



Constrained Laplacian Eigenmap for dimensionality reduction

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1. Introduction

ABSTRACT

Dimensionality reduction is a commonly used tool in machine learning, especially when dealing with high dimensional data. We consider semi-supervised graph based dimensionality reduction in this paper, and a novel dimensionality reduction algorithm called constrained Laplacian Eigenmap (CLE) is proposed. Suppose the data set contains *r* classes, and for each class we have some labeled points. CLE maps each data point into *r* different lines, and each map *i* tries to separate points belonging to class *i* from others by using label information. CLE constrains the solution space of Laplacian Eigenmap only to contain embedding results that are consistent with the labels. Then, each point is represented as a *r*-dimensional vector. Labeled points belonging to the same class are merged together, labeled points belonging to different classes are separated, and similar points are close to one another. We perform semi-supervised document clustering using CLE on two standard corpora. Experimental results show that CLE is very effective.

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In many real life applications, one is often confronted with high-dimensional data (e.g. documents, images). The infamous *curse of dimensionality* states that the performance of most machine learning algorithms degrades rapidly both in effectivity and efficiency as the dimensionality increases [15]. Dimensionality reduction allows one to represent the data in a lower dimensional space, and more importantly, reveal the intrinsic structure of the data [8,45,44,34]. The most popular techniques include principal component analysis (PCA), non-negative matrix factorization (NMF), and graph based dimensionality reduction.

Principal component analysis (PCA) [5,31,30] reduces the dimensionality of the data by finding a few orthogonal linear projections such that the variance of the projected data is maximized. In fact, it turns out that these projections are just the leading eigenvectors of the data's covariance matrix, which are called principal components. PCA is globally optimal in the sense that those projections best preserve the global Euclidean structure of the data. However, it can only discover linear

be handled perfectly. For multi-class problems, the algorithm does map points from the same class into the same point, but cannot guarantee that data points from different classes are mapped separately.

In this paper, we proposed a novel dimensionality reduction algorithm called constrained Laplacian Eigenmap (CLE), which supports multi-class problems. CLE aims to find the projection which respects the intrinsic geometrical structure inferred from all the data points, and also consists with labels. Suppose the data set contains r classes, and for each class we have some labeled points. CLE maps each data point into r different lines, and each map i tries to separate points belonging to class i from others by using label information. The mapping is performed by constraining the solution space of Laplacian Eigenmap to consist with labels. In the resulting r-dimensional space, labeled points belonging to the same class are merged together, labeled points belonging to different classes are separated, and similar points are close to one another.

We perform document clustering using CLE on two standard corpora: TDT2 and Reuters-21578. Experimental results show that this clustering model is very effective, and CLE becomes more competitive as the percentage of labeled documents or the number of clusters increases.

The paper is organized as follows: in Section 2, we give a brief review of related works. Our constrained Laplacian Eigenmap (CLE) algorithm is introduced in Section 3. The experimental results are presented in Section 4. Finally, we provide some concluding remarks and suggestions for future work in Section 5.

2. Related works

2.1. Principal component analysis (PCA)

PCA [5,31,30] can be defined in terms of the orthogonal projections which maximize the variance in the projected space. Given a set of *n*-dimensional data points $\mathbf{x}_1, \ldots, \mathbf{x}_m$, we first consider the projection onto a one-dimensional space using a *n*-dimensional vector **u**. Then, we have the following optimization problem:

$$\mathbf{u}_{opt} = \operatorname*{argmax}_{\mathbf{u}} \mathbf{u}^T S \mathbf{u} \tag{1}$$

with the constraint

 $\mathbf{u}^T \mathbf{u} = 1$

where *S* is the data covariance matrix defined by

$$S = \frac{1}{m} \sum_{i=1}^{m} (\mathbf{x}_i - \overline{\mathbf{x}}) (\mathbf{x}_i - \overline{\mathbf{x}})^T$$
(2)

It can be proved that \mathbf{u}_{opt} equals to the eigenvector of *S* that having the largest eigenvalue _1. This eigenvector is called as the first principal component. If we consider the general case of a *r*-dimensional projection space, the optimal linear projections for which the variance of the projected data is maximized are just the *r* leading eigenvectors $\mathbf{u}_1, \ldots, \mathbf{u}_r$ of *S*.

