

Lecture 10: Supervised Random Walk

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10.1 Supervised Link Prediction Methods

1. Combining scores of different unsupervised methods
 - Computing Scores: $s_w(u, v) = w_{AA}AA(u, v) + w_{CN}CN(u, v) + b$
 - Learning Weights: SVM/Ranking loss on scores
2. Supervised Random Walk
3. Collaborative Filtering based approaches

10.2 Precursor: Random Walks on Graphs

10.2.1 Link Prediction with Random Walks

Consider a source node $s \in V$ in the Graph $G = (V, E)$. Let $D = (d_1, d_2, \dots, d_n)$ be the candidate neighbours (or destination nodes) of node s . We assume that s and d are likely to be connected in the future if:

- The structure around s and d is rich - their surrounding region has a lot of edges.
- There is some affinity between s and d

One way to judge how rich the structure is around s and d is using random walks. Let $count(s, d)$ be the number of times d is reached in a random walk starting at s . Let $p(s, d)$ be the probability of reaching d from a random walk starting at s . We define $score(s, d)$ as:

$$score(s, d) \propto count(s, d) \propto p(s, d)$$

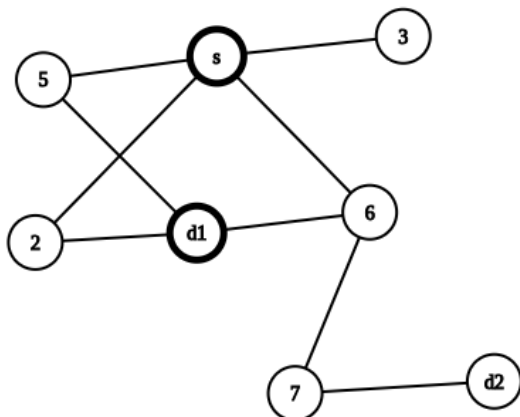
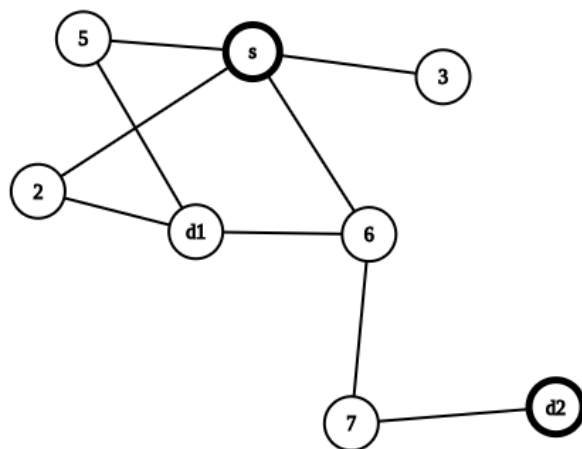
Once we obtain the scores for each $d_i \in D$, we can evaluate this approach for link prediction using MAP or Average Precision.

10.2.2 Constructing a Random Walk

Consider the graph $G = (V, E)$. Let A be the adjacency matrix of the graph. $A_{ij} = 1$ if vertex v_i is connected to v_j , and 0 otherwise. How to perform a random walk on this graph? To do this we define the transition matrix \mathbf{P} .

Transition Matrix: \mathbf{P} is said to be the transition matrix of a graph when $\forall i, j$ P_{ij} denotes the probability that v_i follows v_j in a random walk on the graph. Since you can only move to neighbouring nodes, $(A_{ij} = 0) \implies (P_{ij} = 0)$. P is a stochastic matrix - the sum of entries in each row is 1.

Once we have this matrix \mathbf{P} , we can use it to compute the probability of reaching one node from the other.

Figure 10.1: Edge (s, d_1) - High probabilityFigure 10.2: Edge (s, d_2) - Low probability

Using these probabilities, we can compute $score(s, d)$.

Traditional Random Walks assume you are equally likely to visit each one of your neighbours, and choose P in such a way that this condition is satisfied.

Supervised Random Walks get around this assumption. Using both node features, and the structure of the graph, the transition matrix P is learnt.

