Michael Lan Lingfei Zeng Xunjie Zhu

Stage2 - The Design Stage.

#### • Design Description:

The user is given 3 different similarity measures for searching. We will begin with the first similarity measure, which allows users to find tweets clusters related to search term inputs. (Search phrase and retrieve tweets with similar phrases acc to edit distance)

#### ♦ 1) Tweet clustering based on semantic similarity

We aim to cluster tweets into different groups using Hierarchical Clustering. To do this, we require semantic similarity scores. We compute these using the extended **Edit Distance** algorithm integrated **Word2Vec** embedding to compute the distance between sentences in the tweets, and by grouping by shortest distance we divide tweets into different groups.

In details, we describe tweet sentence  $S_1$  as  $S_1 = \{a_1, a_2, \ldots, a_n\}$  and sentence  $S_2$  as  $S_2 = \{b_1, b_2, \ldots, b_m\}$ .  $S_1$  consists of n words and  $S_2$  consists of m words.  $a_i$  is the i th word of  $S_1$  and  $b_j$  is the j th word of  $S_2$ . We use  $\text{Sim}(S_1, S_2)$  to describe the similarity between  $S_1$  and  $S_2$ , with the range of 0 (no relation) to 1 (semantic equivalence). For edit distance, we utilize Levenshtein Distance algorithm to extend to the word sets in sentences. We define the Levenshtein distance between  $S_1$  and  $S_2$  is L(n,m), where  $0 \le i \le n$  and  $0 \le j \le n$ . And we have:

$$L(i, j) = max(i, j)$$
 if  $min(i, j) = 0$ ,

$$L(i,j) = min \begin{cases} L(i-1,j-1) + c(a_i,b_j) \\ L(i,j-1) + 1 \\ L(i-1,j) + 1 \end{cases} otherwise.$$

Where the function  $c(a_i,b_i)$  is defined as

$$c(a_i, b_j) = \begin{cases} 0 & (a_i = b_j) \\ 1 & (a_i \neq b_j) \end{cases}$$

We also need to consider the circumstances that if two different words have the same or similar meaning (e.g., "cat" and "kitty"), in which case edit distance defines them as a mismatch. We address this issue by introducing Word2vec to measure semantic similarity between words. Word2vec [1] is an unsupervised deep learning algorithm that maps each word to a vector  $\in \mathbb{R}^n$  such that the semantically similar

words are mapped to the vectors that are close to each other in the geometry space. In our settings, given any two words, e.g.,  $a_i$  and  $b_j$  in the above, we first apply word2vec to map them to two vectors,  $\vec{v}_1$  and  $\vec{v}_2$ , and then use the euclidean distance,  $||\vec{v}_1 - \vec{v}_2||$ , to measure the cost of the replacement. We may normalize the cost into a relatively small range.

$$c(a_i, b_j) = \|\vec{v}_1 - \vec{v}_2\|$$

$$where \quad \vec{v}_1 = word2vec(a_i), \vec{v}_2 = word2vec(b_j)$$

## 2a) Recommending tweets and hashtags based on network similarity

Next, we will describe searching by network similarity. Users can input either a tweet or a hashtag to find similar tweets and hashtags from the stored data. If the data does not contain those search terms, the user can choose to 'Gather New Tweets' based on the search terms.

A directed bipartite network G with two different types of nodes, tweets and hashtags, will be created. Instead of storing nodes as strings, we give a unique ID to each node. A tweet will have an outgoing edge to a hashtag if it contains it. The graph  $G^2$  has as its nodes every pair of tweets and every pair of hashtags in G. As explained below, we will prune these node pairs to get an array called computed\_pairs. In Algorithm 3, we apply SimRank on computed\_pair to find a similarity score for every pair of nodes which are not 'far apart', a subjective measure.

SimRank, shown in Algorithm 3, is calculated using these equations: Let A and B be tweets. O(v) are the outgoing neighbors of v. The similarity s(A,B) is:

$$s(A, B) = \frac{C_1}{|O(A)||O(B)|} \sum_{i=1}^{|O(A)|} \sum_{j=1}^{|O(B)|} s(O_i(A), O_j(B))$$

C is a constant between 0 and 1. It is the decay factor and is the only adjustable parameter of the algorithm; we will set it as 0.8.