PCA is closely related to latent semantic indexing (LSI) [14] which projects documents onto a lower dimensional space through singular value decomposition (SVD).

2.2. Non-negative matrix factorization (NMF)

NMF [22,40,23] aims to find the non-negative factorization of

2.5. Constrained spectral clustering (CSC)

CSC is one closely related topic, since most clustering algorithms perform dimensionality reduction before clustering. We discuss two typical algorithms as follows.

Coleman et al. [13] have proposed one constrained spectral clustering algorithm, which supports must-link and cannot-link advices. Different from pervious methods, the proposed algorithm can only be applied to two-class clustering problems. Ji et al. [20] have proposed a document clustering model that enables the user to provide must-link constraints. Specifically, they introduce a constraint matrix U to encode user's prior knowledge. A penalty term that depends on U^TU is added to the objective function of spectral clustering. It is easy to check that U^TU is just the graph Laplacian of one graph constructed by adding an edge between any two must-link points. So, this algorithm is essentially equivalent to the semi-supervised graph embedding algorithm that increases the weight of the edge between any two must-link points.

3. Constrained Laplacian Eigenmap

Definition 1. The semi-supervised dimensionality reduction problem considered in this paper is defined as:

- 1. *Input*: a set of *n*-dimensional data points $\mathbf{x}_1, \dots, \mathbf{x}_m$ that belong to *r* classes, and for each class *i*, there are m_i points labeled.
- Output: the projection that respects the intrinsic geometrical structure inferred from all the data points and also consists with labels.

Denote class *i* as c_i , and the set of labeled points belonging to c_i as l_i .

3.1. Motivation

We can translate one multi-class dimensionality reduction problem into a set of two-class problems. This idea can be justified by considering the indicator matrix. $M \in \{0, 1\}^{m \times r}$ is an indicator matrix if $M_{ij} = 1$ iff $\mathbf{x}_i \in c_j$. Checking columns of Mindividually, we can see that the *i*-th column of M can only separate points belonging to c_i from others, so it can be seen as the embedding result of a two-class dimensionality reduction problem. However, by combining r such vectors to form M, it is sufficient to solve r-class problems, no matter r is 2 or greater.

Based on the above discussion, we propose a novel semisupervised dimensionality reduction algorithm called constrained Laplacian Eigenmap (CLE). If the data set contains r classes, CLE maps each data point into r different lines, and each map i tries to separate points belonging to c_i from others by using the label information. The mapping is performed based on Laplacian Eigenmap, whose solution space is constrained by *modifying the similarity matrix* and *using the subspace trick*. Putting the result of each map as a column, we can form a matrix $Y \in \mathbf{R}^{m \times r}$, where the *i*-th row gives the embedding coordinate of the *i*-th point.

3.2. Constraining the solution space of LE

Suppose we are going to map each point into the *k*-th line. The data set is treated as containing two classes: c_k and $\overline{c_k}$. Then, this map is required to separate points belonging to c_k from others by using the label information. We use two steps to constrain the solution space of Laplacian Eigenmap. Firstly, we incorporate label

information into the graph structure by modifying the similarity matrix. Since we are facing a two-class dimensionality reduction problem, the subspace trick is used later.

3.2.1. Modifying the similarity matrix

Since the data set is treated as containing two classes, we modify the similarity matrix *W* to make it more consist with this assumption. Assume the maximum similarity is 1. *W* is modified as follows:

1.
$$\forall i, j, \text{ if } \mathbf{x}_i, \mathbf{x}_j \in l_k, \text{ set } W_{ij} = 1.$$

2. $\forall i \ i \ \mathbf{if } \mathbf{x}_i \in l_a \ \mathbf{x}_i \in l_b, \ a \neq k \text{ and } b \neq k$

2. $\forall i, j, \text{ if } \mathbf{x}_i \in l_a, \mathbf{x}_j \in l_b, a \neq k, \text{ and } b \neq k, \text{ set } W_{ij} = 1.$ 3. $\forall i, j, \text{ if } \mathbf{x}_i \in l_k, \mathbf{x}_j \in l_a, \text{ and } a \neq k, \text{ set } W_{ij} = W_{ji} = 0.$

After modification, within class $(c_k \text{ or } \overline{c_k})$ links become tighter, and between class $(c_k \text{ and } \overline{c_k})$ links become looser. Unlike previous algorithm [21], the modification here is different for different map. Let W_k be the similarity matrix after modification, D_k be the corresponding diagonal matrix, and L_k be the corresponding Laplacian matrix.

