

Gene	Marker	Position	Group	Allele	Count	Freq	CI-Lower	CI-Upper	
1	ALL	rs709	?	ALL	1	2534	0.998	0.997	1.000
2				ALL	2	4	0.002	0.000	0.003
3				ALL	NA	0			
4									
5	rs534	?		ALL	1	1638	0.647	0.629	0.666
6				ALL	2	888	0.333	0.334	0.371
7				ALL	NA	0			
8									
9	rs47	?		ALL	1	924	0.371	0.382	0.390
10				ALL	2	1264	0.629	0.610	0.648
11				ALL	NA	0			
12									
13	rs45	?		ALL	1	990	0.393	0.376	0.414
14				ALL	2	1518	0.608	0.586	0.624
15				ALL	NA	0			
16									
17	rs79	?		ALL	1	1556	0.625	0.607	0.644
18				ALL	2	932	0.375	0.386	0.390
19				ALL	NA	0			
20									
21	rs31	?		ALL	1	346	0.230	0.214	0.247
22				ALL	2	1812	0.769	0.753	0.786
23				ALL	NA	0			
24									
25	rs91	?		ALL	1	2499	0.989	0.983	0.993
26				ALL	2	27	0.011	0.007	0.013
27				ALL	NA	0			
28									
29	rs23	?		ALL	1	2030	0.799	0.783	0.814
30				ALL	2	511	0.200	0.186	0.217
31				ALL	NA	0			
32									

Confidence intervals width is 95%, computed using the exact quantiles for the binomial distribution.