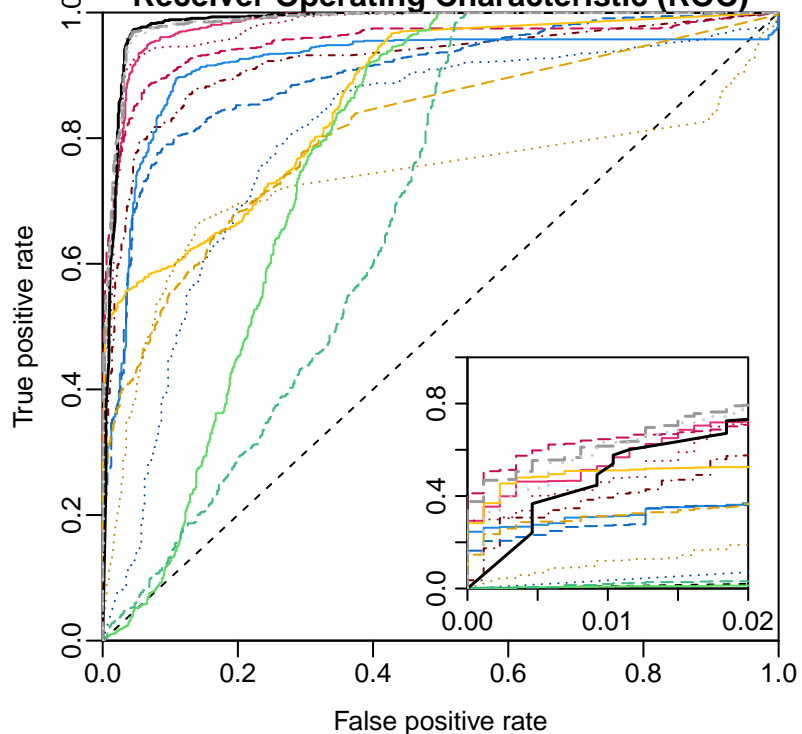


Receiver Operating Characteristic (ROC)



	AUROC	AUPRC
Phylogenetic Profiling		
— P/A Jaccard	0.981	0.978
- - G/L Distance	0.952	0.964
..... P/A Overlap	0.975	0.973
- . . . G/L MI	0.922	0.934
Phylogenetic Structure		
— RP ContextTree	0.916	0.926
- - RP MirrorTree	0.901	0.909
..... Tree Distance	0.801	0.870
Gene Organization		
— Gene Distance	0.866	0.883
- - Orientation MI	0.811	0.842
..... Moran's I	0.731	0.786
Sequence Level		
— Sequence Info	0.774	0.658
- - Gene Vector	0.687	0.606
Ensemble Methods		
— Random Forest	0.984	0.978
- - Logistic Regression	0.986	0.984
..... Neural Network	0.985	0.982
Other		
- - - - Random Guessing	0.500	0.500

Precision–Recall Curve (PRC)