3.2.2. Using the subspace trick

Ideally, we want to map all the points belonging to the same class (c_k or $\overline{c_k}$) into a single point. Since we only have a few labeled points for each class, the best we can do is to represent labeled points belonging to c_k by one 1-dimensional vector, represent other labeled points by another vector, and map similar points as closely as possible. Let $\mathbf{y} = [y_1, \dots, y_m]^T$ be the result of *k*-th map. We have the following objective function:

$$\mathbf{y}_{opt} = \operatorname*{argmin}_{\mathbf{y}} \sum_{i,j} (y_i - y_j)^2 (W_k)_{ij} = \operatorname*{argmin}_{\mathbf{y}} \mathbf{y}^T L_k \mathbf{y}$$
(7)

with constraints

$$\begin{array}{ll} y_i = y_j & \text{if } \mathbf{x}_i, \mathbf{x}_j \in l_k \\ y_i = y_j & \text{if } \mathbf{x}_i \in l_a, \mathbf{x}_j \in l_b, a \neq k, \text{ and } b \neq k \\ y_i \neq y_j & \text{if } \mathbf{x}_i \in l_k, \mathbf{x}_j \in l_a \text{ and } a \neq k \\ \mathbf{y}^T D_k \mathbf{y} = 1 \end{array}$$

Without loss of generality, assume the first m_1 points d_1, \ldots, d_{m_1} are labeled points belonging to c_1 , the next m_2 points $d_{m_1+1}, \ldots, d_{m_1+m_2}$ are labeled points belonging to c_2 , and so on. All the rest points are unlabeled. In order to meet the above constraint, we introduce the label constraint matrix $P_k \in \{0, \pm 1\}^{m \times (2+m-p)}$, where $p = m_1 + \cdots + m_r$. Each row *i* of P_k corresponds to point \mathbf{x}_i , and the matrix is represented as follows:

$$P_{k} = \begin{pmatrix} \mathbf{1}_{m_{1}} & -\mathbf{1}_{m_{1}} & \mathbf{0}_{m_{1} \times (m-p)} \\ \vdots & \vdots & \vdots \\ \mathbf{1}_{m_{k-1}} & -\mathbf{1}_{m_{k-1}} & \mathbf{0}_{m_{k-1} \times (m-p)} \\ \mathbf{1}_{m_{k}} & \mathbf{1}_{m_{k}} & \mathbf{0}_{m_{k} \times (m-p)} \\ \mathbf{1}_{m_{k+1}} & -\mathbf{1}_{m_{k+1}} & \mathbf{0}_{m_{k+1} \times (m-p)} \\ \vdots & \vdots & \vdots \\ \mathbf{1}_{m_{r}} & -\mathbf{1}_{m_{r}} & \mathbf{0}_{m_{r} \times (m-p)} \\ \mathbf{1}_{(m-p)} & \mathbf{0}_{(m-p)} & I_{(m-p) \times (m-p)} \end{pmatrix}$$

Using the label constraint matrix P_k , we can map labeled points into two different points by introducing an auxiliary vector z and equating:

$\mathbf{y} = P_k \mathbf{z}$

Substituting it into Eq. (7), we obtain the following optimization problem:

$$\mathbf{z}_{opt} = \arg\min_{\mathbf{z}} \mathbf{z}^T P_k^T L_k P_k \mathbf{z}$$
(8)

with the constraint

$$\mathbf{z}^T P_k^T D_k P_k \mathbf{z} = 1$$

Note that three constraints of Eq. (7) are dropped. They are automatically guaranteed by equation $\mathbf{y} = P_k \mathbf{z}$, which we will explain later. By the Rayleigh-Ritz theorem [28], we know the solution of this problem is the smallest eigenvector of the following generalized eigenvector problem:

$$P_k^T L_k P_k \mathbf{z} = P_k^T D_k P_k \mathbf{z}$$
(9)

Let \mathbf{z}_0 be the smallest eigenvalue of Eq. (9). We have the following proposition:

Proposition 1. The smallest eigenvalue \mathbf{z}_0 is 0 with eigenvector $[1,0,\ldots,0]^T$, and the corresponding **y** is the constant vector **1**.

