

Supplementary Table 1. “ROH” region description

Region	N.ind	CHR	SNP1	SNP2	BP1	BP2	KB
S1	9	5	rs301286	rs414536	82638711	84935202	2296.492
S2	9	11	rs4625425	rs11040730	46289469	48315844	2026.376
S3	8	2	rs7585990	rs17049898	59086213	59123264	37.052
S4	8	5	rs1966938	rs9314227	93608749	97581046	3972.298
S5	8	5	rs26522	rs974073	102512298	105493632	2981.335
S6	8	5	rs12109973	rs7724997	167204911	167305246	100.336
S7	7	4	rs3796585	rs1375138	156858624	159061598	2202.975
S8	7	5	rs17828812	rs9986158	64886188	67344385	2458.198
S9	7	10	rs7916320	rs2505731	26421818	31556425	5134.608
S10	6	1	rs601354	rs2877978	179663757	181763281	2099.525
S11	6	1	rs7554297	rs12563435	188030954	190163376	2132.423
S12	6	3	rs721213	rs157297	5774685	6354792	580.108
S13	6	5	rs7731628	rs13175654	76647020	79673555	3026.536
S14	6	5	rs248998	rs3777041	79837276	82536051	2698.776
S15	6	5	rs6861829	rs12658199	109911233	113349823	3438.591
S16	6	5	rs7731162	rs11744584	170503280	173348013	2844.734
S17	6	6	rs12662158	rs1107495	6671926	7671056	999.131
S18	6	10	rs10786961	rs7080675	83954103	88935321	4981.219
S19	6	15	rs1834212	rs10520810	93876557	94450872	574.316
S20	6	17	rs3760371	rs4794295	43050913	47099992	4049.08

N. ind = Number of Individuals sharing that Region; **Chr** = Chromosome; **SNP1**= SNP where the Region starts; **SNP2**= SNP where the Region ends; **BP1**= genomic position where the Region starts; **BP2**= genomic position where the Region ends; **N. SNPs** = Number of SNPs within the Region.