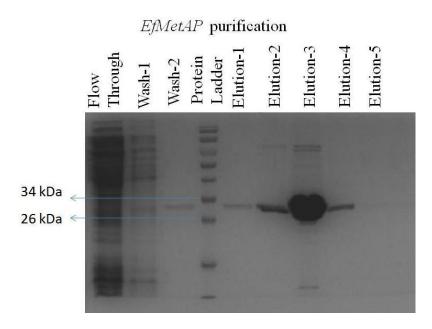
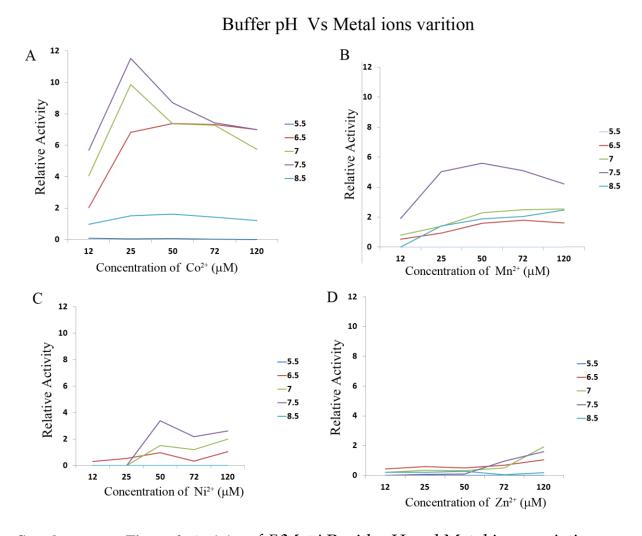
Supplementary material:

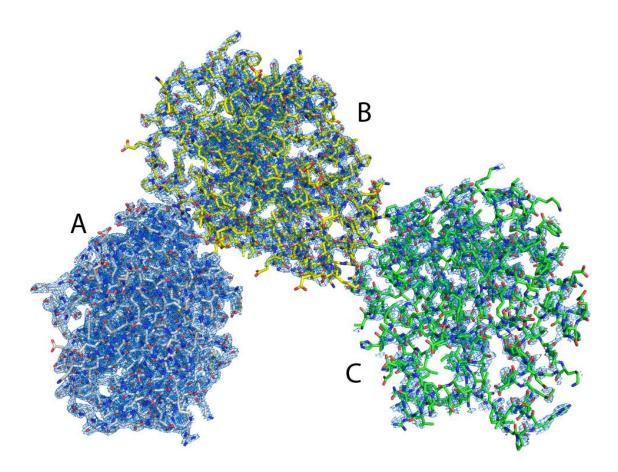
Purification of *Ef***MetAP**: C-terminus penta-his tagged 30 KDa *Ef*MetAP was expressed and purified from BL21 (DE3) *E. coli* strain. More than 95% pure protein was obtained from the single nickel affinity column judged by coomassie blue-stained SDS-polyacrylamide gel.



Supplementary Figure 1. Purification of PolyHis-tagged *Ef*MetAP (~30 kDa).



Supplementary Figure 2. Activity of *Ef*MetAP with pH and Metal ions variation

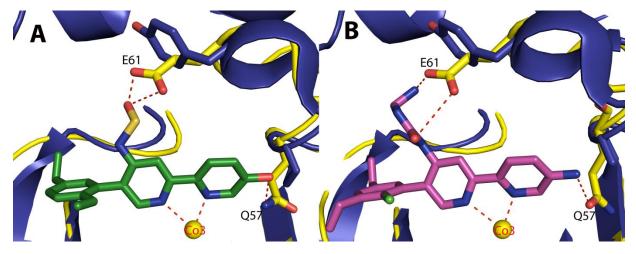


Supplementary Figure 3. Crystal structure of *Ef*MetAP at 2.3 Å with three molecules in the asymmetric unit (2Fo-2Fc electron density contoured at 1.5 σ). Molecule A, B and C represented by white, yellow and green sticks respectively. Good density was observed for all residues in molecules A and B while parts of the molecule C could not be traced.

Virtual Screening

Virtual screening of 126 bipyridine derivatives was carried against *Ef*metAP and *Hs*MetAp to find out microbes specific inhibitor. Two molecules have good hit against *Ef*MetAP.

Compoun d Id	R1	R2	R3	R4	Binding energy with EfMetAP (kcal/mol)	Binding energy with HsMetAP (kcal/mol)
1	ОН	Н	HN		-8.0	-7.2
2	NH2	Н	H ₂ N HN NH ₂	CI	-8.1	-7.0



Supplementary Figure 4. possible binding mode of bipyridine derivatives with *Ef*MetAP. Yellow cartoon represent *Ef*MetAP, while blue cartoon represents *Hs*MetAP. (A) Interaction of compound 1 and (B) compound 2 with *Ef*MetAP.

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