Proof. It is easy to check that L_k is positive semi-definite and D_k is positive definite. Since the column vectors of P_k are independent, for any non-zero **z**, $P_k z$ is not a zero vector. Therefore, $P_k^T L_k P_k$ is still positive semi-definite and $P_k^T D_k P_k$ is positive definite. This implies that ≥ 0 . Actually, the smallest eigenvalue is 0 with eigenvector $[1, 0, \dots, 0]^T$, since

$$P_k^T L_k P_k [1, 0, ..., 0]^T = P_k^T L_k \mathbf{1} = P_k^T \mathbf{0} = 0$$

The corresponding $\mathbf{y} = P_k [1, 0, \dots, 0]^T = \mathbf{1}$. \Box

Vector 1 is useless, since all the data points have the same representations. So we choose the second smallest eigenvector \mathbf{z}_1 . Let $\mathbf{z}_1 = [z_1^1, z_2^1, ..., z_{2+m-p}^1]^T$, it is easy to check that

$$\mathbf{y} = \begin{bmatrix} (\underline{z_1^1 - z_2^1}), \dots, (\underline{z_1^1 - z_2^1})_{n_1 + \dots + n_{k-1}}, \underbrace{(\underline{z_1^1 + z_2^1}), \dots, (\underline{z_1^1 + z_2^1})_{n_k}}_{n_k, \dots, (\underline{z_1^1 - z_2^1}), \dots, (\underline{z_1^1 - z_2^1})_{n_{k+1} + \dots + n_r}}, (\underline{z_1^1 + z_3^1), \dots, (\underline{z_1^1 + z_2^1 + m - p})}_{n_k} \end{bmatrix}^T$$

So labeled points belongs to c_k are represented by $z_1^1 + z_2^1$, and other labeled points are represented by z_1^1 - z_2^1 . Since points belonging to same class are usually similar to one another, then in theory unlabeled points belonging to c_k will be mapped around $z_1^1 + z_2^1$, and other unlabeled points will be mapped around $z_1^1 - z_2^1$.

3.3. The algorithm

Given a set of data points $\mathbf{x}_1, \ldots, \mathbf{x}_m \in \mathbf{R}^n$ belonging to *r* classes and some labeled points for each class, CLE projects each points onto a r-dimensional discriminative space. It is performed as follows:

- 1. Constructing the adjacency graph. Let G denote a graph with m nodes, and the *i*-th node corresponds to the point \mathbf{x}_i . There are two variations [2]:
 - (1) ε neighborhoods [parameter $\varepsilon \in \mathbf{R}$]: Nodes *i* and *j* are connected by an edge if $\|\mathbf{x}_i - \mathbf{x}_i\|^2 < \varepsilon$ where the norm is the usual Euclidean norm in Rⁿ.
 - (2) k nearest neighbors [parameter $k \in \mathbf{N}$]: Nodes i and j are connected by an edge if \mathbf{x}_i is among k nearest neighbors of \mathbf{x}_i or \mathbf{x}_i is among k nearest neighbors of \mathbf{x}_i .
- Choosing the weights. There are many kinds of weighting methods [6]:
 - (1) 0-1 weighting,
 - (2) Gaussian kernel weighting,
 - (3) dot-product weighting, and
 - (4) polynomial kernel weighting.
- 3. Mapping each data point into r different lines. Each map i tries to separate points belonging to c_i from others by using the label information.

Suppose we are going to map each point into the *k*-th line. We use the following two steps to constrain the solution space of Laplacian Eigenmap (Section 3.2).