Let c and be be hashtags. I(v) are the ingoing neighbors of v. The similarity s(c,d) is:

$$s(c,d) = \frac{C_2}{|I(c)||I(d)|} \sum_{i=1}^{|I(c)|} \sum_{i=1}^{|I(d)|} s(I_i(c), I_j(d))$$

The similarity between two nodes is the average similarity of their neighbor. Initially, for node x, all node pairs (x,x) have score 1, while other node pairs have score 0. Next, the similarity scores are calculated according to the equations above. This step is iterated until the updated scores do not differ much from the previous iteration. Since s(a,b) and s(b,a) are symmetric, so only need to store one of their scores. More information can be found in the original paper, the powerpoints, or by contacting this report's authors.

For memory efficiency, pairs of nodes which are far away are pruned out. Let M be the adjacency matrix of G. If entry (n,m) in  $M^k$  is nonzero, then there exists a path between n and m of length k. Let Kmax be a subjective upper bound; we will set it as 4. Consider:

$$Z = \sum_{k=1}^{Kmax} M^{2k}$$

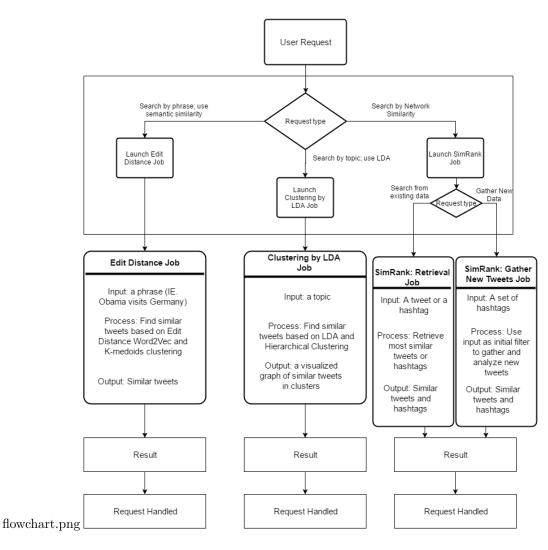
Only even powers of M are included in Z because G is bipartite. Then we will only compute similarity scores on the nonzero entries of Z, which are gathered into computed\_pairs.

## 2b)Time and Space Complexity for pruning node pairs and running SimRank

The bulk of the running time depends on the matrix multiplication used to obtain computed\_pairs; this takes  $\Theta(n^3)$ . The second pruning approach, computing the shortest path for all pairs, takes  $O(n \log n)$  for each pair, and since there are  $n^2$  pairs, its total running time would be  $O(n^3 \log n)$ . Thus, the adjacency matrix multiplication approach is preferred.

Now let d = the average of |I(a)| \* |I(b)| over all nodes in  $G^2$ . Let K = average number of iterations the algorithm performs. Let w be the number of pairs in computed\_pairs . Using a 'naive' implementation of SimRank, average running time of Algorithm 3 is  $O(K^*w^*d)$ . Its worst case mainly depends on K. We store node\_scores in a hash table, so its space complexity is O(n).

