

Results under standard GWAS and MultiPhen approaches for genome-wide significant SNPs: TRIG-LDL combination

Sig. SNPs	CHOL	TRIG	HDL	LDL	MultiPhen	Order diff
rs629301	-	-0.08	-	<b>12.43</b>	<b>12.19</b>	-0.24
rs4420638	-	1.08	-	<b>12.91</b>	<b>9.73</b>	-3.18
rs174546	-	2.94	-	5.11	<b>9.67</b>	4.56
rs1042034	-	5.26	-	6.91	<b>9.04</b>	2.13
rs964184	-	<b>10.96</b>	-	1.46	<b>8.65</b>	-2.31
rs1367117	-	0.22	-	<b>9.49</b>	<b>8.12</b>	-1.37
rs6511720	-	0.63	-	<b>8.61</b>	6.98	-1.63
rs1260326	-	<b>8.04</b>	-	0.46	6.71	-1.33