

**Supplemental Table 1** Annotation and minor allele frequencies for TRIM SNPs of interest.

Gene	Chromosome	rsID	Chromosomal position	Gene location	Major allele	Minor allele	MAF Caucasians	MAF African-Americans
TRIM25	17	rs11540270	52321803	3UTR	A	G	7%	12%
		rs2525998	52322526	3UTR	G	A	29%	41%
		rs2525997	52326239	intron	A	C	9%	14%
		rs205499	52327399	intron	G	A	21%	24%
		rs2525996	52331302	intron	G	C	25%	13%
		rs2525994	52331920	intron	A	C	25%	18%
		rs2525993	52332778	intron	A	G	25%	14%
		rs205498	52333792	coding	A	G	24%	23%
		rs7225205	52335979	intron	G	A	12%	25%
		rs9909750	52337801	intron	G	A	12%	18%
		rs2525992	52343180	intron	A	G	25%	16%
		rs169530	52343239	intron	G	A	19%	11%
		rs11869863	52348353	flanking_5UTR	T	A	13%	18%
TRIM5	11	rs7943397	5639521	flanking_3UTR	C	G	41%	24%
		rs7122620	5640142	flanking_3UTR	A	G	46%	47%
		rs11038628	5645515	coding	G	A	6%	24%
		rs12808061	5646550	intron	C	A	12%	27%
		rs12271333	5647266	intron	A	C	12%	16%
		rs16934050	5648364	intron	C	A	6%	25%
		rs904375	5648555	intron	G	A	18%	47%
		rs3740994	5656376	intron	A	C	15%	11%
		rs3740995	5656915	coding	G	A	14%	22%
		rs10769175	5657224	intron	G	A	20%	48%
		rs10838525	5657576	coding	G	A	37%	20%
		rs11601507	5657649	coding	C	A	7%	1%
		rs3740996	5657856	coding	G	A	12%	8%
		rs3824949	5657984	UTR	C	G	44%	74%
		rs7124435	5658394	intron	A	G	2%	11%
		rs17305868	5658458	intron	G	A	48%	48%
		rs7127767	5658674	intron	A	G	47%	48%

Gene	Chromosome	rsID	Chromosomal position	Gene location	Major allele	Minor allele	MAF Caucasians	MAF African-Americans
		rs12278842	5660133	intron	G	C	47%	46%
		rs3802981	5663098	flanking_5UTR	A	G	48%	48%

3UTR, three-prime untranslated region; 5UTR, five-prime untranslated region; MAF, minor allele frequency; one SNP having a MAF of 74% in African-Americans is the result of requiring consistent alleles between cohorts.