Figure S1. Genes associated with *M. tuberculosis* PZA susceptibility by TnSeq. Libraries of $2 \times 10^5$ independent himar1 mutants (4-fold saturation) were plated on 7H9 agar without (top panels, 7H9) or with POA (middle panels, 7H9 + POA). Genomic DNA was extracted, processed and sequenced as described in Minato, *et al* 2019 *mSystems*. Bottom panels show all TA dinucleotides of the region that is illustrated. Mean read count comparisons from two independent replicates are shown for *pstS3C2A1* (a) *pckA* (b), *aceE* (c), and *subl-cysTWA1* (d).