

S5 Table: Allelic diversity of VNTR loci in the recurrent leprosy cases.

Locus ID	Coordinates	1126		2188		3208	
		1126-2007	1126-2011	2188-2007	2188-2014	3208-2007	3208-2015
(T)8, & (A)6	337466-337473, 337474-337479	8 & 6	8 & 6	8 & 6	8 & 6	8 & 6	8 & 6
(T)6 & (N)7 & (T)8	514181-514186, 514187-514193, 514194-514200	6 & 7 & 8	6 & 7 & 8	6 & 7 & 8	6 & 7 & 8	6 & 7 & 8	6 & 7 & 8
(A)9	1414666-1414674	8	8	8	8	8	8
(G)9	976857-976865	9	10	9	9	9	9
(C)9	2658192-2658200	9	11	9/10	9/10	8/9	8/9
(G)10a	347280-347289	10	9	9	9	13	13
(G)10b	442993-443002	NA	?	9	9	?	9/10
(G)11	1309544-1309554	NA	11	9	9	10/11	12
(G)12	1116443-1116454	9/10	10	9/10	9	11	11
(CG)6	2947291-2947302	6	6	6	6	6	6
(AC)8a	1531185-1531200	9	8	8	8	9	9
(AC)8b	2211035-2211050	7	7/8	7/8	7/8	7	7
(AC)9	1452573-1452590	8	NA	7/8	7/9	8	8
(CA)6	2507097-2507108	6	6	6	6	6	6
(TA)8	3221617-3221632	9	NA	10	10	?	8/9
(TA)9	2844971-2844988	7	6	?	?	7	7
(TA)10	1744091-1744110	8/10	8/9	10/11	10/11	10	10
(AT)10	2951821-2951840	NA	NA	5	5	6	6
(AT)15	948935-948964	NA	NA	?	?	?	?
(AT)17	2597735-2597768	NA	NA	?	13	13	13
(ACC)5	1980049-1980063	5	5	5	5	5	5
(GGT)5	2567251-2567265	4	4	4	4	4	4
(AGT)5a	1237528-1237542	5	5	5	5	5	5
(GTA)9	2583814-2583840	9	10	9/10	11	9	9
(CACCG)3	2562391-2562405	3	3	3	3	3	3
6-3a	1190341-1190358	3	3	3	3	3	3
6-3b	2302531-2302548	3	3	3	3	3	3
6-7	1816857-1816892	6	6	?	8	7	7
7-3	285076-285096	3	3	3	3	3	3
10-4	1139035-1139074	4	NA	4	4	4	4
12-5	1381661-1381723	5	5	5	5	5	5
15-3	2928131-2928175	3	3	3	3	3	3
27-5	687026-687160	5	5	5	5	5	5

VNTR: variable number tandem repeat; NA: not available because of low coverage at that locus; mixture of multiple alleles; loci where VNTR number varied between the first and second strain are highlighted.