



Component Algorithms

- G/L MI
- G/L Distance
- RP MirrorTree
- P/A Conservation
- P/A Jaccard
- RP ContextTree
- Gene Vector
- Sequence Info
- · - Gene Distance
- · - Moran's I
- · - Orientation MI
- · - Tree Distance

AUROC's (CORUM)

0.774
0.713
0.701
0.691
0.688
0.685
0.585
0.522
0.504
0.498
0.497
0.491

Ensemble Models

- Logistic Regression
- Random Forest
- Neural Network
- - Logistic Regression (Transfer)
- - Random Forest (Transfer)
- - Neural Network (Transfer)

AUROC's (CORUM)

0.819
0.838
0.857
0.695
0.681
0.688

AUROC's (KEGG)

0.952
0.955
0.942
0.841
0.786
0.795