

^b Clade designations according to SITVIT2 using revised SpoIDB4 rules; Unk: Unknown pattern within any of the major clades described in SITVIT2 database.

^a In this study, 129 distinct spoligotype patterns were obtained from 592 *M. tuberculosis* isolates; 523/592 (88.3%) were clustered (60 clusters, 2-155 strains/cluster), and 69 (11.7%) were unclustered. A total of 96 SITs containing 543 isolates matched a pre-existing shared type in SITVIT2, whereas 9 SITs (containing 25 isolates) were newly-created either within the present study or after a match with an orphan in the database (showed by an asterisk; SIT2836, SIT3051, SIT3072, SIT3073, SIT3074, SIT3075, SIT3076, SIT3077, SIT3078).

^b Clade designations according to SITVIT2 using revised SpoIDB4 rules; Unk: Unknown pattern within any of the major clades described in SITVIT2 database