

Supplementary File 6: Isoniazid resistance mutations

Isoniazid resistance mutations among the 161 strains that underwent successful genotyping and their associated degree of phenotypic resistance to isoniazid. No.- number.

Genotype	Total	Phenotype					
		Highly resistant		Resistant		Borderline	
		No.	Row %	No.	Row %	No.	Row %
ahpC_C-72T	1	1	100.0	0	0.0	0	0.0
fabG1_C-15T	79	63	79.7	10	12.7	6	7.6
inhA_I194T/fabG1_C-15T	1	1	100.0	0	0.0	0	0.0
inhA_I21T/fabG1_C-15T	1	1	100.0	0	0.0	0	0.0
inhA_S94A/fabG1_C-15T	1	1	100.0	0	0.0	0	0.0
katG_S315T/N	72	72	100.0	0	0.0	0	0.0
katG_S315T/fabG1_C-15T	5	5	100.0	0	0.0	0	0.0
katG_S315T/fabG1_G-17T	1	1	100.0	0	0.0	0	0.0
Total	161	145	90.1	10	6.2	6	3.7