Supplemental Table 1 Drug susceptibility patterns of 183 clinical *M. tuberculosis* for whole-genome sequencing

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| Susceptibility or resistance | Number of strains |
| Fully susceptible\* | 46 |
| MDR except XDR | 113 |
| HR | 10 |
| HRE | 12 |
| HRO | 3 |
| HRS | 31 |
| HREC | 2 |
| HREO | 6 |
| HRSE | 20 |
| HRSO | 10 |
| HRSCK | 1 |
| HRSEC | 1 |
| HRSEO | 17 |
| Over all XDR | 24 |
| HRKO | 1 |
| HRSCO | 2 |
| HRSKO | 1 |
| HRSECO | 1 |
| HRSEKO | 5 |
| HRECKO | 3 |
| HRSCKO | 1 |
| HRSECKO | 10 |

Note, \* isolates susceptible to isoniazid, rifampicin, ethambutol, streptomycin, kanamycin, ofloxacin, and capreomycin, simultaneously; H, isoniazid; R, rifampicin; S, streptomycin; E, ethambutol; K, kanamycin; O, ofloxacin; C, capreomycin.