Variation at the NFATC2 Locus Increases the Risk of Thiazolidinedione-Induced Edema in the Diabetes REduction Assessment with ramipril and rosiglitazone Medication (DREAM) Study

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ON BEHALF OF THE DREAM INVESTIGATORS

OBJECTIVE — Thiazolidinediones are used to treat type 2 diabetes. Their use has been associated with peripheral edema and congestive heart failure—outcomes that may have a genetic etiology.

RESEARCH DESIGN AND METHODS — We genotyped 4,197 participants of the multiethnic DREAM (Diabetes REduction Assessment with ramipril and rosiglitazone Medication) trial with a 50k single nucleotide polymorphisms (SNP) array, which captures \sim 2000 cardiovascular, inflammatory, and metabolic genes. We tested 32,088 SNPs for an association with edema among Europeans who received rosiglitazone (n=965).

RESULTS — One SNP, rs6123045, in *NFATC*2 was significantly associated with edema (odds ratio 1.89 [95% CI 1.47–2.42]; $P = 5.32 \times 10^{-7}$, corrected P = 0.017). Homozygous individuals had the highest edema rate (hazard ratio 2.89, $P = 4.22 \times 10^{-4}$) when compared with individuals homozygous for the protective allele, with heterozygous individuals having an intermediate risk. The interaction between the SNP and rosiglitazone for edema was significant ($P = 7.68 \times 10^{-3}$). Six SNPs in *NFATC*2 were significant in both Europeans and Latin Americans (P < 0.05).

CONCLUSIONS — Genetic variation at the *NFATC2* locus contributes to edema among individuals who receive rosiglitazone.

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Ithough changes in lifestyle can prevent or delay diabetes (1), the majority of patients require multiple therapeutic strategies to prevent or treat the disease. Thiazolidinediones (TZDs) are a class of drugs used in the treatment

of diabetes that derive their insulin sensitizing effects from the activation of the peroxisome proliferator–activated receptor γ (PPAR γ) (2). TZDs can effectively control glycemia among diabetic patients (3). However, their use has been shown to

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cause an increase in peripheral edema and congestive heart failure (CHF) (4,5). Edema is the most commonly reported adverse drug reaction associated with TZDs, and this has been partly attributed to the 6–8% increase in plasma volume that occurs with their use (6). In addition, the observed increase in CHF associated with rosiglitazone may derive from a shared etiology.

An aim of the Diabetes REduction and Assessment with ramipril and rosiglitazone Medication (DREAM) trial was to determine whether rosiglitazone could prevent progression to diabetes among individuals with impaired fasting glucose (IFG) and/or impaired glucose tolerance (IGT) (7). Consistent with previous findings, a significant increase in edema and CHF among individuals receiving rosiglitazone was observed (7). The identification of genetic variants that predispose individuals to edema or CHF could lead to pretherapeutic screening procedures. To determine whether genetic variation contributes to the etiology of TZDinduced edema, we tested common single nucleotide polymorphisms (SNPs), capturing ~2,000 cardiovascular/metabolic genes in DREAM trial participants receiving rosiglitazone.

RESEARCH DESIGN AND

METHODS— The DREAM trial has been described in detail elsewhere (7). We tested 32,088 SNPs for an association with TZD-induced peripheral edema in 965 European individuals receiving rosiglitazone. Edema was defined as the presence of pitting edema at both ankles reported at any clinic visit, and individuals that withdrew from treatment due to edema were included. We used logistic regression for each SNP adjusted for age, sex, BMI, and the use of ACE inhibitors and calcium channel blockers (CCBs). Individuals taking diuretics were excluded from all analyses. The first 10 principal components of the shared alleles (identical by state) were included as covariates.

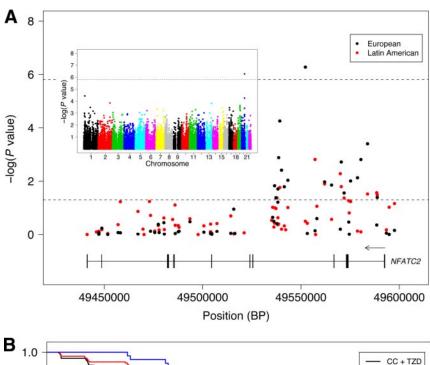
To test for interaction between the SNP and rosiglitazone, we performed a logistic regression analysis that included the main effects of the SNP, rosiglitazone, and their interaction term. Survival curves for each genotypic class and the corresponding hazard ratios (HRs) were calculated from a Cox proportional hazard analysis. A detailed description of all materials and methods is available in in an online appendix available at http://care.diabetesjournals.org/cgi/content/full/dc10-0452/DC1.

RESULTS — In our genetic substudy of DREAM, we observed an increase in edema among individuals who received rosiglitazone (n = 390 [22.3%]) versus placebo (n = 256 [14.5%]). Among the Europeans, 253 (26.2%) individuals receiving rosiglitazone experienced edema compared with 154 (16.1%) receiving placebo ($P = 8.63 \times 10^{-7}$) (supplementary Table 1). The clinical characteristics of the Europeans were not significantly different between the rosiglitazone and placebo arms (supplementary Table 2).

We tested 32,088 SNPs against edema in the Europeans receiving rosiglitazone. One SNP, rs6123045, in the nuclear factor of activated T-cells cytoplasmic calcineurin-dependent 2 (NFATC2) gene was significantly associated with edema (odds ratio [OR] 1.89 [95% CI 1.47–2.42]; $P = 5.32 \times 10^{-7}$, corrected P = 0.017) (Fig. 1A). The distribution of the observed versus the expected P values is shown in supplementary Fig. 1. We detected a significant interaction between rs6123045 and rosiglitazone treatment for edema in Europeans ($P = 7.68 \times 10^{-3}$). The effect of rs6123045, although in the same direction, was not significantly associated with edema in the placebo group (OR 1.16, P = 0.29) (Fig. 1B).

