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ydjF predicted DNA-binding transcriptional regulator ydhV predicted oxidoreductase yejF YejA/YejB/YejE/YejF ABC transporter gfcC conserved protein ygjR predicted NAD(P)-binding dehydrogenase rffA dTDP-4-oxo-6-deoxy-D-glucose transaminase yhdR fkpA peptidyl-prolyl cis-trans isomerase; in protein folding recN protein used in recombination and DNA repair yneK predicted protein yejE YejA/YejB/YejE/YejF ABC transporter yfgC predicted peptidase wxxE lipid III flippase b3245 pykF pyruvate kinase I monomer mdoG periplasmic glucan (MDO) biosynthesis protein acrR AcrR transcriptional repressor rcsC RcsC-P his mdoH membrane glycosyltransferase; synthesis of membrane-derived oligosaccharide (MDO) cutA copper binding protein CutA yrbE YrbF/YrbE ABC transporter greA transcription elongation factor GreA flhC FlnDC transcriptional dual regulator cysE serine acetyltransferase monomer rsxD integral membrane protein of SoxR-reducing complex aceE subunit of E1p component of pyruvate dehydrogenase complex ppc phosphoenolpyruvate carboxylase sbmA predicted transporter yrbB predicted NTP-binding protein serA alpha;-ketoglutarate reductase / D-3-phosphoglycerate dehydrogenase yaiW predicted DNA-binding transcriptional regulator transporter ptsH HPr
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Figure S8. Loci whose disruption was significant in bleomycin.

Key

Z-score

Yellow (blue) indicates that transposon insertions in or near a gene were beneficial (deleterious). Z-scores were calculated as described in *Materials and Methods*.