

Supplemental Table A. Regions of homozygosity found in controls.

#	Cohort	Region	Size
1	HapMap	2q32.3q33.1	7.7 Mb
2	HapMap	3q26.1	5.6 Mb
3	HapMap	6q22.1q22.31	6.5 Mb
4	CCF	4q34.1q35.1	7.25 Mb
5	CCF	9p24.1p23	4.2 Mb
6	CCF	13q14.11q21.31	22.4 Mb
7	CCF	13q21.3-q32.1	4.95 Mb
8	CCF	13q32.1q33.3	13.6 Mb
9	CCF	15q23q26.1	20.3 Mb