

Supplementary Table 1. Association between secretor MBL2 polymorphisms and survival in white patients*

Genotype	No. (%)	5-year overall survival, RR of death† (95% CI)
Promoter SNP		
mb12_11 (<i>rs11003125</i>)	215 (41.1)	
C/C (<i>L/L</i>)	238 (45.5)	1.0 (referent)
C/G (<i>L/H</i>)	70 (13.4)	0.86 (0.65 to 1.15)
G/G (<i>H/H</i>)		1.43 (0.99 to 2.07)
mb12_12 (<i>rs7096206</i>)	329 (61.7)	
G/G (<i>Y/Y</i>)	180 (33.8)	1.0 (referent)
G/C (<i>Y/X</i>)	24 (4.5)	0.59 (0.45 to 0.79)
C/C (<i>X/X</i>)	204 (38.3)	0.81 (0.44 to 1.50)
G/C + C/C‡		0.62 (0.47 to 0.81)
Exon 1 SNP		
mb12_03 (<i>rs5030737</i>)		
C/C (<i>Non-D/Non-D</i>)	447 (85.0)	1.0 (referent)
C/T (<i>Non-D/D</i>)	75 (14.2)	0.94 (0.66 to 1.35)
T/T (<i>D/D</i>)	4 (0.8)	2.60 (0.64 to 10.66)
mb12_01 (<i>rs1800450</i>)		
G/G (<i>Non-B/Non-B</i>)	391 (75.5)	1.0 (referent)
G/A (<i>Non-B/B</i>)	120 (23.2)	0.84 (0.62 to 1.14)
A/A (<i>B/B</i>)	7 (1.3)	2.45 (0.90 to 6.70)
mb12_02 (<i>rs1800451</i>)		
G/G (<i>Non-C/Non-C</i>)	479 (97.0)	1.0 (referent)
G/A (<i>Non-C/C</i>)	14 (2.8)	0.91 (0.37 to 2.22)
A/A (<i>C/C</i>)	1 (0.2)	—
Haplotype§		
GGCGG (<i>HYA</i>)	294 (28.6)	1.0 (referent)
CGCGG (<i>LYA</i>)	280 (27.3)	1.03 (0.82 to 1.31)
CCCGG (<i>LXA</i>)	220 (21.4)	0.71 (0.54 to 0.92)
CGCAG (<i>LYB</i>)	135 (13.2)	0.90 (0.67 to 1.21)
CGCGA (<i>LYC</i>)	16 (1.6)	0.75 (0.31 to 1.84)
GGTGG (<i>HYD</i>)	81 (7.9)	0.95 (0.67 to 1.36)
Diplotype		
YA/YA	159 (31.0)	1.0 (referent)
YA/XA	124 (24.2)	0.66 (0.46 to 0.95)
XA/XA	23 (4.5)	0.84 (0.44 to 1.58)
YA/B	78 (15.2)	0.95 (0.65 to 1.39)
XA/B	30 (5.8)	0.38 (0.19 to 0.74)
YA/C	8 (1.5)	1.25 (0.45 to 3.43)
YA/D	46 (9.0)	0.91 (0.57 to 1.45)
XA/D	17 (3.3)	0.57 (0.26 to 1.23)
XA/C	3 (0.6)	—
O/O	25 (4.9)	1.02 (0.55 to 1.88)

*RR = risk ratio; CI = confidence interval; SNP = single-nucleotide polymorphism; — = insufficient data.

Statistically significant associations are in boldface type.

†Risk ratio of death was adjusted for sex, stage (III–IV vs. I–II), age at diagnosis, current smoking status, and pack-years of smoking.

‡A dominant model was assumed.

§SNPs are ordered in the haplotypes by location on the gene, as L/H, Y/X, A/D, A/B, and A/C.