher exact test, except *calculated by t-test

Table 2 Characteristics of medication at monthly intervals after transplantation

	1 month	2 months	3 months	4 months	5 months	6 months
patients with gingival overgrowth (n=84)						
Cyclosporine concentration (ng/ml)	420.0±234.2	380.1±206.7	308.3±170.2	306.3±198.2	276.5±193.3	280.8±179.4
Cyclosporine dose (mg/day) (n=84)	300.0±84.1	280.9±78.8	260.7±79.3	251.2±76.9	245.2±68.6	235.4±65.4
Diltiazem dose (mg/day) (n=56)	173.8±24.5	173.8±24.5	173.8±24.5	173.8±24.5	173.8±24.5	173.8±24.5
Verapamil dose (mg/day) (n=28)	174.5±72.2	174.5±72.2	174.5±72.2	174.5±72.2	174.5±72.2	174.5±72.2
Prednisone dose (mg/day) (n=84)	18.8±7.4	16.3±5.2	15.1±4.3	12.4±3.7	10.6±3.0	9.5±3.1
patients without gingival overgrowth (n=140)						
Cyclosporine concentration (ng/ml)	413.2±258.1	370.8±288.2	326.8±263.5	276.6±159.8	296.9±195.6	249.8±172.2
Cyclosporine dose (mg/day) (n=140)	282.4±99.1	274.4±88.2	256.5±85.4	254.9±83.6	246.2±77.2	238.4 ± 73.0
Diltiazem dose (mg/day) (n=104)	172.2±30.4	172.2±30.4	172.2±30.4	172.2±30.4	172.2±30.4	172.2±30.4
Verapamil dose (mg/day) (n=36)	184.3±60.5	184.3±60.5	184.3±60.5	184.3±60.5	184.3±60.5	184.3±60.5
Prednisone dose (mg/day) (n=140)	17.9±4.5	15.2±3.0	13.4±3.4	12.1±2.5	11.9±2.6	11.9±2.6

Table 3 Distribution of *tgfb1* alleles and genotypes among kidney transplant patients with and without gingival overgrowth

		Gingival Overgrowth (n=84)		Healthy gingiva (n=140)		p value	OR (95%CI)
		n	(%)	n	(%)		
Genotype -800G>A							
	GG	78	(92.9)	120	(85.7)	-	-
	GA	6	(7.1)	19	(13.6)	0.188	0.48 (0.18-1.27)
	AA	0	(0.0)	1	(0.7)	1.000	-
	minor allele carriers (GA+AA)	6	(7.1)	20	(14.3)	0.132	0.46 (0.17-1.20)
	minor allele frequency	6	(3.6)	21	(7.5)	0.103	
Genotype -509C>T							
	CC	43	(51.2)	61	(43.6)	-	-
	CT	32	(38.1)	63	(45.0)	0.306	0.72 (0.40-1.28)
	TT	9	(10.7)	16	(11.4)	0.657	0.80 (0.32-1.97)
	minor allele carriers (CT+TT)	41	(48.8)	79	(56.4)	0.272	0.74 (0.43-1.27)
	minor allele frequency	50	(29.8)	95	(33.9)	0.404	
Genotype +869T>C							
	TT	34	(40.5)	50	(35.7)	-	-
	TC	38	(45.2)	65	(46.4)	0.652	0.86 (0.47-1.55)
	CC	12	(14.3)	25	(17.9)	0.424	0.71 (0.31-1.55)
	minor allele carriers (TC+CC)	50	(59.5)	90	(64.3)	0.480	0.82 (0.47-1.42)
	minor allele frequency	62	(36.9)	115	(41.1)	0.404	
Genotype +915G>C							
	GG	72	(85.7)	120	(85.7)	-	-
	GC	12	(14.3)	20	(14.3)	1.000	1.00 (0.46-2.16)
	CC	0	(0.0)	0	(0.0)	1.000	-
	minor allele carriers (GC+CC)	12	(14.3)	20	(14.3)	1.000	1.00 (0.46-2.16)
	minor allele frequency	12	(7.1)	20	(7.1)	1.000	

All calculations performed by means of Fisher exact test, using homozygotes for a major (more frequent) allele or major allele frequency as reference.

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Table 4 Frequency of *TGFB1* haplotypes among kidney transplant patients with and without gingival overgrowth.

	<i>Gingival Overgrowth</i> (2n=168)		<i>Healthy gingiva</i> (2n=280)		<i>p value</i>
	<i>n</i>	(%)	<i>n</i>	(%)	
G_C_T_G	100	(59.5)	144	(51.4)	*0.116
G_T_C_G	50	(33.9)	95	(29.8)	0.236
G_C_C_C	12	(7.2)	20	(7.1)	0.848
A_C_T_G	6	(7.5)	21	(3.6)	0.063

All calculations performed by means of Fisher exact test, using frequency of major haplotype as reference, except* - in relation to the sum of other haplotypes.

EH program (Jurg Ott, Rockefeller University, New York) was used to estimate haplotype frequencies based on results of typing for following SNPs: -800G>A, -509T>C, +869T>C, +915G>C. Among 16 potentially possible haplotypes derived from 4 SNPs, only four were observed in the studied sample of kidney transplant patients, with respective frequencies: G_C_T_G (0.545), G_T_C_G (0.324), G_C_C_C (0.071), and A_C_T_G (0.060).

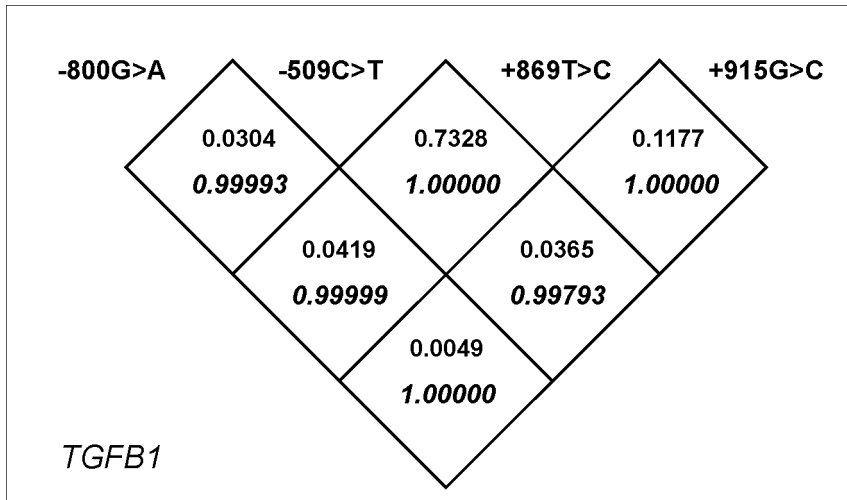


Figure 1. Pairwise LD between the studied *TGFB1* SNPs, estimated on base of analysis of all study subjects (n=224). Numbers represent r^2 (square correlation coefficient), and D' values (*italics*).