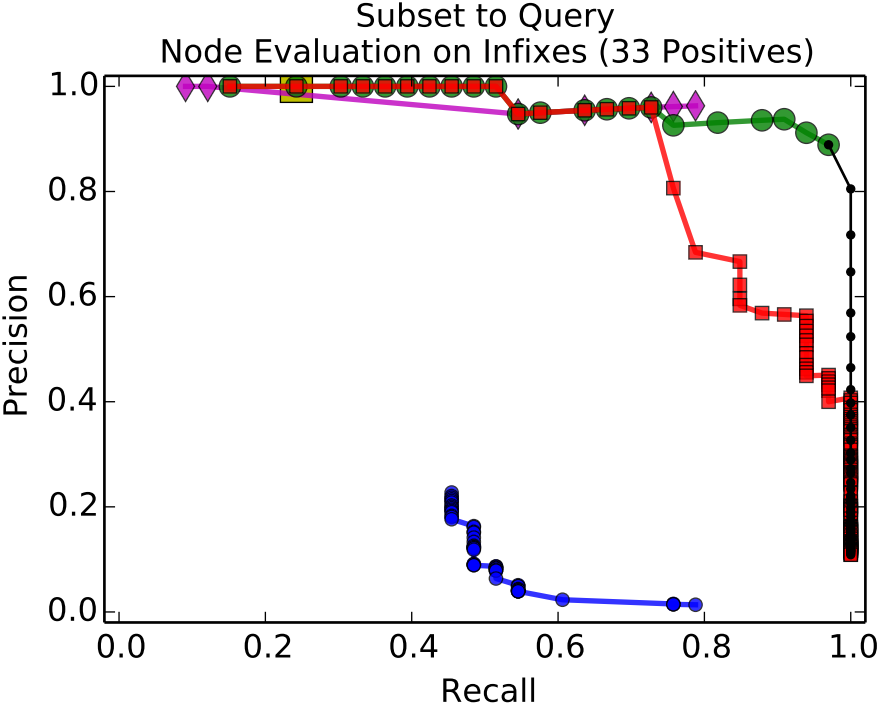
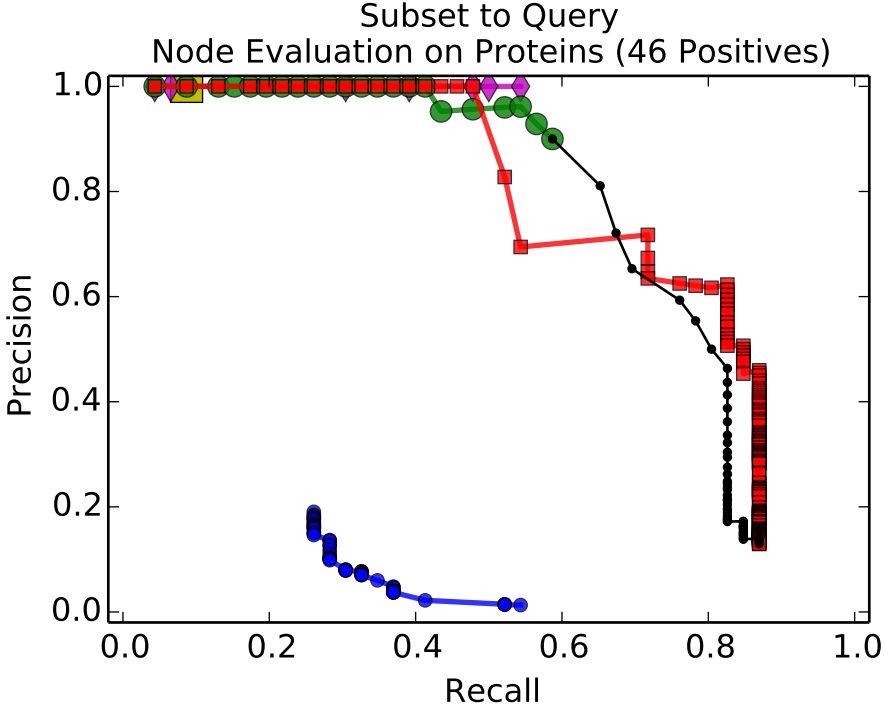
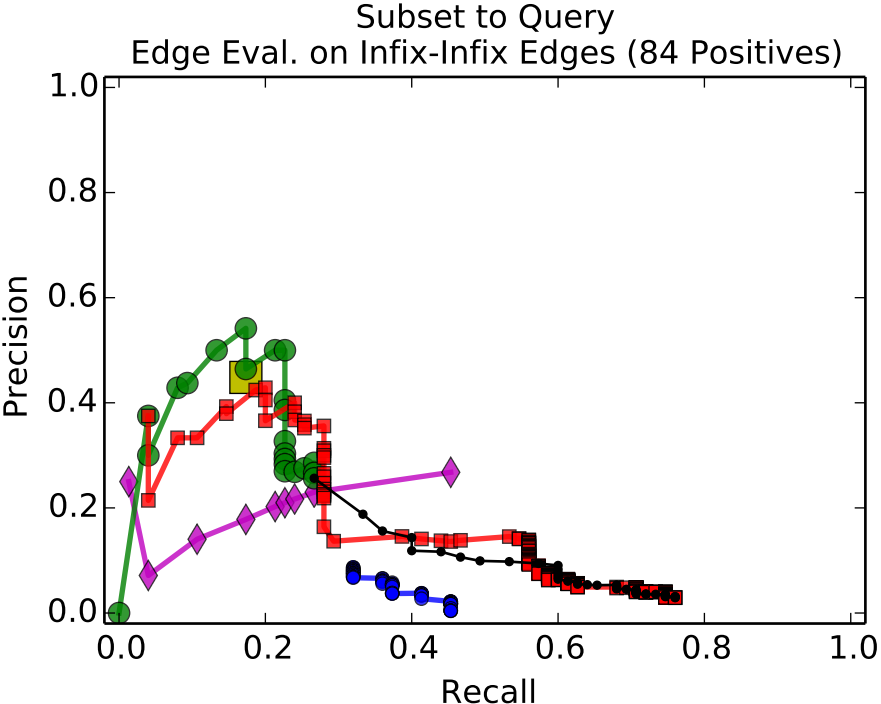
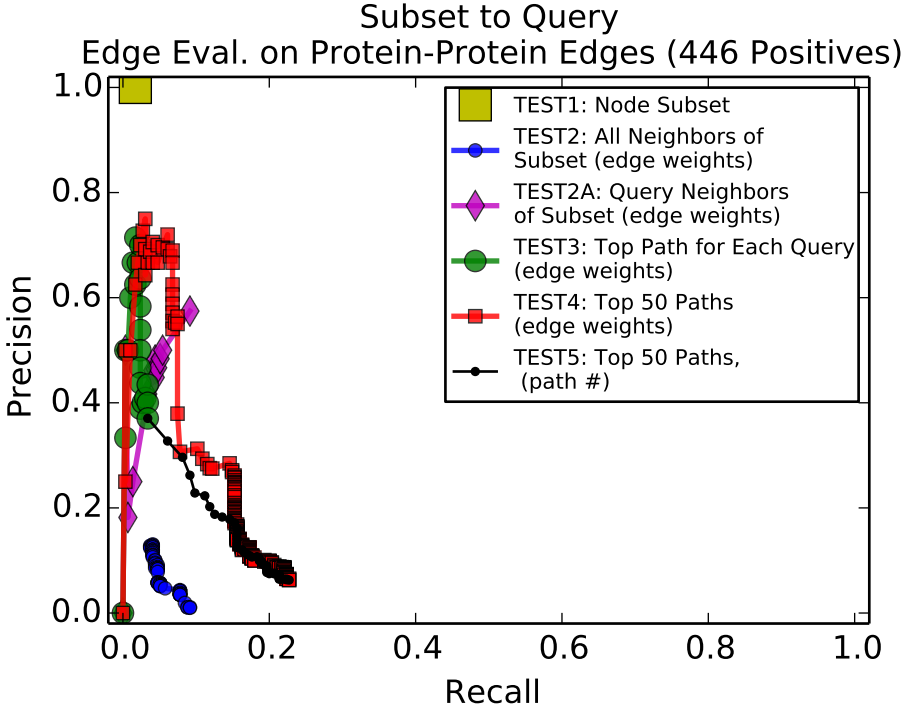

Subset to Query
Node Evaluation on Infixes (33 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Subset to Query
Node Evaluation on Proteins (46 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Subset to Query
Edge Eval. on Infix-Infix Edges (84 Positives)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0


Subset to Query
Edge Eval. on Protein-Protein Edges (446 Positives)
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: Top 50 Paths (edge weights)
TEST5: Top 50 Paths, (path #)
Precision
Recall
0.0
0.2
0.4
0.6
0.8
1.0