

SUPPLEMENTARY TABLE S3. TOP SINGLE NUCLEOTIDE POLYMORPHISMS ASSOCIATED WITH DRUG-INDUCED HEPATOTOXICITY IN PHARMACOKINETIC GENES AND HUMAN LEUKOCYTE ANTIGEN REGION FOR ANTIRETROVIRAL DRUG-ALONE AND ANTIRETROVIRAL AND ANTI-TUBERCULOSIS DRUGS CO-TREATMENT GROUPS

<i>Group</i>	<i>SNP</i>	<i>Chr (loci)</i>	<i>Alleles</i>	<i>p_min</i>	<i>p_adj</i>	<i>OR (95% CI)</i>	<i>MAF</i>	<i>Gene/loci</i>
ARV	rs11642957	16 (16124008)	T/C	9.5×10^{-5}	1.1×10^{-4}	4.9 (2.3–10.7)	0.18	<i>ABCC1</i>
	rs9380345	6 (33080359)	T/C	4.6×10^{-3}	1.0×10^{-3}	8.6 (2.4–31.2)	0.03	<i>HLA-DPB2</i>
	rs2857204	6 (32744347)	A/C	5.3×10^{-3}	8.8×10^{-3}	3.3 (1.1–10.0)	0.42	<i>HLA-DQB2</i>
	rs13199787	6 (32705276)	T/C	6.6×10^{-3}	2.3×10^{-2}	1.9 (0.6–6.4)	0.43	<i>HLA-DQA2</i>
	rs2860975	10 (96766934)	A/C	2.0×10^{-3}	2.5×10^{-2}	7.8 (1.7–35.7)	0.27	<i>CYP2C9</i>
	rs1569343	4 (70341463)	A/C	4.6×10^{-3}	2.6×10^{-2}	4.9 (1.6–15.2)	0.46	<i>UGT2B4</i>
	rs12024396	1 (97880883)	A/G	1.1×10^{-3}	2.8×10^{-2}	6.7 (2.0–21.8)	0.46	<i>DPYD</i>
ARV-ATD	rs9276370	6 (32707295)	A/C	2.8×10^{-4}	1.1×10^{-3}	5.1 (2.0–12.7)	0.36	<i>HLA-DQA2</i>
	rs7773149	6 (32706042)	A/G	3.5×10^{-4}	1.2×10^{-3}	3.8 (1.7–8.3)	0.36	<i>HLA-DQA2</i>
	rs9341244	2 (38303095)	A/G	3.8×10^{-3}	2.1×10^{-3}	8.8 (2.2–35.3)	0.02	<i>CYP1B1</i>
	rs7592624	2 (234602906)	A/G	3.3×10^{-3}	5.2×10^{-3}	4.5 (1.7–11.6)	0.38	<i>UGT1A8</i>
	rs11203943	8 (18070927)	T/C	3.9×10^{-3}	6.6×10^{-3}	4.7 (1.7–12.9)	0.06	<i>NAT1</i>
	rs7670819	4 (70481880)	T/C	2.9×10^{-3}	6.0×10^{-2}	16.8 (9.2–30.9)	0.45	<i>UGT2A1</i>