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Abstract

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A network with *n* nodes contains $O(n^2)$ possible links. Even for networks of modest size, it is often difficult to evaluate all pairwise possibilities for links in a meaningful way. Hence, we propose an ensemble enabled approach to scaling up link prediction by decomposing traditional link prediction problems into subproblems of smaller size. These subproblems are each solved with the use of latent factor models, which can be effectively implemented over networks of modest size. Furthermore, the ensemble enabled approach has several advantages in terms of performance. Experiments on large networks demonstrate the effectiveness and scalability of our approach.

Structural Bagging Methods

Renjun Hu¹,

To scale up link prediction on very large networks, we develop three bagging methods:

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(1) Select an ensemble component N_s form G by one of the node, edge and biased edge bagging methods; (2) Construct a reduced adjacency matrix W_s from N_s ; (3) Apply NMF to W_s and use top-*k* method to predict.

If a node pair appears in multiple ensemble components, the maximum predicted value is considered.

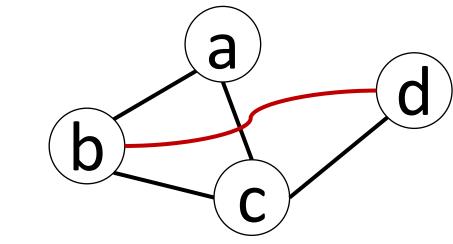
The O(n²) Problem in Link Prediction

| Network Sizes | 1 GHz | 3 GHz | 10 GHz | |
|-----------------------|--------------|-------------|-------------|--|
| 10 ⁶ nodes | 1000 sec. | 333 sec. | 100 sec. | |
| 10 ⁷ nodes | 27.8 hrs | 9.3 hrs | 2.78 hrs | |
| 10 ⁸ nodes | > 100 days | > 35 days | > 10 days | |
| 10 ⁹ nodes | > 10000 days | > 3500 days | > 1000 days | |

Table 1: Time required to allocate a single machine cycle to every node-pair possibility in networks.

Most current link prediction algorithms evaluate the link propensities only over a subset of possibilities rather than exhaustively search over the entire network [1, 2].

Bound of node bagging ensembles: The expected times of each node pair included in μ / f^2 ensemble components is at least μ , where f is the sampling fraction.



Scaling up Link Prediction with Ensembles

Figure 1: Triangle-closing model: The edge (*b*, *d*) is a triangle-closing edge. When the node c is selected, its neighbors a, b and d are also put into the same ensemble component.

Jinpeng Huai¹

Using link prediction characteristics: Since most of all new links in social networks span within very short distances [4], typically closing triangles shown in Figure 1, we design particular bagging approaches such that a node is always sampled together with all its neighbors, which guarantees the possibility of forming triangles.

Experimental Results

| AA · 🖬 · | NMF(Node) · • · | AA · 🖃 · | NMF(Node) · • · |
|-----------|-----------------|-----------|-----------------|
| BIGCLAM 😽 | NMF(Edge) 📥 | BIGCLAM 🛪 | NMF(Edge) 📥 |
| NMF 🔶 | NMF(Biased) + | NMF 🔶 | NMF(Biased) + |

Latent Factor Model for Scalable Link Prediction

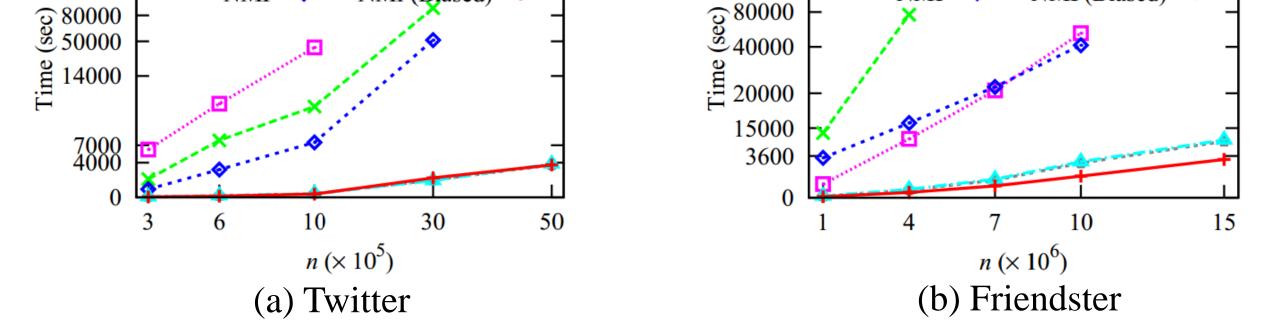
The link prediction ranking problem: Given a network G = (N, A) with node set N and edge set A, determine the top-k node-pair recommendations such that these node pairs are not included in A.

