

**Supplementary Table 1**

Strain Name	Genetic Lineage	Obtained from	MIC ( $\mu\text{g/ml}$ )		Presence/absence of Tap580 insertion & method of confirmation	Reference
			RIF	INH		
H37Rv	4	C Sasetti	0.15	0.08	PCR/Xhol digest	
		NIRT	0.4*	0.2*		
CDC1551	4	WR Bishai	0.075	0.08	Genome sequencing and PCR/Xhol digest	
M1338	1	NIRT Clinical Isolate	0.4*	0.2*		[3]
NITR203	2 Beijing	NIRT Clinical Isolate	0.4*	>1.6*	Genome sequencing	[4]
SA161	2 Beijing	I Orme	0.15	0.16	Genome sequencing and PCR/Xhol digest	
M4100A	2 non-Beijing	S Gagneux	0.075	0.04	PCR/Xhol digest	[5,6]
SG1	3	S Gagneux	0.075	0.08	PCR/Xhol digest	[5]
GM1503	4	S Gagneux	0.08	0.03	PCR/Xhol digest	[5]

\*MIC determined at the National Institute for Research in Tuberculosis in Chennai, India