

Table S6. Number of Ensembl genes with independent *cis*-eQTLs at the 0.01 permutation threshold, as determined by stepwise association model.

No. of independent <i>cis</i> - eQTLs	0.01 permutation threshold							
	CEU	CHB	GIH	JPT	LWK	MEX	MKK	YRI
1	645	765	697	788	772	472	940	786
2	10	6	1	6		0	7	11
3	2	2						2
4		1		1				
5					1			
<b>Total</b>	657	774	698	795	773	472	947	799
<b>genes with <math>\geq 2</math> <i>cis</i>-eQTLs</b>	12	9	1	7	1	0	7	13
<b>% genes with <math>\geq 2</math> <i>cis</i>-eQTLs</b>	1.83	1.16	0.14	0.88	0.13	0.00	0.74	1.63