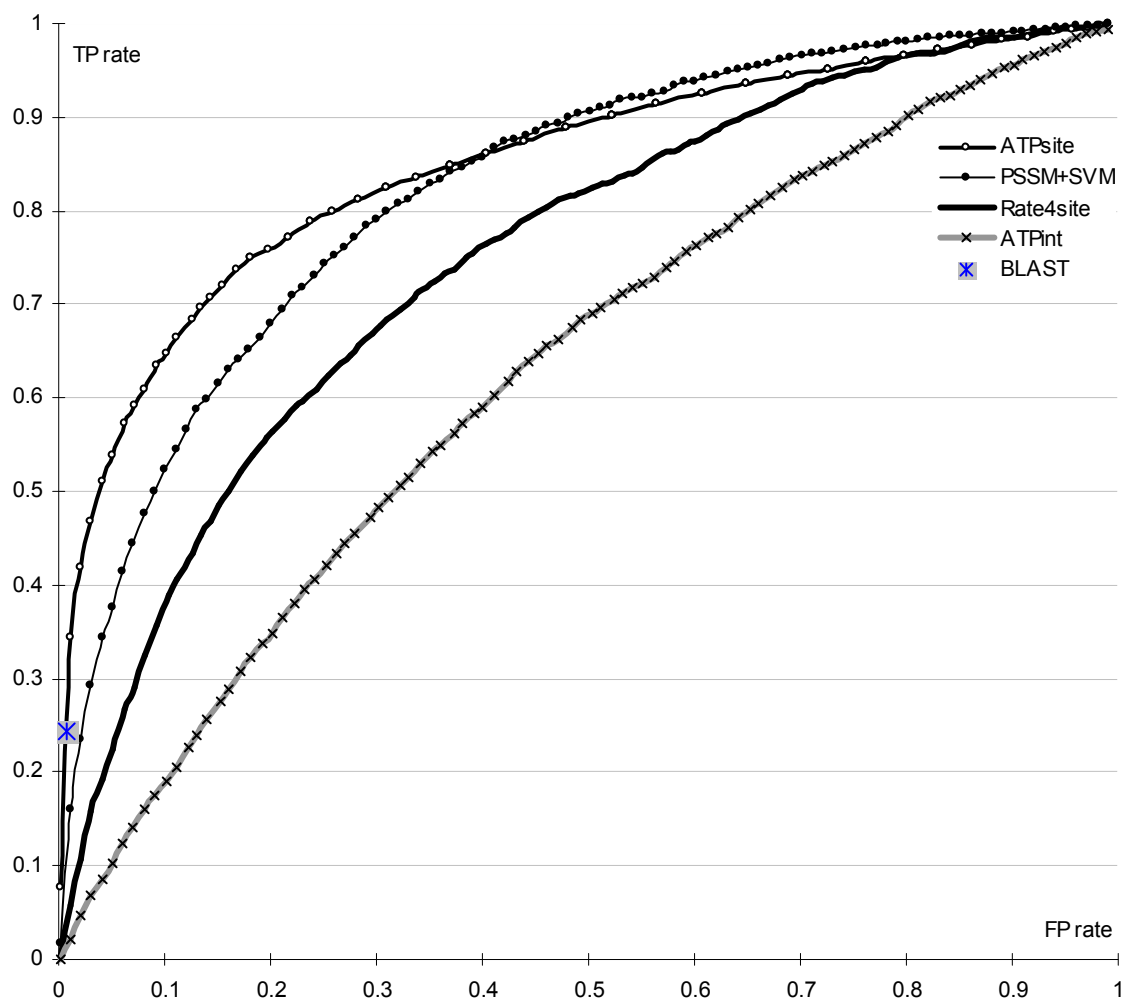
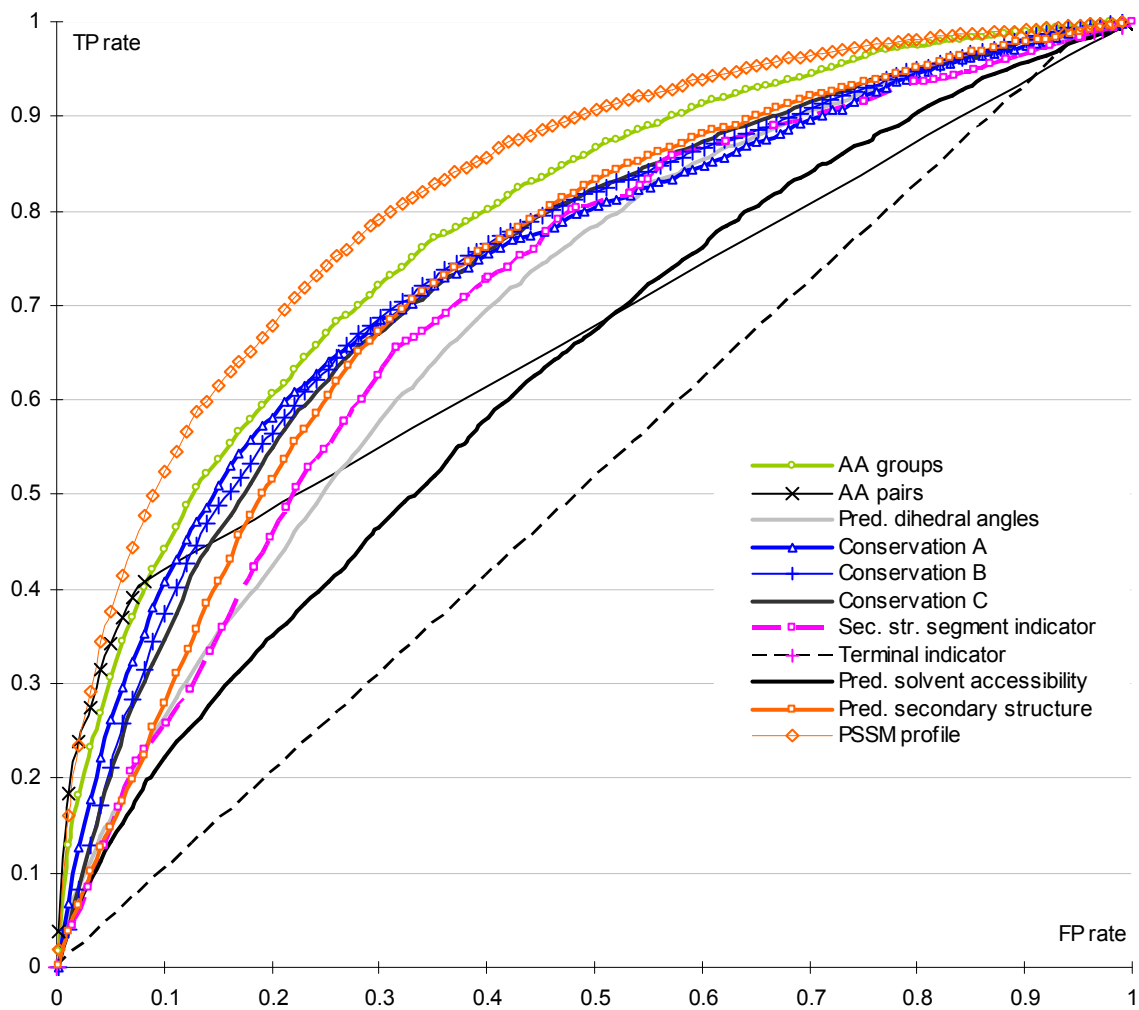


Supplement for “ATPsite: sequence-based prediction of ATP-binding residues”

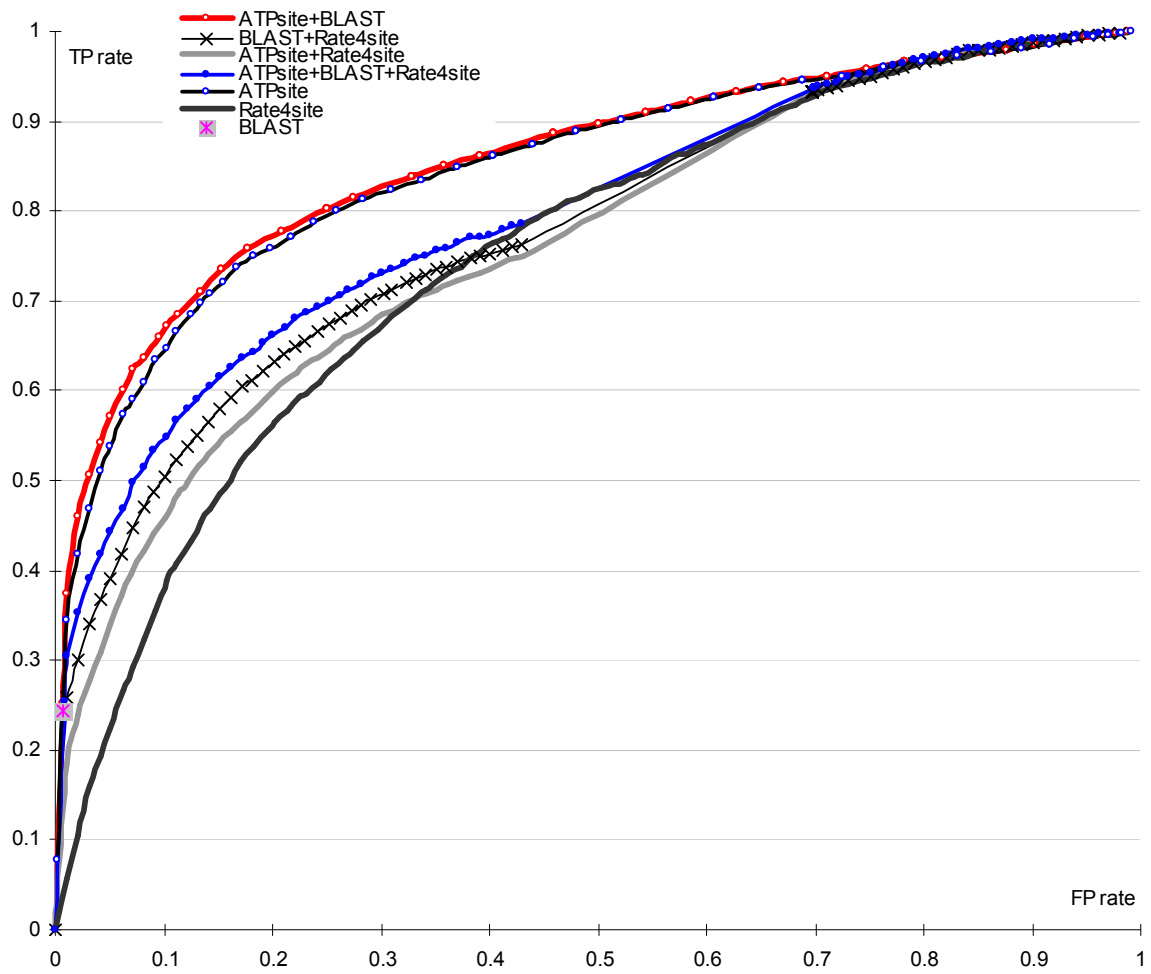
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Supplementary Figure 1. ROC curves for ATPsite, ATPint, Rate4site and the predictor based on PSSM with SVM classifier. The BLAST-based solution is shown using a single point that corresponds to binary predictions.



Supplementary Figure 2. ROC curves calculated based on predictions generated using individual input types.



Supplementary Figure 3. ROC curves of the ensemble predictors.