

S6 Table: Specific SNPs restricted to 3208-2007 and 3208-2015 strains compared to 24 other *M. leprae* genomes

Position	Name	Type	TN base (ref)	Alternate allele	SNP effect	Amino-acid change
489941	<i>ML0394</i>	protein coding	C	T	missense	Leu163Phe
630056	<i>aroB</i>	protein coding	C	A	synonymous	Leu163Leu
1552528	<i>ansP1</i>	protein coding	C	A	synonymous	Ser496Ser
1552530	<i>ansP1</i>	protein coding	T	C	missense	Ser496Pro
1552533	<i>ansP1</i>	protein coding	A	C	missense	Lys495Gln
1785100	-	Intergenic	G	A	-	-
1844335	<i>ML1529</i>	Pseudogene	C	T	-	-
1900827	-	Intergenic	G	A	-	-

SNP: single nucleotide polymorphism.