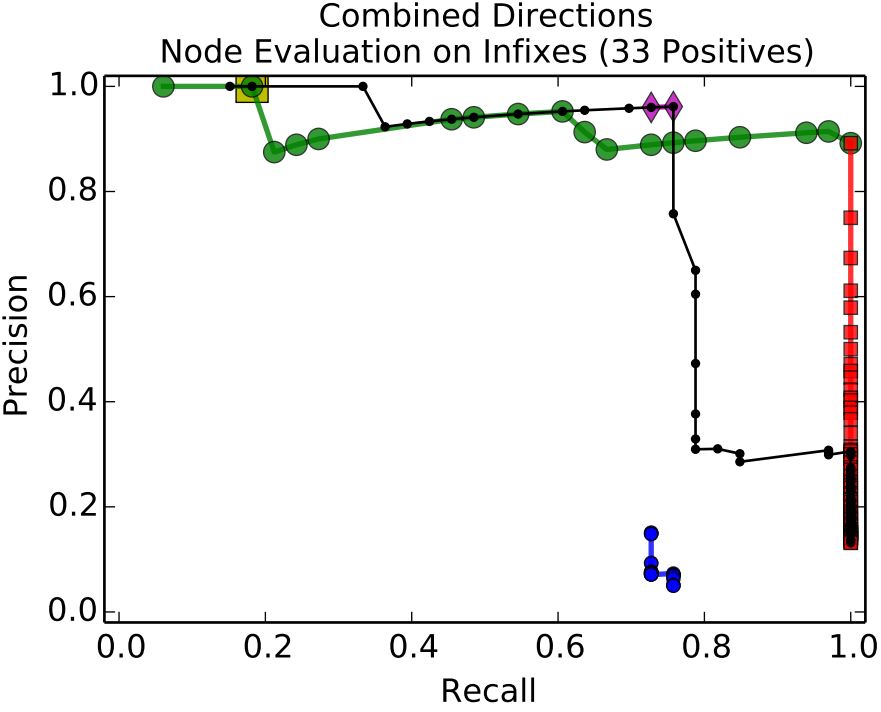
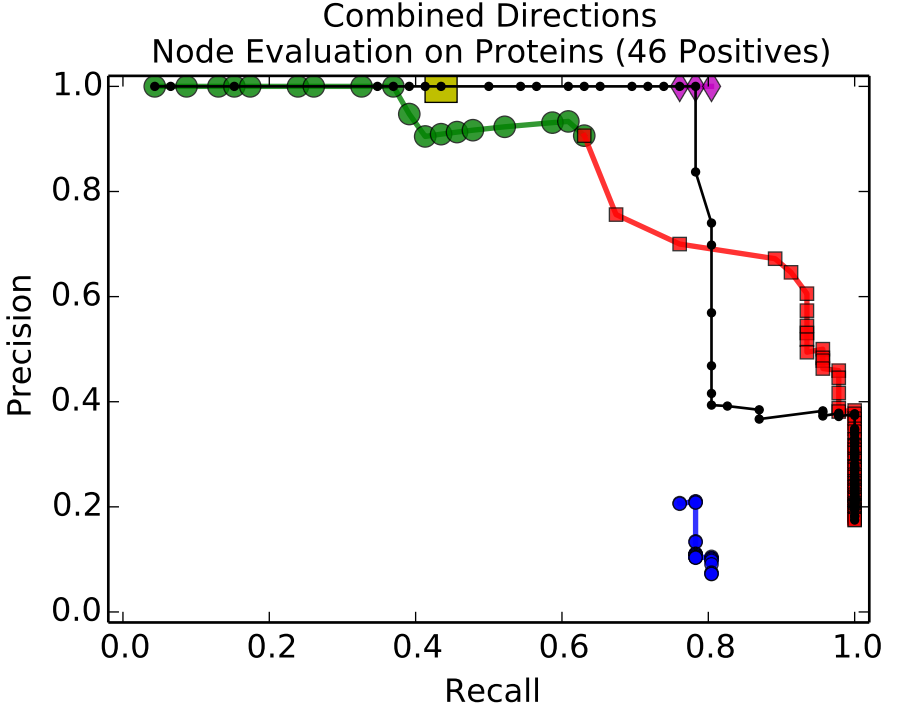
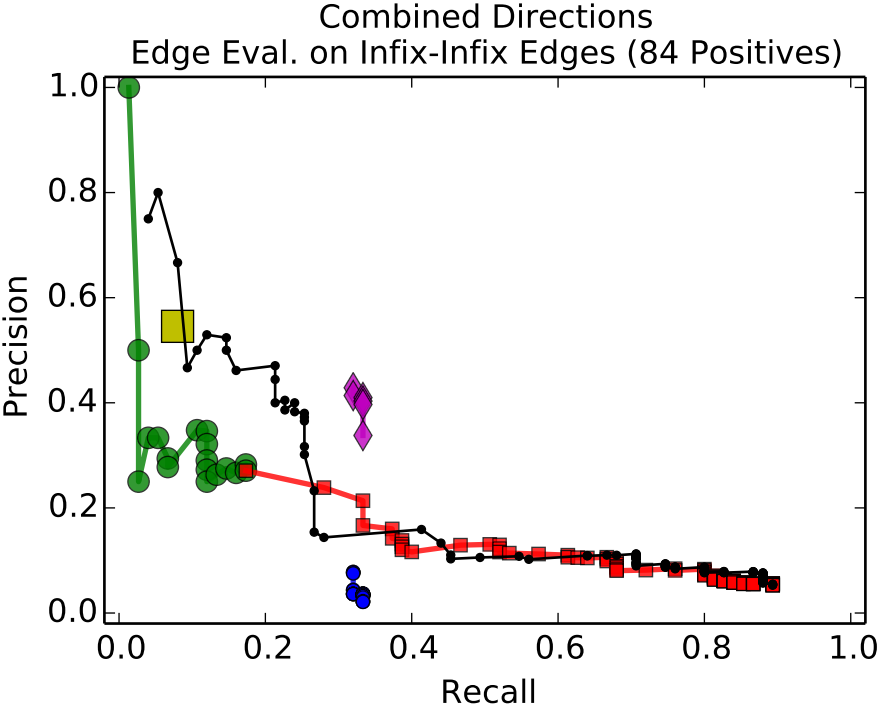
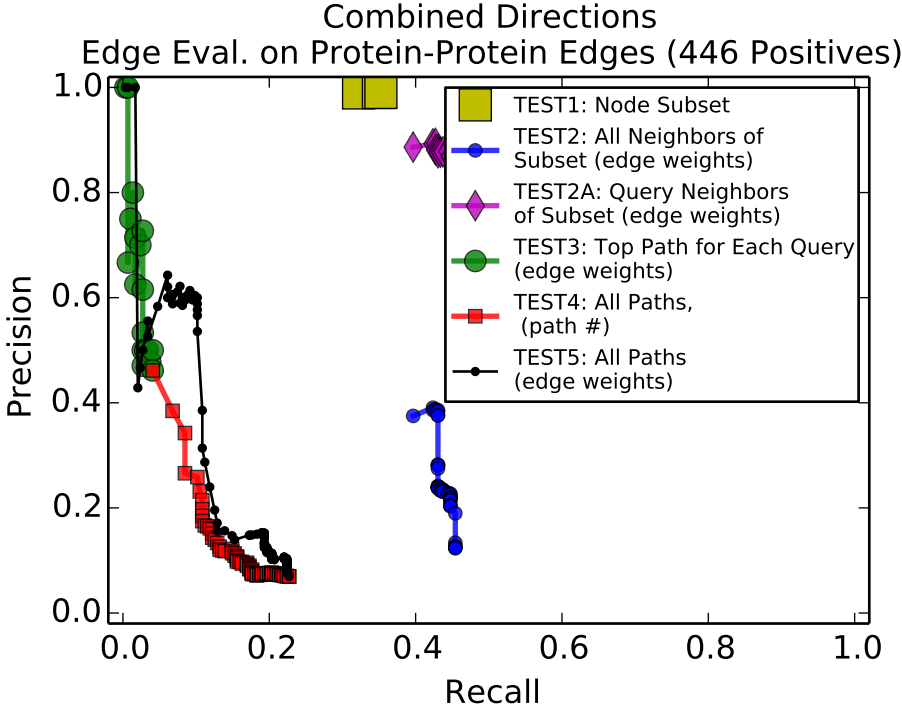

Combined Directions
Node Evaluation on Infixes (33 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Combined Directions
Node Evaluation on Proteins (46 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Combined Directions
Edge Eval. on Infix-Infix Edges (84 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Combined Directions
Edge Eval. on Protein-Protein Edges (446 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0
TEST1: Node Subset
TEST2: All Neighbors of Subset (edge weights)
TEST2A: Query Neighbors of Subset (edge weights)
TEST3: Top Path for Each Query (edge weights)
TEST4: All Paths, (path #)
TEST5: All Paths (edge weights)