Supplementary Table S3: Geographical distribution of individual lineage 3 clonal complexes (CCs) in Asia, Africa, and Europe, BG=genetic background

UN Geo-region	MTBC L3-CCs						Total
	1	**2**	**3**	**4**	**5**	**L3-BG**	
Australia–New Zealand	35 (4.7%)	1 (0.3%)	92 (14.0%)	18 (10.4%)	4 (4.0%)	113 (17.0%)	**263**
Central Asia	3 (0.4%)	1 (0.3%)	18 (2.7%)	1 (0.6%)	0 (0.0%)	3 (0.5%)	**26**
Eastern Africa	188 (25.3%)	242 (70.8%)	19 (2.9%)	12 (6.9%)	41 (41.4%)	16 (2.4%)	**518**
Eastern Asia	11 (1.5%)	0 (0.0%)	1 (0.2%)	0 (0.0%)	2 (2.0%)	10 (1.5%)	**24**
Middle Africa	1 (0.1%)	3 (0.9%)	1 (0.2%)	1 (0.6%)	1 (1.0%)	0 (.0%)	**7**
Northern Africa	120 (16.0%)	7 (2.0%)	1 (0.2%)	0 (0.0%)	2 (2.0%)	2 (0.3%)	**132**
Northern America	53 (7.1%)	11 (3.2%)	140 (21.2%)	19 (11.0%)	4 (4.0%)	110 (16.5%)	**337**
Northern Europe	30 (4.2%)	23 (6.7%)	16 (2.4%)	8 (4.6%)	12 (12.1%)	13 (2.0%)	**102**
Southern Africa	0 (0.0%)	3 (0.9%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	**3**
Southern Asia	79 (10.6%)	2 (0.6%)	216 (32.8%)	75 (43.4%)	6 (6.1%)	259 (38.9%)	**637**
Southern Europe	21 (2.8%)	4 (1.2%)	25 (3.8%)	9 (5.2%)	6 (6.1%)	9 (1.4%)	**74**
Western Africa	6 (0.8%)	1 (0.3%)	5 (0.8%)	0 (0.0%)	2 (2.0%)	21 (3.2%)	**35**
Western Asia	87 (11.7%)	2 (0.6%)	48 (7.3%)	0 (0.0%)	1 (1.0%)	56 (8.4%)	**194**
Western Europe	109 (14.7%)	42 (12.3%)	77 (11.7%)	30 (17.3%)	18 (18.2%)	54 (8.1%)	**330**
Total	**743**	**342**	**659**	**173**	**99**	**666**	**2,682**