

Article

Structural evidence of active site adaptability towards different sized substrates of aromatic amino acid aminotransferase from *Psychrobacter* sp. B6

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For both subunits A and B of *PsyArAT*/FOH dimer (Figure S1A,B) volume differences are not big but noticeable (Table 1). In *PsyArAT*/YOH volume of the active pocket of monomer A (Figure S1C) is bigger than in monomer B (Figure S1D). For the *PsyArAT*/WOH differences between active site volume are the most visible. Only one monomer of each dimer bound the ligand. The volume of active pocket of monomers A and C with bound inhibitor is much smaller (Figure S1E) in comparison to empty subunits (monomers B and D) (Figure S1F). *PsyArAT*/DOH crystallized as monomer in asymmetric unit and its active site (Figure S1G) is the smallest, especially comparing to the native *PsyArAT* (PDB ID: 4RKC) (Figure S1H).

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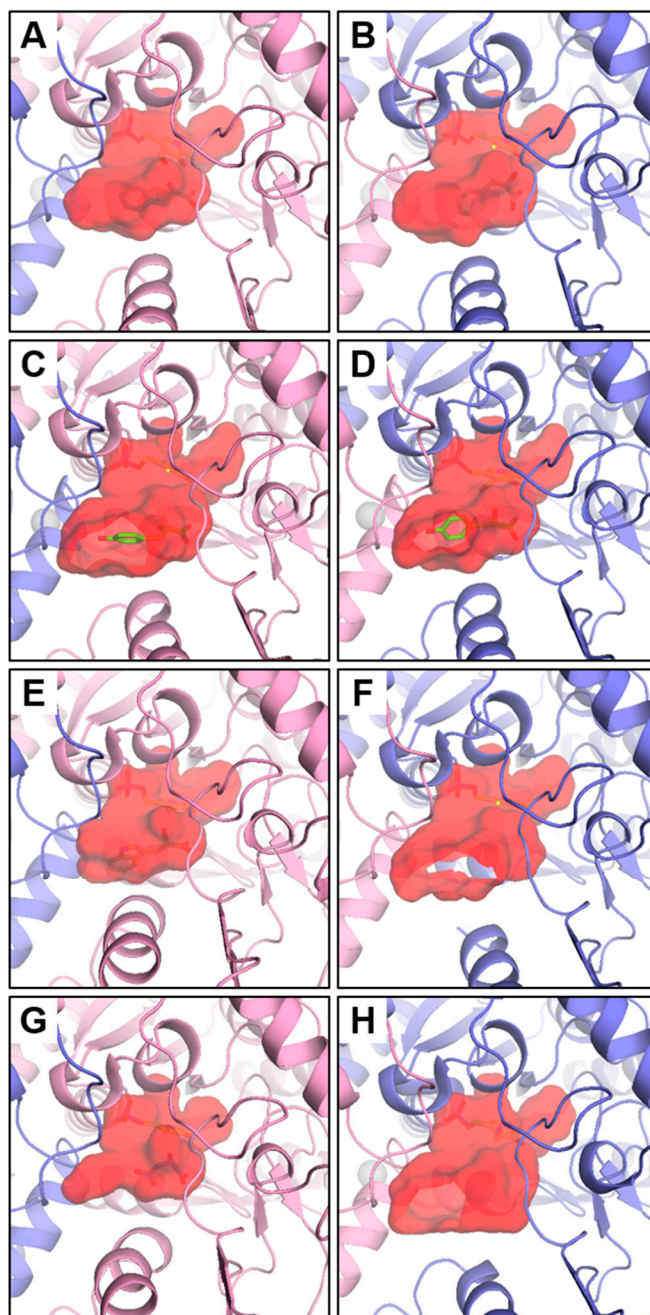


Figure S1. Graphical volume representations of *PsyArAT* active sites. (A, B) *PsyArAT*/FOH monomer A and B, respectively; (C, D) *PsyArAT*/YOH monomer A and B, respectively; (E, F) *PsyArAT*/WOH - monomer A and B, respectively; (G) *PsyArAT*/DOH monomer; (H) *PsyArAT* (PDB ID: 4RKC) monomer B.