

Supplementary Table S6. Resampling results of studies included for meta-analysis of MBL-2 polymorphisms [A>B (Asian), A>O (European), A>C, A>D, Y>X and H>L).

Genotype/allele	HC	SLE	OR	95% CI	P-value
Codon 52 A>B (Asian)					
AA	805063	624585	1	--	ref
AB	227133	224256	1.273	1.264 to 1.281	<0.0001
BB	37014	54844	1.910	1.884 to 1.936	<0.0001
AB+BB	264147	279100	1.362	1.353 to 1.370	<0.0001
A	1837259	1473426	1	--	ref
B	301161	333944	1.383	1.375 to 1.390	<0.0001
Codon combined A>O (European)					
AA	633958	570510	1	--	ref
AO	367852	378165	1.142	1.136 to 1.149	<0.0001
OO	61972	75543	1.355	1.339 to 1.370	<0.0001
AO+OO	429824	453708	1.173	1.167 to 1.179	<0.0001
A	1635768	1519185	1	--	ref
O	491796	529251	1.159	1.154 to 1.164	<0.0001
Codon 57 A>C					
AA	874454	756035	1	--	ref
AC	86753	78895	1.052	1.041 to 1.063	<0.0001
CC	6875	5988	1.007	0.9730 to 1.043	0.684
AC+CC	93628	84883	1.049	1.038 to 1.059	<0.0001
A	1835661	1590965	1	--	ref
C	100503	90871	1.043	1.034 to 1.053	<0.0001
Codon 52 A>D					
AA	774693	663839	1	--	ref
AD	49611	54962	1.293	1.277 to 1.309	<0.0001
DD	9001	7894	1.023	0.9928 to 1.055	0.1368
AD+DD	58612	62856	1.251	1.237 to 1.266	<0.0001
A	1598997	1382640	1	--	ref
D	67613	70750	1.210	1.197 to 1.223	<0.0001
-221 Y>X					
YY	192376	273776	1	--	ref
YX	120927	106873	0.621	0.6148 to 0.6273	<0.0001
XX	38012	39036	0.721	0.7107 to 0.7327	<0.0001
YX+XX	158939	145909	0.6451	0.6392 to 0.6510	<0.0001
Y	505679	654425	1	--	ref
X	196951	184945	0.725	0.7203 to 0.7309	<0.0001
-550 H>L					
HH	142296	152472	1	--	ref
HL	214682	203046	0.882	0.874 to 0.891	<0.0001
LL	205919	224585	1.018	1.008 to 1.027	0.0002
HL+LL	420601	427631	0.948	0.940 to 0.956	<0.0001
H	499274	507990	1	--	ref
L	626520	652216	1.023	1.018 to 1.029	<0.0001

Note: Data are number of subjects calculated by 1000 re-sampling groups. HC= healthy controls; SLE: systemic lupus erythematosus; OR = odds ratio; 95% CI = 95% confidence interval.