

Appendix 2. Composition of the family sample used for linkage analysis of chromosome 17 and association analysis of the RUNX1 and RAPTOR loci.

Cohort	# families informative for linkage*			# families informative for association*			total # families
	old cohort†	new cohort	pooled cohort	TDT	PDT	FBAT	
German	42	41	83	71	84	86	102
American	73	118	191	397	421	431	477
Total	115	159	274	468	505	517	579

*All families informative for the TDT are also informative for the PDT, and all families informative for the PDT are also informative for the FBAT. In contrast, 305 families informative for the FBAT are not informative for linkage, and 62 families informative for linkage are not informative for FBAT.

†The old cohort refers to families previously analyzed for linkage¹

Reference

1. Nair RP, Henseler T, Jenisch S, *et al.* Evidence for two psoriasis susceptibility loci (HLA and 17q) and two novel candidate regions (16q and 20p) by genome-wide scan. *Hum Mol Genet* 1997;**6**(8):1349-1356.