

Figure S1. Fluorescence microscope images of SF8300 bacteria stained with vancomycin-bodipy conjugate in presence or absence of antibacterial agent.

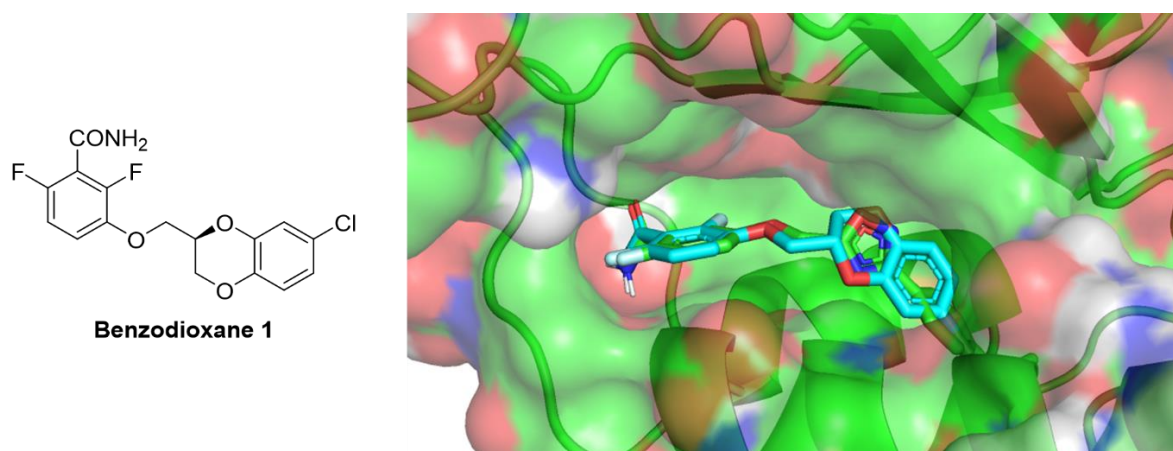
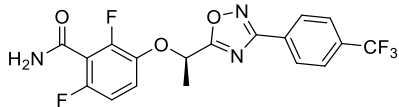
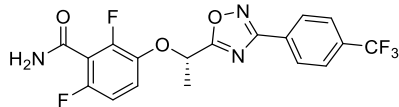
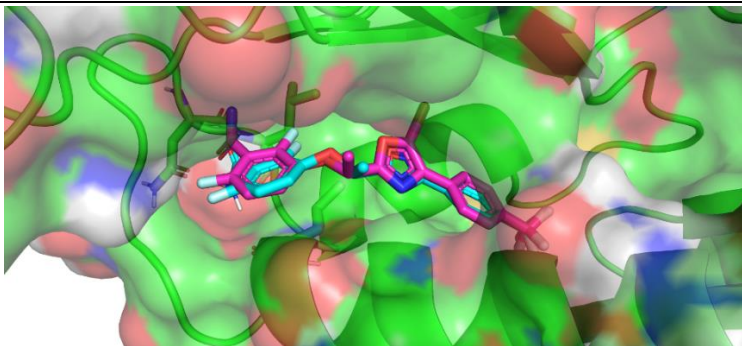
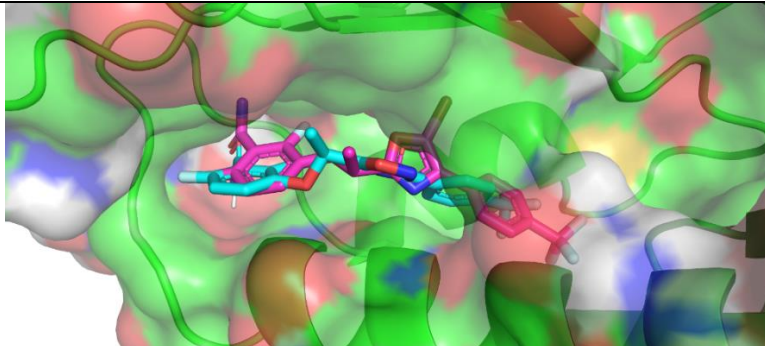


Figure S2: Superimposition of docking results of compound **II.c** (green) and **benzodioxane 1** (cyan) within the allosteric binding site of FtsZ.

Compound	V.b	
Enantiomer	(R)	(S)
Structure		
Docking and superimposition with co-crystallized inhibitor		

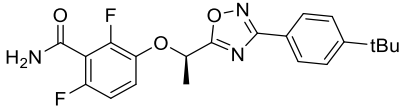
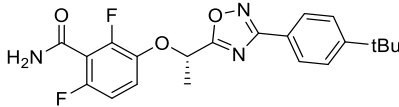
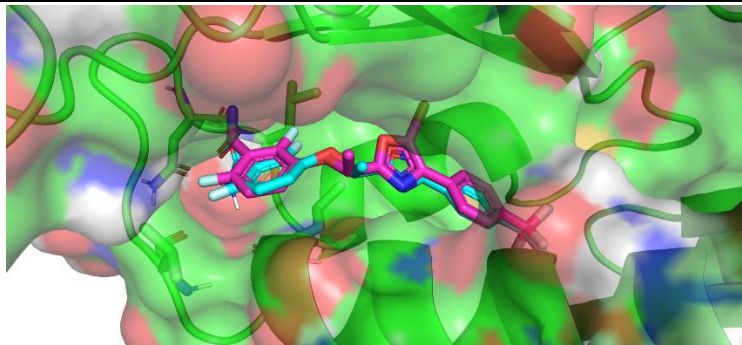
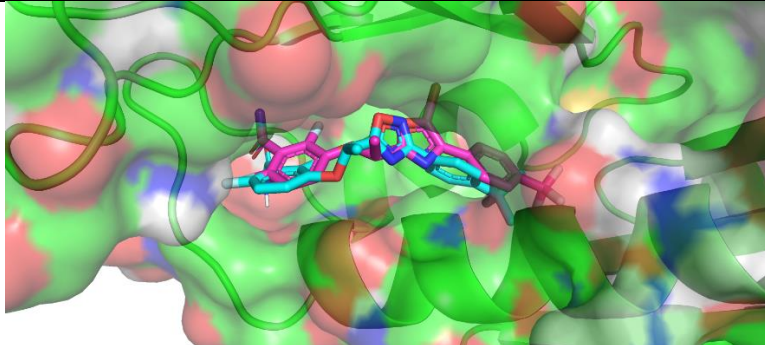
Compound	V.c	
Enantiomer	(R)	(S)
Structure		
Docking and superimposition with co-crystallized inhibitor		

Figure S3: Docking of both enantiomers of compound **V.b** and **V.c** within the allosteric binding site of FtsZ (PDB code: 6KVP) [6] . Docking results (cyan) are superimposed with the co-crystallized oxazole inhibitor (magenta).

