



Phylogenetic Profiling		AUROC	AUPRC
—	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
- - -	Neural Network	0.985	0.982
...	Logistic Regression	0.986	0.984
Other			
- - - -	Random Guessing	0.500	0.500

