









































Population Set 1: CLM (N=60), MXL (N=66)

SNP	Population Set 1 (N=126)		1000 Genomes Phase I summaries
rs1077835	A: 0.587 G: 0.413		
rs11216230	G: 0.944 A: 0.056		
rs13392272	C: 0.631 T: 0.369		
rs1532624	C: 0.639 A: 0.361		
rs2126259	T: 0.242 C: 0.758		
rs2238675	C: 0.901 T: 0.099		
rs2278426	C: 0.813 T: 0.187		
rs2367970	G: 0.730 A: 0.270		
rs2472386	G: 0.532 A: 0.468		
rs261334	C: 0.631 G: 0.369		
rs28526159	C: 0.456 T: 0.544		
rs4791641	C: 0.671 T: 0.329		
rs5880	G: 0.933 C: 0.067		
rs660240	T: 0.187 C: 0.813		
rs7121538	A: 0.889 C: 0.111		
rs7499892	C: 0.857 T: 0.143		
rs884366	G: 0.575 A: 0.425		
rs9282541	G: 0.952 A: 0.048		
rs7528419	A: 0.821 G: 0.179		
rs964184	G: 0.238 C: 0.762		

rs1250229	T: 0.194 C: 0.806	T C	T C	EUR AME EAS AFR
rs2235776	C: 0.726 T: 0.274	C T	C T	EUR AME EAS AFR
rs118146573	G: 0.893 A: 0.107	G A	G A	EUR AME EAS AFR
rs3902354	C: 0.278 A: 0.722	C A	C A	EUR AME EAS AFR
rs79236614	C: 0.929 G: 0.071	C G	C G	EUR AME EAS AFR
rs28680850	G: 0.611 A: 0.389	G A	G A	EUR AME EAS AFR
rs78536982	C: 0.921 T: 0.079	C T	C T	EUR AME EAS AFR
rs62436827	A: 0.909 G: 0.091	A G	A G	EUR AME EAS AFR
rs4360309	C: 0.242 T: 0.758	C T	C T	EUR AME EAS AFR
rs72925845	G: 0.948 A: 0.052	G A	G A	EUR AME EAS AFR
rs8102280	G: 0.952 A: 0.048	G A	G A	EUR AME EAS AFR
rs780093	T: 0.421 C: 0.579	T C	T C	EUR AME EAS AFR
rs2954031	G: 0.623 T: 0.377	G T	G T	EUR AME EAS AFR
rs4588572	A: 0.540 G: 0.460	A G	A G	EUR AME EAS AFR
rs11571087	T: 0.000 A: 1.000	T A	T A	EUR AME EAS AFR
rs12721308	A: 0.996 G: 0.004	A G	A G	EUR AME EAS AFR
rs34052647	G: 0.984 A: 0.016	G A	G A	EUR AME EAS AFR
rs10902456	G: 0.714 T: 0.286	G T	G T	EUR AME EAS AFR
rs1042034	C: 0.234 T: 0.766	C T	C T	EUR AME EAS AFR
rs4841132	A: 0.179 G: 0.821	A G	A G	EUR AME EAS AFR
rs3208305	A: 0.706 T: 0.294	A T	A T	EUR AME EAS AFR
rs66462329	G: 0.762 A: 0.238	G A	G A	EUR AME EAS AFR
rs2954038	C: 0.270 A: 0.730	C A	C A	EUR AME EAS AFR
rs2575876	G: 0.714 A: 0.286	G A	G A	EUR AME EAS AFR
rs174537	G: 0.413 T: 0.587	G T	G T	EUR AME EAS AFR
rs180360	A: 0.647	A G	A G	EUR AME EAS AFR

			EUR	AME	EAS	AFR
rs3135506	G: 0.353 G: 0.913 C: 0.087					
rs662799	G: 0.139 A: 0.861					
rs35853021	G: 0.659 T: 0.341					
rs8033940	G: 0.520 A: 0.480					
rs247617	C: 0.726 A: 0.274					
rs11076175	A: 0.865 G: 0.135					
rs5883	C: 0.964 T: 0.036					
rs16942887	G: 0.841 A: 0.159					
rs1800961	C: 0.964 T: 0.036					
rs12740374	G: 0.821 T: 0.179					
rs668948	G: 0.159 A: 0.841					
rs934197	G: 0.702 A: 0.298					
rs4245791	C: 0.179 T: 0.821					
rs12916	T: 0.575 C: 0.425					
rs3798168	C: 0.841 A: 0.159					
rs2980869	C: 0.567 T: 0.433					
rs6511720	G: 0.901 T: 0.099					
rs2074304	G: 0.956 C: 0.044					
rs7412	C: 0.929 T: 0.071					
rs631106	C: 0.659 A: 0.341					
rs11902417	G: 0.746 A: 0.254					
rs780094	T: 0.421 C: 0.579					
rs33951980	C: 0.929 T: 0.071					
rs10096633	C: 0.881 T: 0.119					
rs60223219	A: 0.933 G: 0.067					

[rs28927680](#)

G: 0.099
C: 0.901



[rs2266788](#)

G: 0.135
A: 0.865



[rs118029212](#)

C: 0.944
T: 0.056



[rs58542926](#)

C: 0.940
T: 0.060



[rs12721054](#)

A: 0.992
G: 0.008



[rs439401](#)

T: 0.433
C: 0.567



Population Set 1
(N=126)

6 populations/groups were queried, and 77 SNPs are displayed
database query ended in 8.899 seconds · results page generated in 8.952 seconds

SPSmart: [v5.1.1](#) · dbSNP version: [build 132](#)

selected database: **1000 Genomes Phase I** · data source: [1000 Genomes](#) · data version: [Phase I May 2011](#)

available databases on SPSmart: [1000 Genomes](#) · [HapMap](#) · [Perlegen](#) · [HGDP-CEPH](#) · [copyright and disclaimer](#)

