



Figure S1. Clustal W 2.0.8 multiple sequence alignment of DacB from *E. coli* K12 and *P. aeruginosa* PAO1. The sequences corresponding to the signal peptides are shown in italics. SXXK, SXN, and KTG motifs are shown within boxes and the delimitations of the three domains are indicated with arrows. Asteriks, colons, and periods indicate identical, conserved, and semiconserved residues, respectively.