Independent Study

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Abstract. This is independent study mid-way report

- 1. Abstract. Hil!
- 2. Introduction. Hi2!
- **3. Baseline Approach.** Here is the baseline approach which can be roughtly divided into two parts: the first part is constructing symmetric matrix from the feature similarity along time, and the second part is to construct symmetric normalized Laplacian matrix and obtain the top m eigenvectors with the top m smallest eigenvalues, and then perform k-means for boundary detection.

Algorithm 1 Baseline Approach

Input: number of top m smallest eigenvalues

- 1: M = getSymmetricMatrix()
- 2: L = scipy.sparse.csgraph.laplacian()
- 3: eigVals, eigVecs = np.linalg.eig(L)
- 4: Y = getMthSmallest(eigVals, eigVecs, m)
- 5: return boundaryDetection(Y)

3.1. Symmetric and Laplacian Matrix. Hi2sub!

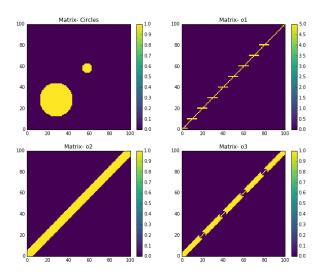


Figure 3.1. The training and test accuracy with epochs

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3.2. Boundary Detection. As the pseudocode in Algorithm 2, once the Laplacian matrix is constructed, each row is the representation of eigen-features at specific time. Therefore, running k-means for eigen-features of all time points will yield the results of which centroids of this time point belongs to, and therefore the place where the $t_i \neq t_{i+1}$ is where the boundary is. As showed in figure 3.1, boundary is correctly detected at each time point, but when doing experiments I noticed the initiation and number of iteration during k-means will affect the correctness of boundary, and the more iteration usually will gives better boundary.

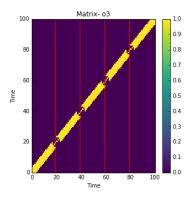


Figure 3.2. The original symmetric matrix and detected boundary

Algorithm 2 boundary Detection

Input: Laplacian eigenvectors $Y \in \mathbb{R}^{n \times m}$

Output: Boundary b, Centroids c

- 1: $\overline{y}_i = \frac{Y_i}{\|Y\|}$ //normalize each row Y_i
- 2: Run k-means on $\{\overline{y}_i\}_{i=1}^n$
- 3: Let c_i denote the cluster containing \overline{y}_i
- 4: $b \leftarrow \{i | c_i \neq c_{i+1}\}$
- 5: return b, c

3.3. ToDos. Hi3!

REFERENCES

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