



Figure S5. Comparison of the ochratoxinase-like enzymes based on the individual analysis of electrostatic surfaces in the atomic models. Hierarchical clustering analysis of the electrostatic distance matrix for the selected group of carboxylic ester hydrolases structures was calculated by the webPIPSA algorithm and the Poisson-Boltzman equation. The enzymes are identified by their producing fungal strain, where the ochratoxin-producing species are depicted in red color.