10.2.3 Computing Scores: PageRank

Given a transition matrix, also called weighted adjacency matrix, \mathbf{P} , and nodes s and d , how to compute $score(s, d)$? Let us analyse \mathbf{P} a little closely. Let $P(j|i) = P_{ij}$ be the probability of the moving to node j from the node i in a random walk.

$$P_i = [P(1|i) \quad P(2|i) \quad \dots \quad P(n|i)]$$

\mathbf{P} is a stochastic matrix, having sum of each row as 1. Therefore,

$$\mathbf{P}^M \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = \mathbf{P} \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix}$$

From the equation above, we can see that $\lambda(\mathbf{P}) = 1$.

Given a starting state s , let $p_t = [p_{1t} \quad \dots \quad p_{nt}]$ where p_{it} is the probability of being at node i after t steps.

$$p_t^T \mathbf{P} = p_{t+1}^T$$

Stationary Distribution: As $t \rightarrow \infty$, the probability of being at any node p_∞^T will be:

$$p_\infty^T = \lim_{t \rightarrow \infty} (p_0^T \mathbf{P}^t)$$

In most cases, this limit exists, and $p_\infty = p_{st}$, the steady state probability distribution on the graph. Clearly, $p_\infty^T = p_{st}^T \mathbf{P}$. This implies p_{st} is the left eigenvector of the eigenvalue $\lambda = 1$ of \mathbf{P} .

This steady state distribution, p_{st} is called the *PageRank* of the graph.

We have defined $score(s, d)$ to be proportional to the probability of reaching a node d in a random walk. Hence we define:

$$score(s, d) = p_{st}[d]$$

10.2.4 Better Metric: Personalised PageRank

An interesting observation is that $score(s, d) = p_{st}[d]$ is independent of the start node s . Therefore, for the purpose of link prediction, simply using PageRank is not enough to make user-specific predictions. We correct for this by introducing personalised page rank.

The idea is to introduce a restart probability, with which, a node can jump back to the source vertex s . This biases the random walks to the starting node s , and hence more localized, or personalized, compared to PageRank.

$$P_{ij}^* = (1 - \alpha)P_{ij} + \alpha I(j = s)$$

where α is the restart probability.

Hence,

$$P_s^* = (1 - \alpha)P + \alpha R_s$$

where only the entries in s^{th} column of R_s are 1, and all others are 0.

Note that

$$\mathbf{P}^* \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = (1 - \alpha)\mathbf{P} \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} + \alpha R_s \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = (1 - \alpha) \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} + \alpha \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix}$$

Similar to the above derivation, we can derive p_s^T to be the solution of

$$p^T \mathbf{P}_s^* = p^T$$

10.3 Supervised Random Walk

10.3.1 Modelling the Transition Matrix

One way to model the transition matrix P with the parameter w is as follows:

$$P_{ij} = \frac{w^T f(x_i, x_j)}{\sum_{j'} w^T f(x_i, x_{j'})}$$

where x_i, x_j are the node features of nodes i and j respectively.

Here, f should be a non-negative function, for example

$f(x, y) = \|x - y\|$ (Norm function)

$f(x, y) = x \odot y$ (Element-wise dot product)

The requirement of a non negative function can be bypassed using a hinge as follows:

$$P_{ij} = \frac{[w^T f(x_i, x_j)]_+}{\sum_{j'} [w^T f(x_i, x_{j'})]_+}$$

Incorporating constraints:

For a fixed node s , we choose w as per the following objective

$$\min ||w||^2$$

such that $\forall l, d \ (s, l) \notin E, (s, d) \in E \quad p_s^T[d] = P_\infty(d|s) > P_\infty(l|s) = p_s^T[l]$

i.e. the min norm solution of w such that the probabilities, as computed by Personalized PageRank using parameter w , for neighbours of s are more than non-neighbours.

Hence, for a node s , we have $N(s) \times (|V| - N(s))$ constraints.

Note that, the optimal w can be calculated either individually for all source nodes s , or a common w can be calculated across all nodes.

In practice, the above optimization can be realised using hinge loss and some form of regularization on w . i.e.

$$\min \lambda ||w||^2 + \sum_{l, d \ (s, l) \notin E, (s, d) \in E} [p_s^T[l] - p_s^T[d]]_+$$

10.3.2 Recommended reading

Supervised Random Walks: Predicting and Recommending Links in Social Networks[1]

References

- [1] BACKSTROM, L., AND LESKOVEC, J. Supervised random walks: Predicting and recommending links in social networks. In *Proceedings of the Fourth ACM International Conference on Web Search and Data Mining* (New York, NY, USA, 2011), WSDM '11, Association for Computing Machinery, p. 635–644.