

Supplementary Table 3. Results of the Family-based Association Analysis in 156 MS Families from the Southern Ostrobothnia

Gene	SNP	Location ^a	P-value ^b
<i>C7</i>	rs1901167	40996921	0.094
<i>C7</i>	rs324061	40999587	0.530
<i>C7</i>	rs13157656	41000609	0.122
<i>C7</i>	rs3805221	41001075	0.148
<i>C7</i>	rs3805226	41009762	0.056
<i>C7</i>	rs3828511	41012092	0.029
<i>C7</i>	rs6860438	41015419	0.006
<i>C7</i>	rs1055021	41015923	0.018
Intergenic	rs7732104	41019193	0.210
Intergenic	rs4957361	41021361	0.571
Intergenic	rs13161930	41021841	0.041
Intergenic	rs2122564	41023131	0.087
Intergenic	rs1376174	41023856	0.031
Intergenic	rs7736582	41027101	0.181
<i>FLJ40243</i>	rs896117	41035176	0.061
<i>FLJ40243</i>	rs1530812	41035909	0.029
<i>FLJ40243</i>	rs3817324	41035949	0.008
<i>FLJ40243</i>	rs2167229	41036780	0.019
<i>FLJ40243</i>	rs16870603	41038515	0.497
<i>FLJ40243</i>	rs7712140	41040153	0.511
<i>FLJ40243</i>	rs2305314	41040564	0.079
<i>FLJ40243</i>	rs1450658	41044999	0.756
<i>FLJ40243</i>	rs1450659	41051060	0.918
<i>FLJ40243</i>	rs10058056	41055699	0.331

^a Location on Chromosome 5 according to UCSC Genome Browser, Mar 2006 freeze.

^b Gamete Competition Association Results.