

Supplementary Information

Supplementary Table 1. Allele and genotype frequencies of the nine variants genotyped in both the discovery and the replication cohorts. In the replication cohort, DNA was insufficient to genotype the nine variants in all patients; therefore, the number of genotyped patients varies by variant. HWE=Hardy Weinberg Equilibrium; MAF=minor allele frequency

			Discovery Cohort (n=67)			Replication Cohort (n=74-104)		
Gene	Variant (rs#)	Function	N (%)	MAF	HWE	N (%)	MAF	HWE
ABCB1	IVS9 -44 A>G (rs10276036)	Intronic		0.46	0.01	77	0.35	0.82
AA			25 (37)			32 (41)		
AG			23 (34)			36 (47)		
GG			19 (28)			9 (12)		
ABCC1	IVS11 -48 C>T (rs3765129)	Intronic		0.17	0.08	74	0.17	0.08
CC			48 (72)			49 (66)		
CT			15 (22)			25 (34)		
TT			4 (6)			0 (0)		
ABCC1	1684 T>C (rs35605)	Synonymous, Exon 13		0.23	0.78	95	0.22	0.03
TT			40 (60)			54 (57)		
TC			23 (34)			40 (42)		
CC			4 (6)			1 (1)		
ABCC2	-24 C>T (rs717620)	5' UTR		0.18	0.34	94	0.17	0.21
CC			44 (66)			63 (67)		
CT			22 (33)			30 (32)		
TT			1 (1)			1 (1)		
ABCC2	3972C>T (rs3740066)	Synonymous, Exon 28		0.34	0.81	104	0.36	0.84
CC			30 (45)			43 (41)		
CT			29 (43)			47 (45)		
TT			8 (12)			14 (14)		
SLCO1B1*1b	388 A>G (rs2306283)	Nonsynonymous, Exon 5 (N130D)		0.46	0.58	84	0.40	0.73
AA			21 (31)			29 (35)		
AG			31 (46)			42 (50)		
GG			15 (22)			13 (15)		
SLCO1B1*5	521 T>C (rs4149056)	Nonsynonymous, Exon 6 (V174A)		0.19	0.79	83	0.13	0.74
TT			44 (66)			63 (76)		
TC			21 (31)			19 (23)		
CC			2 (3)			1 (1)		
UGT1A1*93	-3156 G>A (rs10929302)	Intronic		0.32	0.24	103	0.28	0.57
GG			33 (49)			52 (50)		
GA			25 (37)			44 (43)		
AA			9 (13)			7 (7)		
UGT1A1*28	TA6>TA7 (rs8175347)	TA repeat, Promoter		0.34	0.25	75	0.33	0.11
*1/*1			31 (46)			31 (41)		
*1/*28			26 (39)			39 (52)		
*28/*28			10 (15)			5 (7)		