A Cox proportional hazards analysis revealed that individuals homozygous for the risk allele had a decrease in the time to the first report of edema in comparison with individuals heterozygous or homozygous for the protective allele (HR $1.76, P=3.43\times 10^{-5}$ and HR $2.89, P=4.22\times 10^{-4}$, respectively) (Fig. 1*B*). The effect appears to be additive because heterozygous individuals had an increased rate of edema compared with homozygous individuals (HR 1.64, P=0.11).

Among Europeans receiving rosiglitazone, rs6123045 was not significantly associated with diabetes or death or with



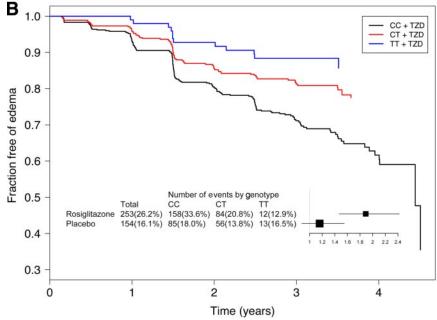


Figure 1—*A*: Results of the association analysis between SNPs and TZD-induced peripheral edema at the NFATC2 locus. The $-\log$ of the P values are plotted against SNP location. P values were calculated from a logistic regression analysis adjusted for age, sex, BMI, and use of ramipril and CCBs, as well as the first 10 principal components of the alleles shared identity by state among the European and Latin American individuals. Individuals taking diuretics were excluded from the analysis. The dashed lines indicate Bonferroni corrected and nominal significance. Inset: Results of the initial association scan of 32,088 SNPs and TZD-induced peripheral edema in Europeans (n = 965) receiving rosiglitazone. The -log of the P values are plotted against SNP location for each chromosome. B: Survival curves estimated from the Cox proportional hazards model of time to the first occurrence of edema according to the rs6123045 genotype. European individuals homozygous for the risk allele (CC) have an increase in the rate to the first report of edema in comparison with the individuals heterozygous (CT) or homozygous (TT) for the protective allele (adjusted HR 1.76, $P = 3.43 \times 10^{-5}$ and adjusted HR 2.89, P = 4.22×10^{-4} , respectively). Inset: The effect of the rs6123045 SNP on peripheral edema among European individuals receiving rosiglitazone or placebo. 33.6% (158 of 470) of individuals homozygous for the risk allele, 20.8% (84 of 403) of heterozygous individuals, and 12.9% (12 of 93) of individuals homozygous for the protective allele developed edema while receiving rosiglitazone compared with 18.0% (85 of 473), 13.8% (56 of 404), and 16.5% (13 of 79), respectively, while receiving placebo. The per-allele OR and 95% CI of the logistic regression analysis are shown.

cardiovascular end points, including CHF, myocardial infarction, stroke, angina, or a composite of these outcomes (data not shown).

rs6123045 was not significantly associated with TZD-induced edema in Latin Americans. However, six SNPs in NFATC2 were significant in both Europeans and Latin Americans (Fig. 1A). A haplotype defined by these SNPs is significant in both populations (OR 0.45 [95% CI 0.30-0.66]; $P = 2.26 \times 10^{-5}$ and OR $0.34 [95\% CI 0.13-0.90]; P = 1.47 \times$ 10^{-2} in Europeans and Latin Americans, respectively) (data not shown). All significantly associated SNPs were in Hardy-Weinberg equilibrium (P > 0.05).

CONCLUSIONS — We demonstrated that variation within NFATC2 contributes to TZD-induced edema. rs6123045 was significantly associated with TZDinduced edema among Europeans when corrected for multiple testing. The significant interaction between rs6123045 and rosiglitazone treatment for edema (P = 7.68×10^{-3}) among the European DREAM participants highlights its contribution to the etiology of TZD-induced edema. rs6123045 was not significantly associated with TZD-induced edema in Latin Americans. However, six SNPs in the same region are associated with TZDinduced edema and define a significantly associated haplotype in both populations.

Previous studies of either TZDinduced or dual PPAR agonist-induced edema (8–11) were smaller in size ($n \le$ 730) and tested fewer genes (\leq 222). None of these studies analyzed the NFATC2 gene. Of the 38 SNPs previously associated with edema, 20 are captured by the ITMAT-Broad-CARE (IBC) array used in our study, and we were thus able to directly test them. However, we were unable to replicate any of these associations in Europeans (P > 0.05).

The NFATC2 gene encodes a cytoplasmic component of the NFAT transcription complex. Four NFAT cytoplasmic component proteins (NFATc1c4) are known, and these are translocated to the nucleus after being dephosphorylated by the phosphatase calcineurin (12). Treatment of cardiomyocytes with rosiglitazone inhibited endothelin-1 induced calcineurin activity, suppressed the nuclear translocation of NFATc4, and enhanced the association of PPARy with calcineurin/NFATc4 (13). The constitutive activation of either calcineurin or NFATc4 in mice leads to cardiac hyper-

trophy and heart failure (14). In addition, Nfatc2 null mice are protected from calcineurin-induced cardiac hypertrophy (15). In the context of these findings, our results are provocative and constitute a step toward elucidating the etiology of the CHF associated with TZD use (5).

Identifying the specific genetic variants interacting with TZDs and resulting in edema or cardiovascular events may have important clinical consequences and enable genetic variant directed use of TZD drugs among people with dysglycemia.

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