ALLERGY, GENETICS, VITAMINS

## RS4711, UH RS7041

28.04.2010

The <u>first GWAS of human vitamin serum D</u> level finds the most important SNPs:

In a genome-wide association study (GWAS) of 4,501 persons of European ancestry drawn from five cohorts, we identified single nucleotide polymorphisms (SNPs) in the gene encoding group-specific component (vitamin D binding) protein, GC, on chromosome 4q12-13 that were associated with 25(OH)D concentrations: rs2282679 (P=2.0 x  $10\hat{a}$ €"30), in LD with rs7041, a nonsynonymous SNP (D432E; P=4.1 x 10-22), and rs1155563 (P =  $3.8 \times 10\hat{a}$ €"25).

Funny, rs7041 is the same variant that <u>we typed earlier</u> with <u>limited success</u> – maybe I should have also tested for FEV1 in adults only?? At least a <u>new COPD study this month in Thorax</u> arrives at this conclusion.

In patients with COPD, 25-OHD levels correlated significantly with forced expiratory volume in 1 s (FEV1) (r=0.28, p<0.0001) [...] Logistic regression corrected for age, gender and smoking history further revealed that homozygous carriers of the rs7041 T allele exhibited an increased risk for COPD (OR 2.11; 95% CI 1.20 to 3.71; p=0.009).

What really worries me – all these lines are not drawn in the original papers. A most recent review on COPD genetics even fignores GC as a candidate, nay, nay.

## Addendum 12-6-2010

The 2nd genome screen confirms GC with GWAS significance at

4p12 (overall p=1 $\hat{A}\cdot 9\tilde{A}$ —10 $^-9$  for rs2282679, in GC); 11q12 (p=2 $\hat{A}\cdot 1\tilde{A}$ —10 $^2$ 27 for rs12785878, near DHCR7); and 11p15 (p=3 $\hat{A}\cdot 3\tilde{A}$ —10 $^2$ 20 for rs10741657, near CYP2R1). Variants at an additional locus (20q13, CYP24A1) were genome-wide significant in the pooled sample (p=6 $\hat{A}\cdot 0\tilde{A}$ —10 $\hat{A}$ 10 for rs6013897).

https://www.wjst.de/blog/sciencesurf/2010/04/rs4711-uh-rs7041/ Page 2

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