

Suppl. Table S2. Mtb polymorphic sites associated with multidrug resistance.

1633538 PE_PGRS27	364	C > CACGGCCGCC, C > CANNNNNNNNN, C > CATG, C > CATGGCC, C > CATGGCCGCC, C > CATGGCGCCCC, and others AGC > A, A > AACCCCTTCG, A > AAGCAGAAG, A > ACG, A > AGCGC, A > AGG, A > AGTCAGCGG, A > C, and others	EMB, INH, PZA, RIF, SM
2534564 Rv2262c	330		AM, CM, MFX, OFX, PTO
2061433 PE_PGRS33	415	T > TCCGCCGGCG C > CAGGGTTGGA, C > CAGGGTTGTC, C > CCGAGTTGCCCG, C > CCGGCTTGCCCCG, C > CCGGCTTGGCGG,	EMB, INH, PZA, RIF, SM
968428 PE_PGRS15	606	CCGGGTTGG > CCGGGTTGGACGGGTTGG, and many others C > CCTGGCGGCC, C > CG, C > CGCCG, C > CGT, C > CGTCG, C > CGTCGCCGCCG, C > CGTCGG,	AM, MFX, OFX, PTO, SM
1632542 PE_PGRS27	696	C > CGTCGCCGCCG, C > CGTCGGCGCGTTGCCGCCGCCACCG, CGG > CG, and many others	EMB, INH, PZA, RIF, SM
2729611 Rv2434c	317	T > TCAGATC, T > TCGGA, T > TCGGATCGG, T > TCGTATCG, T > G	AM, CM, MFX, OFX, PTO

*Locations of polymorphic sites are given as in the reference genome *Mycobacterium tuberculosis* H37Rv (NC_000962.3)

^tAntibiotic abbreviations: amikacin (AM), capreomycin (CM), ethambutol (EMB), isoniazid (INH), moxifloxacin (MFX), ofloxacin (OFX), prothionamide (PTO), pyrazinamide (PZA), rifampicin (RIF), and streptomycin (SM)