

Table 6. *MIP-1α*+113T is in complete linkage with *MIP-1α*+459T.

+459	+113		
	C/C	C/T	T/T
C/C	1,149	0	0
C/T	476	95	0
T/T	55	6	3

All individuals homozygous for *MIP-1α*+113T are also homozygous for *MIP-1α*+459T, and all individuals heterozygous for *MIP-1α*+113T are homozygous or heterozygous for *MIP-1α*+459T. This suggests that *MIP-1α*+113T mutation occurred in the context of a mutated *MIP-1α*+459T allele. Thus, this linkage pattern along with the sequencing data identified three different haplotypes designated CC, CT, and TT. Data is for the combined analysis of U.S.-based HIV+ individuals and control blood donors.