



Supplementary Figure S6. Deeplex Myc-TB results obtained using 10^2 versus 10^4 *M. tuberculosis* genome copies of sample TB-TDR0014 from the TDR collection. A red arrow indicates an erratically detected false positive synonymous low frequency variant (5.2%) *prnCA* Y34Y at 10^2 genomes. This variant was detected only in this experiment. This likely reflects very sporadic emergence over filtered noise level of low frequency amplification errors, sometimes seen with 10^1 or 10^2 genome copies.