

# SUPPLEMENTARY MATERIAL

## OWL-NETS: Abstracting Knowledge for Network Inference

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### 1. Link Prediction Algorithms

The eight local similarity algorithms, defined consistent with the literature,<sup>1,2</sup> are:

- (1) Degree Product: Given two nodes  $i$  and  $j$ , this measure is calculated as the product of the degree (i.e., the number of connected nodes) of nodes  $i$  and  $j$ , where  $k$  is the node degree:

$$score(i, j) = k_i k_j \quad (1)$$

- (2) Common Neighbors: Given two nodes  $i$  and  $j$ , this measure is calculated as the number of neighbors that are common to both nodes  $i$  and  $j$ , where  $\Gamma$  is the set of nodes connected to node  $j$ :

$$score(i, j) = |\Gamma(i) \cap \Gamma(j)| \quad (2)$$

- (3) Jaccard Coefficient:<sup>3</sup> Nodes  $i$  and  $j$ , this measure is calculated as the number of neighbors that are common to both nodes  $i$  and  $j$  normalized by the number of nodes adjacent to either node  $i$  or node  $j$ :

$$score(i, j) = \frac{|\Gamma(i) \cap \Gamma(j)|}{|\Gamma(i) \cup \Gamma(j)|} \quad (3)$$

- (4) Srenson Similarity:<sup>4</sup> Given nodes  $i$  and  $j$ , this measure is calculated as the number of neighbors that are common to both nodes  $i$  and  $j$  normalized by the sum of the degrees of node  $i$  and node  $j$ :

$$score(i, j) = \frac{|\Gamma(i) \cap \Gamma(j)|}{k_i + k_j} \quad (4)$$

- (5) Leicht-Holme-Newman:<sup>5</sup> Two nodes  $i$  and  $j$ , this measure is calculated as the number of neighbors that are common to both nodes  $i$  and  $j$  normalized by the product of the degrees of node  $i$  and node  $j$ :

$$score(i, j) = \frac{|\Gamma(i) \cap \Gamma(j)|}{k_i * k_j} \quad (5)$$

- (6) Shortest Paths: Given two nodes  $i$  and  $j$ , this measure is calculated as the reciprocal of the length of the shortest path from node  $i$  to node  $j$ ,  $\sigma(i, j)$ :

$$score(i, j) = \frac{1}{\sigma(i, j)} \quad (6)$$

A score of zero is given for all node pairs not connected by a path.

- (7) Resource Allocation:<sup>6</sup> given two nodes  $i$  and  $j$ , this measure is calculated as the sum of the reciprocal of the degrees of nodes adjacent to both nodes  $i$  and  $j$ :

$$score(i, j) = \sum_{z \in \Gamma(i) \cap \Gamma(j)} \frac{1}{k_z} \quad (7)$$

- (8) Adamic-Advar:<sup>7</sup> Given two nodes  $i$  and  $j$ , this measure is calculated as the sum of the reciprocal of the log of the degrees of the nodes adjacent to both nodes  $i$  and  $j$ :

$$score(i, j) = \sum_{z \in \Gamma(i) \cap \Gamma(j)} \frac{1}{\log(k_z)} \quad (8)$$

The two global similarity algorithms, defined consistent with the literature,<sup>8</sup> are:

- (1) Katz:<sup>9</sup> Given an unweighted adjacency matrix  $A$ ,  $A_{i,j}$  if there is a link between nodes  $i$  and  $j$  and zero if there is not. Each element of  $A_{i,j}$ ,  $A_k$  has value equal to the number of walks with length  $k$  between nodes  $i$  and  $j$ :

$$score(i, j) = \sum_{k=1}^{\infty} \beta^k A_{ij}^k \quad (9)$$

where  $\beta$ , must be lower than the largest eigenvector of matrix  $A$ , that is used to give shorter paths more weight. Consistent with the literature,<sup>10</sup> a value of  $\beta = 0.001$  was used.

- (2) Rooted Page Rank: A random walker starts from node  $i$  and randomly moves to a neighbor of node  $i$ . The walker then has a probability of  $1 - \alpha$  for teleporting back to node  $i$ . Consistent with the literature,<sup>10</sup> a value of  $\alpha = 0.15$  was used.

