

	USP18	FAM230A	TMEM191B	RIMBP3	PI4KAP1	PRODH	DGCR6	GGT3P
	1	3	0	0	0	1	1	1
	1	5	0	0	0	1	1	2
	1	4	1	1	1	1	1	1
	1	4	0	0	0	2	2	1
	1	6	0	0	0	1	1	2
	1	6	1	1	1	1	1	2
	1	7	0	0	0	1	1	3
	1	8	0	0	0	1	1	3
	1	8	1	1	1	1	1	3
	1	9	0	0	0	1	1	4
	1	10	1	1	1	1	1	4
	1	11	0	0	0	1	1	5
	1	12	1	1	1	1	1	5
	1	16	1	1	1	1	1	7

Fig. S8

Predicted gene copy number variation in the observed LCR22A alleles. Haplotypes from Fig.2A were divided according to their duplicon composition. Coding genes and pseudogenes were stated with their corresponding duplicon color. Gene count is based on the number of duplicons present in the haplotype.'