S3 Table: List of 20 *M. leprae* genomes used to infer unique SNPs in recurrent cases

|  |  |  |
| --- | --- | --- |
| **Name** | **Genotype** | **Described in** |
| TN | 1A | 1 |
| Thai53 | 1A | 2 |
| S2 | 1B | 3 |
| S11 | 1D | 3 |
| 3077 | 2F | 3 |
| Refshale\_16 | 2F | 3 |
| SK8 | 2F | 3 |
| Jorgen\_625 | 3I | 3 |
| SK2 | 3I | 3 |
| NHDP98 | 3I | 4 |
| NHDP55 | 3I | 4 |
| I30 | 3I | 4 |
| NHDP63 | 3I | 2 |
| S9 | 3K | 3 |
| Kyoto-2 | 3K | 5 |
| S10 | 3K | 3 |
| S15 | 3L | 3 |
| S13 | 4N | 3 |
| S14 | 4O | 3 |
| Br4923 | 4P | 2 |
| Mx1-22 (*M. lepromatosis*) | Outgroup | 6 |

SNP: single nucleotide polymorphism.