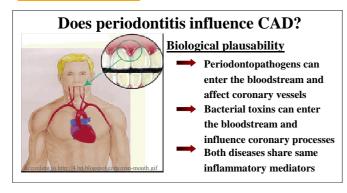
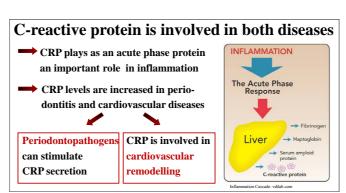
Are Genetic Variants of C-Reactive Protein Prognostic Markers for Further Cardiovascular Events in Patients With Coronary Heart Disease?

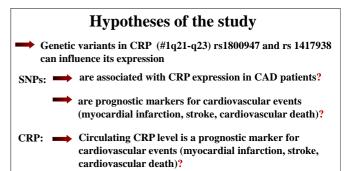
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Introduction







Material and Methods

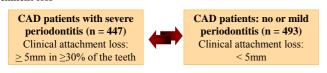
Cardiovascular patients

 $\label{longitudinal cohort study (Clinical Trials.gov Identifier: NCT01045070; n = 940) \\ Period of investigation: 10/2009-02/2011, Follow up: 11/2010-04/2012 \\$

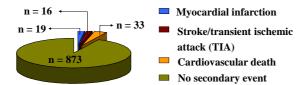
- Inclusion criteria: in-patient stay subjects with ≥50% stenosis of the main coronary artery German caucasian, ≥ 18 years of age, Presence of ≥ 4 teeth
- Exclusion criteria: periodontal treatment during the last 6 months, antibiotic therapy during the last 3 month, pregnancy
- Baseline clinical examination:

 Anamnesis → age, gender, smoking status, medication, existing diseases

 Dental examination → plaque index, bleeding on probing, pocket depth, clinical



1 year follow up:
 Evaluation of possible
 secondary cardiovascular
 events (telephone surveys
 or written surveys)



Genomic investigations

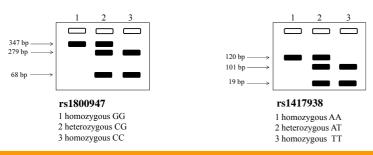
DNA-isolation from EDTA-blood

Preparation of genomic DNA was carried out using the blood extraction kit (Qiagen, Hilden, Germany). Genotyping of rs 1800947 and rs 1417938

Genotyping was performed by polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) PCR reactions were carried out using Mastermix (Promega, Mannheim, Germany)

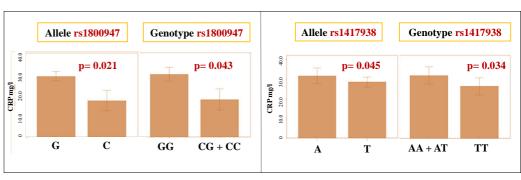
PCR-program (2min 94°C; 10 cycles: 15sec 94°C, 1min 64°C; 20 cycles: 15sec 94°C, 50sec 61°C, 30sec 72°C) $\mathbf{rs1800947} \rightarrow \mathbf{forward}$: cag ttt tac agt ggg tgg gtc, reverse: ccc gcc agt tca gga cat tag, restriction enzyme: BsiHKA I $\mathbf{rs1417938} \rightarrow \mathbf{forward}$: acc ccc at acct cag atc gaa, reverse: gac gtg acc atg gag aag ct, restriction enzyme: Tfi I

Schematic illustration of PCR fragments after restriction cleavage



Results and discussion

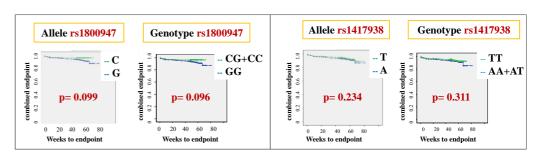
CRP serum level in dependence of SNPs



Allele and genotype of both SNPs rs1800947 (G-allele, GG-genotype) and rs1417938 (A-allele and AA + AT-genotypes) are significantly associated with increased CRP serum level.

CRP SNPs as prognostic markers for CAD

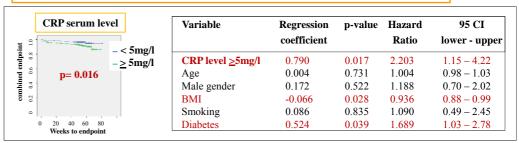
→ Kaplan-Meier survival curves and log-rank-test



The genotype and allele frequencies of both SNPs rs1800947 and rs1417938 were not prognostic markers for adverse cardiovascular events (myocardial infarction, stroke/TIA, cardiovascular death) regarding the one-year outcome.

Increased CRP serum level as a prognostic marker for CAD

→ Kaplan-Meier survival curve, log-rank-test, cox regression



■ In a complex risk model (cox regression) considering age, gender, body mass index (BMI), smoking, and diabetes as potential confounders, the CRP serum level could be proven as an independent prognostic indicator for adverse cardiovascular events regarding the one-year outcome.

Conclusions

CRP polymorphisms rs1800947 and rs1417938 are associated with CRP serum level in our cohort of CAD patients. Despite the CRP level as an independent prognostic marker for adverse cardiovascular events (myocardial infarction, stroke/TIA, cardiovascular death), neither polymorphism could be proven to have prognostic value regarding the one year outcome.