# Flow Diagram For User Requests



# • High Level Pseudo Code

```
Algorithm 1: Edit Distance
                    input: A sentence S_1 with n words as
                                                                              S_1 = \{a_1, a_2, \dots, a_n\} and a sentence S_2
                  with m words S_2 = \{b_1, b_2, \dots, b_m\}

output: The edit distance between S_1 and S_2
                                                                              L(n,m)
       \mathbf{1} \ \mathbf{for} \ i = 0 \ to \ n \ \mathbf{do}
     2 | L(i,0) = i
    з end
    4 for j=0 to m do
    5 | L(0,j) = j
    6 end
    7 for i = 1 to n do
                                        for j = 1to m do
    8
                                                           L(i, j) = \min \{L(i-1, j) + 1, L(i, j-1) + 1, L(i-1, j-1) + 1, L(i-1, j-1) + 1, L(i-1, i-1) 
       9
                                                                     c(a_i,b_j)
                                       end
10
11 end
12 return L(m, n)
```

```
Algorithm 2: Hierarchical Clustering Algorithm
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```
input: An integer n represents the number
            of initial clusters, a clusterMatrix
            stores the sentence vector of every
            cluster, an array, a
            queue(implemented by binary heap)
            stores indices and distances between
            two cluster
   {f output:} clustered nodes
 \mathbf{1} for i=1 to n do
      {\rm initiaDistance}(i,i)=0;
2
      for j = i + 1to n do
          4
           i, j, 1, 1)
      end
 5
 6 end
7 while n \geq miniumClusters do
      entry = queue.pop();
      merge(entry.cluster1, entry.cluster2,
 9
       entry.distance, entry.distance,
       clusterMatrix);
      for i = 1 to n do
10
11
          if i \neq
           entry.cluster1 and cluster Matrix[i].size \neq
           0 then
12
              first = min(entry.cluster1, i);
             second = min(entry.cluster2, i);
13
14
             newDistance =
               {\tt getDistance} ({\tt initialDistance},
               clusterMatrix(first),
               clusterMatrix(second));
              queue add entry(newDistance,
15
               first, second,
               clusterMatrix(first).size,
               {\rm clusterMatrix}({\rm second}).{\rm size})
16
          end
      \mathbf{end}
17
18
      n = n - 1;
19 end
20
```

Algorithm 3: Node pair similarity scores pseudocode input: An array of node pairs called computed\_pairs output: A hash table of similarity scores for every node pair in computed\_pairs 1 Create hash table node\_scores; 2 for (a,b) in  $nodes(G_-2)$ : do if a==b then  $Node\_scores[(a,b)] := 1$ 4 5  $| Node\_scores[(a,b)] := 0$ 6 7 end s end  $9 \text{ stop\_signal} := \text{off};$ 10 while  $stop\_signal == off do$  $stop\_signal := on;$ 11 for (a,b) in  $nodes(G_{-2})$  do if a != b then 13 {COMMENT: Let N(v) denote the 14 neighbors of node v }  $R_k(a,b) :=$ 15  $\frac{C_1}{|N(a)||N(b)|} \sum_{i=1}^{|N(a)|} \sum_{j=1}^{|N(b)|} R_{\mathbf{k}\text{-}\mathbf{1}}(N_{\mathbf{i}}(a), N_{\mathbf{j}}(b))$ end16 if  $|R_k(a,b) - R_{k-1}(a,b)| > 0.05$  then 17  $stop\_signal := off$ 18 19  $node\_scores[(a,b)] := R_k(a,b)$ 20 21 22 end 23 Return node\_scores

• Algorithms and Data Structures: In Algorithm 1, the major algorithm is Edit Distance in Dynamic Programming. The data structure is two-dimensional array, and the time complexity is  $\mathcal{O}(mn)$ . In Algorithm 2, we use Agglomerative Hierarchical Clustering Algorithm to cluster the documents. The time complexity of this algorithm is  $\mathcal{O}(n^2logn)$ . In Algorithm 3, the nodes with their IDs are stored in hash tables. The graphs, represented as adjacency lists, are also stored in hash tables. The node pairs and their scores are stored in hash tables. Similar tweets and hash-tags are stored in arrays.

# References

[1] T. Mikolov, I. Sutskever, K. Chen, G. Corrado, and J. Dean. Distributed representations of words and phrases and their compositionality. In Proceedings of the 26th International Conference on Neural Information Processing Systems, NIPS'13, pages 3111-3119, USA, 2013. Curran Associates Inc.

[2] Jeh, Glen, and Jennifer Widom. "SimRank: a measure of structural-context similarity." In Proceedings of the eighth ACM SIGKDD international conference on Knowledge discovery and data mining, pp. 538-543. ACM, 2002.