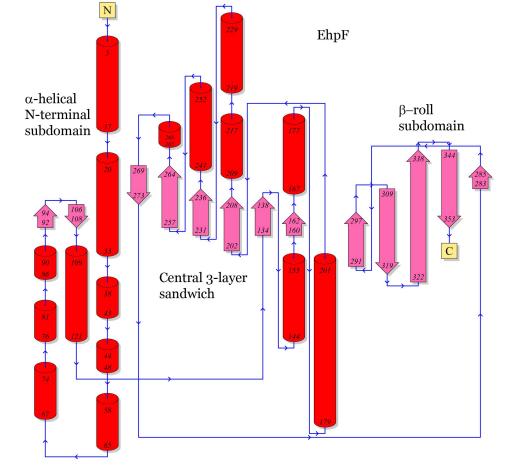


EhpF 361 VIEGVY GriC 347 IIEGVY



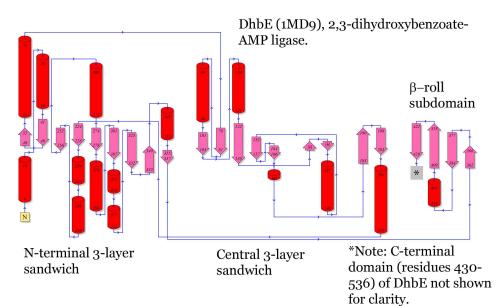


Figure S1. Alignment illustrating the sequence similarity between EhpF and GriC from S. griseus. The structure of GriC is not currently known but it is believed to participate in the conversion of 2-amino-3-hydroxybenzoic acid to 2-amino-3hydroxybenzaldehyde in the grixazone biosynthetic pathway.

Figure S2. Annotated topological diagrams for EhpF and DhbE illustrating the similarities and differences between the folds of EhpF and a typical ANL superfamily member. The figure was generated using the PDBSum server (www.ebi.ac.uk/pdbsum).