We make a latent factor model, i.e., nonnegative matrix factorization (NMF), practical for link prediction by factorizing the weight matrix W of G into FF^{T} , where F is a set of latent factors and can be determined by using the following multiplicative update rule [3]:

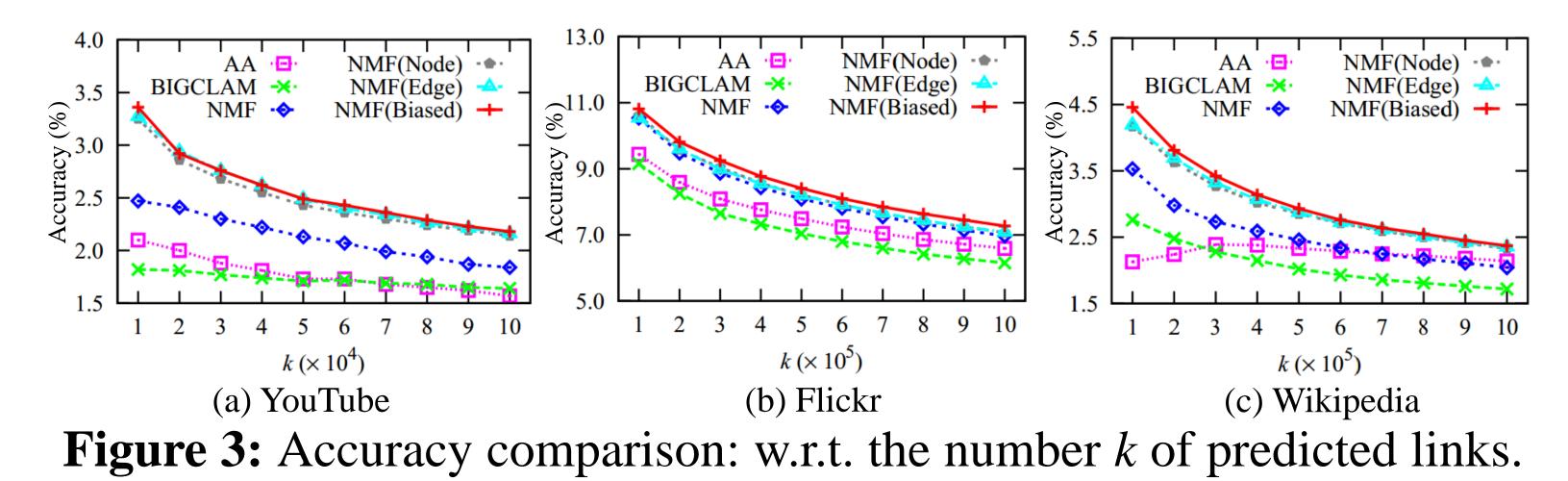
$$F_{ij} \leftarrow F_{ij} (1 - \beta + \beta \frac{(WF)_{ij}}{(FF^T F)_{ij}}), \beta \in (0, 1]$$

The positive values of entries in FF^{T} can be viewed as the predictions of noisy 0-entries in W.

Efficient top-k prediction searching: We design an ε approximate top-k method to speed up searching the k largest values in FF^{T} . The following nested loop is executed for each (say *p*-th) column of S:







| Dataset | Accuracy | Dataset | Speedup |
|-----------|----------|------------|---------|
| YouTube | 18% | Twitter | 20x |
| Flickr | 4% | | |
| Wikipedia | 16% | Friendster | 31x |

for each i = 1 to f_p do for each j = i + 1 to f_p , do *II* S is obtained by sorting the columns of *F* in a descending order if $S_{iD} \cdot S_{iD} < \varepsilon / r$ then break inner loop; else increase the score of node-pair (R_{ip} , R_{ip}) by an amount of $S_{ip} \cdot S_{ip}$; // R is an inverted index matrix end for end for

The complexity of NMF is $O(r \cdot (m + n \cdot r))$. For large sparse networks, the $O(nr^2)$ term might be the bottleneck.

Table 2: Summaries of the accuracy and efficiency improved by NMF(Biased) compared with NMF.

The ensemble-enabled approach not only increases the efficiency, but also improves the accuracy.

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