

Table 1. Genotype distribution for the H324Q (C→A) SNP in *LARS2* in four different cohorts.

Cohort	NGT		T2DM		<i>P</i>	<i>P</i> _{adj}	OR _{adj} (95% CI)
	CC	CA+AA	CC	CA+AA			
Hoorn Study	323	13 (0.039)	193	18 (0.085)	0.04	0.05	2.19 (1.00 – 4.85)
ERGO (R'dam) Study	432	25 (0.055)	422	35 (0.077)	0.23	0.27	1.37 (0.77 – 2.42)
DK1, Steno case-control study	491	23 (0.045)	667	39 (0.055)	0.43	0.11	1.58 (0.90 – 2.78)
DK2, Inter99 case-control study	4248	253 (0.056)*	605	49 (0.075)	0.06	0.05	1.43 (1.00 – 2.05)

*P*_{adj} and OR_{adj} values were obtained after regression analysis with adjustment for age and gender. * including 3 AA homozygotes.