

SUPPLEMENTARY TABLE S1. Resistance profiles and genotypes of investigated clinical isolates of *M. tuberculosis*

Strain	INH (0.1 µg/ml)			RIF (1.0 µg/ml)		ETH (2.5 µg/ml)	
	Phenotype	Genetic loci		Phenotype	Genetic loci	Phenotype	Genetic loci
		<i>katG</i>	<i>inhA</i>				
Isoniazid-resistant strains							
186133	R	wt	C-15T	S	wt	R	wt
186136	R	wt	C-15T	S	wt	R	wt (R245R)
179932	R	wt	C-15T	S	wt	R	wt
500090	R	S315T1	wt	S	wt	R	bp 637 and 638 del, stop at aa pos. 317
Multidrug-resistant strains (<i>katG</i> -mutation)							
295	R	S315T1	wt	R	S531L	R	L478R
130	R	S315T1	wt	R	S531L	R	bp 1054 del, stop at aa pos.372
177836	R	S315T1	wt	R	S531L	R	S266R & bp 1047 del, stop at aa pos. 372
186038	R	S315T1	wt	R	S531L	R	W256Stop
186137	R	S315T1	wt	R	S531L	R	bp 8 del, stop at aa pos. 62
176231	R	S315T1	wt	R	S531L	R	S266R & bp 1047 del, stop at aa pos. 372
186071	R	S315T1	wt	R	S531L	R	A-7G
186075	R	S315T1	wt	R	H526D	R	N298T
186033	R	S315T1	wt	R	S531L	R	S266R & bp 849 del, stop at aa pos. 313
186032	R	S315T1	wt	R	H526Y	R	L229R
186003	R	S315T1	wt	R	S531L	R	bp 107 del, stop at aa pos. 62
186014	R	S315T1	wt	R	S531L	R	M1R
Multidrug-resistant strains (<i>inhA</i> -promoter mutation)							
179	R	wt	C-15T	R	S531L & D516Y	R	wt
181783	R	wt	C-15T	R	S531L	R	P192S
186110	R	wt	C-15T	R	S531L	R	wt
176743	R	wt	C-15T	R	S531L	R	P192S
181976	R	wt	C-15T	R	S531L	R	P192S
Multidrug-resistant strains (<i>katG</i> -mutation and <i>inhA</i> -promoter mutation)							
1703	R	S315T1	T-8C	R	H526Y	R	F431L
186044	R	S315T1	C-15T	R	H526D	R	wt
186025	R	S315T1	C-15T	R	S531L	R	H281P
186031	R	S315T1	C-15T	R	S531L	R	S15P