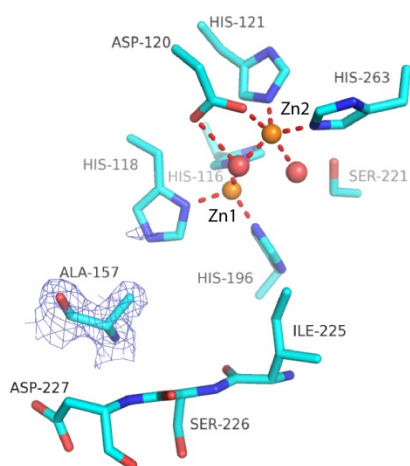


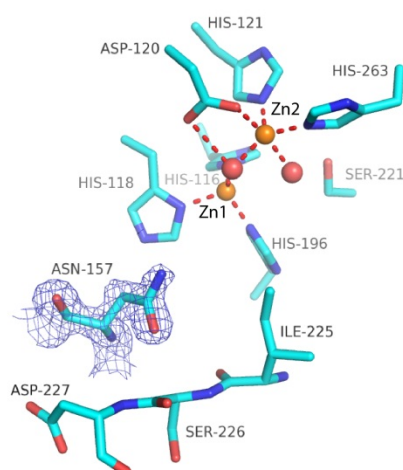
Supplementary to
Leiros et al., Crystal Structure of the Mobile Metallo- β -Lactamase AIM-1 from *Pseudomonas aeruginosa*: Insights into Antibiotic Binding and the role of Gln157.

Figure S1 2Fo-Fc electron density map at 1.2 σ for the a) AIM-1-Q157A and b) AIM-1-Q157N crystal structures. c) The AIM-1-Q157A (cyan) and d) the AIM-1-Q157N (cyan) structures with docked hydrolysed cefoxitin (magenta) and the AIM-1 (white) structure (Q157) superimposed.

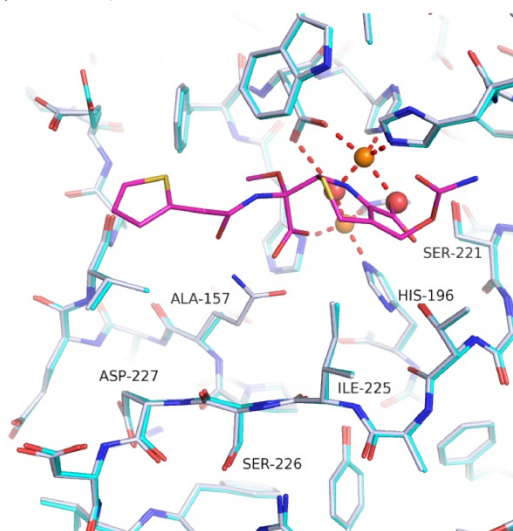
a) AIM-1-Q157A



b) AIM-1-Q157N



c) AIM-1-Q157A



d) AIM-1-Q157N

