

Supplemental Table 2. Minor allele frequencies (MAF) of single nucleotide polymorphisms (SNPs) selected for genetic association analysis of bosentan-induced liver injury in Caucasians.

Gene	SNP¹	MAF (%)	Reason for filtering
<i>ABCC2</i>	rs717620	23.2	
	rs2273697	21.4	
	rs17222723	2.7	MAF<5%
	rs8187710	2.7	MAF<5%
	rs1885301	42.9	
	rs7910642	8.0	
	rs2804402	57.1	
<i>ABCB11</i>	rs2287622	45.5	
<i>SLCO1B1</i>	rs11045819	11.6	
	rs2306283	34.8	
	rs4149056	18.8	
<i>SLCO1B3</i>	rs414917	17.0	
	rs7311358	16.1	In LD with rs414917
<i>CYP2C9</i>	*2/ rs1799853	12.5	
	*3/ rs1057910	8.0	

¹Fifteen SNPs in five genes encoding proteins involved in bosentan disposition were selected based on literature review. Twelve SNPs (in bold) were analyzed after filtering for MAF (<5%) and linkage disequilibrium ($r^2 > 0.8$).