Table 1. Table 1. Descriptive Characteristics by Network Representation

Property	OWL	OWL-NETS	p-value
Nodes	1578.400 (155.850)	247.950 (22.741)	< 0.0001
Edges	4110.930 (445.103)	1130.100 (153.514)	< 0.0001
Average Degree	5.204 (0.091)	9.083 (0.473)	< 0.0001
Density	0.003 (0.000)	0.037 (0.002)	0.002
Diameter	10.000 (0.000)	5.880 (0.325)	< 0.0001
Clustering Coefficient	0.067 (0.005)	0.338 (0.013)	0.013
Degree Assortativity	-0.223 (0.001)	-0.122 (0.019)	0.019
Degree Heterogeneity	13.684 (1.542)	2.354 (0.185)	< 0.0001
Number of Shortest Paths	5694.170 (1054.645)	683.680 (116.755)	< 0.0001
Average Shortest Path Length	3.760 (0.039)	2.934 (0.048)	< 0.0001
Number of Cliques	3532.050 (376.901)	703.110 (95.554)	< 0.0001

Table 2. Table 2. Descriptive Characteristics by Network Representation and Query

Property	Q2:OWL	Q2:OWL-NETS	Q3:OWL	Q3:OWL-NETS
Nodes	840	59	22,679	1783
Edges	1426	59	33,848	3940
Average Degree	3.419	2.000	2.980	4.420
Density	0.0040	0.0344	0.0001	0.0020
Diameter	13	4	9	18 <sup>a</sup>
Degree Assortativity	-0.193	-0.630	-0.124	-0.308
Degree Heterogeneity	8.789	4.533	558.463	6.673
Number of Shortest Paths	2,683	202	91,483	8,986 <sup>a</sup>
Average Shortest Path Length	4.06	3.19	4.13	6.54 <sup>a</sup>

\*Query (Q). All descriptives were run on the undirected versions of the networks.

<sup>a</sup>Query 3:OWL-NETS statistics were derived on the largest connected component.

Table 3. Table 3. Queries 2 Link Prediction Run-Time

Algorithm	OWL	OWL-NETS
Degree Product	00:00:07	00:37:42
Shortest Path	00:00:10	01:49:08
Common Neighbors	00:00:07	00:31:47
Jaccard Coefficient	00:00:07	00:32:04
Sorenson Similarity	00:00:07	00:32:09
Leicht-Holme-Newman	00:00:07	00:32:09
Adamic Advar	00:00:08	00:32:51
Resource Allocation	00:00:07	00:32:36
Katz	00:00:30	08:12:51
Rooted Page Rank	00:01:00	12:16:21

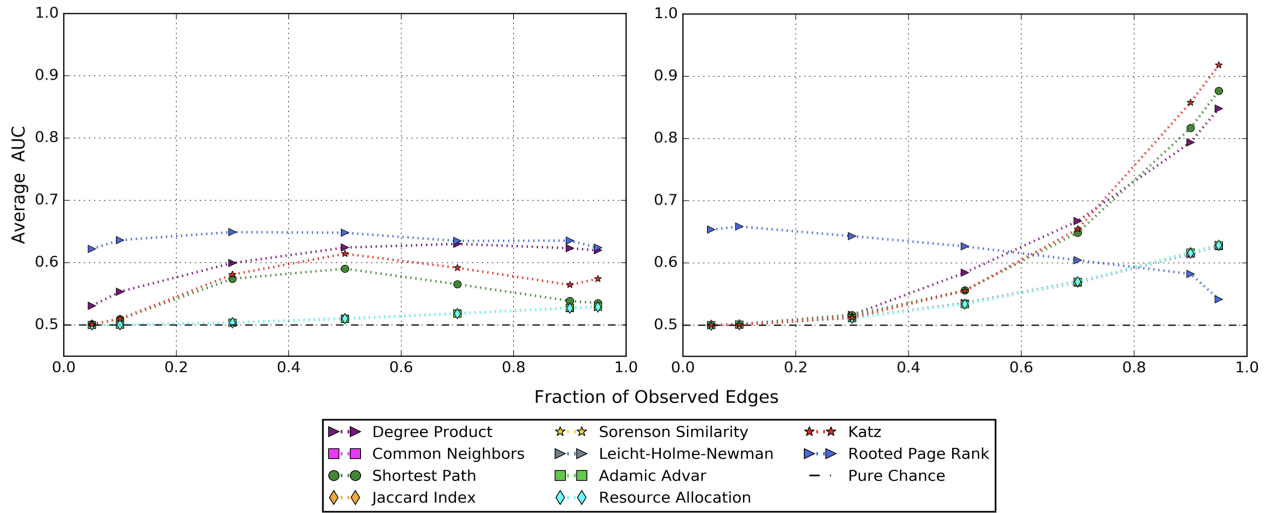


Fig. 1. Comparison of Link Prediction Methods by Network. (left) The original OWL representation network and (right) the OWL-NETS abstraction networks created from Query 2 (Table1).

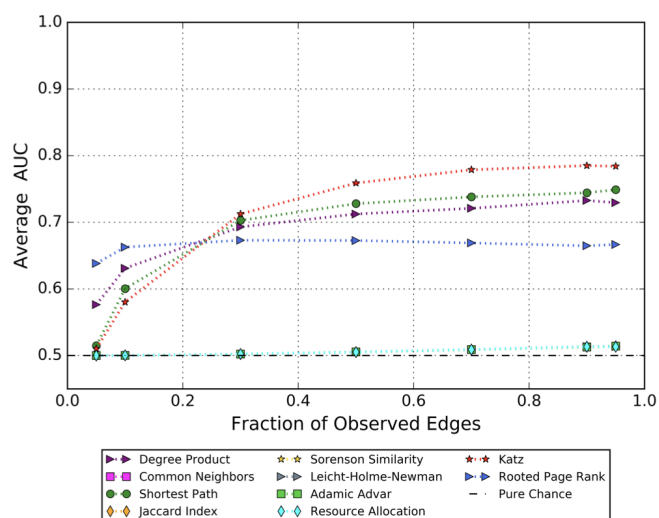


Fig. 2. Link Prediction Methods. The OWL-NETS abstraction networks created from Query 3 (Table1).

Table 4. Table 4. Top Scoring Edges from the Query 3 OWL-NETS Abstraction Networks (n=6 edges)

Node 1	Node 2	Description
AG-1067 <sup>a</sup>	MMP2 <sup>b</sup>	AGI-1067 is derived from probucol, which has been shown to decrease MMP-2 expression and activity in Apolipoprotein E-deficient mice. <sup>11</sup>
DB03683 <sup>a</sup>	APAF1 <sup>b</sup>	DB03683a targets MMP9 through an unknown mechanism. Downregulation of MMP9 induces APAF1 expression. <sup>12</sup>
celiprolol <sup>a</sup>	CYCS <sup>b</sup>	Celiprolol is an investigational drug used to treat hypertension. Cytochrome c has been shown to mediate hypertension in rats and in humans. <sup>13,14</sup>
1454838 <sup>c</sup>	TF <sup>b</sup>	TF binds to and transports iron. Iron is required for the proliferation of multiple myeloma cells. CD147 (1454838) is overexpressed in multiple myeloma cells and is positively correlated with cell proliferation. <sup>15,16</sup>
DB04513 <sup>a</sup>	RAF1 <sup>b</sup>	DB04513 targets Calmodulin 1, which can regulate the threshold for activation of the Ras/Raf/MEK/ERK signaling pathway. <sup>17</sup>
CXCL12 <sup>b</sup>	DB07691 <sup>a</sup>	DB07691 is an n-phenylbenzamide, which can inhibit the Mitochondrial Permeability Transition Pore whose continual opening is associated with mitochondrial dysfunction. CXCL12 regulates mitochondria association around the MTOC (microtubule organizing center). <sup>18,19</sup>

<sup>a</sup>DrugBank entity (DrugBank ID used for experimental compounds); <sup>b</sup>Uniprot entity (gene symbol shown ); <sup>c</sup>Reactome entity (database identifier).

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