- (1) Modifying the similarity matrix:
 - If both of two labeled points belong to c_k , the edge weight is increased. If neither of two labeled points belong to c_k , the edge weight is also increased. If one labeled point belongs to c_k , and the other labeled point does not, the edge weight is decreased.
- (2) Using the subspace trick: Firstly, we construct the label constraint matrix P_k for the k-th map. Secondly, we calculate the second smallest eigenvector \mathbf{z}_1 of the generalized eigenvector problem (9). Then, $\mathbf{y} = P_k \mathbf{z}_1$ gives the result of the *k*-th map.
- 4. Combining the result of each mapping. Denote the results of all the *r* maps as $\mathbf{y}_1, \ldots, \mathbf{y}_r$. The embedding coordinate of point \mathbf{x}_i is given by the *i*-th row of $Y = [\mathbf{y}_1, \dots, \mathbf{y}_r]$.

After CLE, we get r-dimensional representations of the original data points. It is easy to check that labeled points belonging to the same class are merged together, labeled points belonging to different classes are separated, and similar points are close to one another. So, in this space, better classification or clustering performance can be obtained.

4. Experimental result

In this section, we perform document clustering using CLE to show the effectiveness of our algorithm. Document clustering has received a lot of attention as a fundamental tool for organization, summarization and retrieval of large volumes of text documents. Two standard document collections were used in the experiments: TDT2 and Reuters-21578.

4.1. Data corpora

We used the TDT2 and Reuters-21578 corpora as [6]. The TDT2 corpus¹ consists of documents collected from six sources (APW, NYT, VOA, PRI, CNN, and ABC) during 1998. It consists of 11,201 documents and 96 categories. In our experiments, we removed those documents belonging to two or more categories and used the largest 30 categories. This led to a data set with 9394 documents in 30 categories as described in Table 1. In this table, CluID means cluster's ID and DocNum means number of documents contained in the cluster.

Reuters-21578 corpus² contains 21.578 documents in 135 categories. In our experiments, we excluded those documents with multiple labels, and chose the largest 30 categories. It left us with 8067 documents in 30 categories as described in Table 2.

We removed the stop words, and represented each document as a term-frequency vector. Each document vector is normalized to unit length.

4.2. Evaluation metric

The clustering performance is evaluated by comparing the label obtained from our clustering algorithm with that provided by the document corpus. Two metrics, the accuracy (AC) and the normalized mutual information metric (\overline{MI}), are used to measure the clustering performance [40,6]. Given a document d_i , let p_i and

¹ Nist Topic Detection and Tracking corpus is at http://www.nist.gov/speech/

tests/tdt/1998/ ² Reuters-21578 corpus is at http://www.daviddlewis.com/resources/testcol lections/reuters21578/

- 3. *k*-means after LSI (LSI),
- 4. k-means after Laplacian Eigenmaps (LE),
- 5. k-means after semi-Laplacian Eigenmap (Semi-LE).

Since *k*-means algorithm can only find local minimum, and is sensitive to initial points. When we need to perform *k*-means, we apply it 10 times with different start points and the best result in

 q_i be the obtained cluster label and the label provided by the corpus, respectively. The *AC* is defined as follows:

$$AC = \frac{\sum_{i=1}^{m} \delta(q_i, \max(p_i))}{m}$$
(10)

where *m* is the total number of documents, $\delta(x, y)$ is the delta function that equals one if x = y and equals zero otherwise, and map(p_i) is the permutation mapping function that map each cluster label p_i to the equivalent label from the data corpus. The best mapping can be found by using the Kuhn–Munkres algorithm [27].

Let *C* denote the set of clusters provided by the document corpus and *C'* obtained from our algorithm. Their mutual information metric MI(C, C') is defined as following:

$$MI(C, C') = \sum_{c_i \in C, c'_j \in C'} p(c_i, c'_j) \cdot \log_2 \frac{p(c_i, c'_j)}{p(c_i) \cdot p(c'_j)},$$
(11)

where $p(c_i)$ and $p(c'_j)$ are the probabilities that a document arbitrarily selected from the corpus belongs to the clusters c_i and c'_j , respectively, and $p(c_i, c'_j)$ is the joint probability that the arbitrarily selected document belongs to the clusters c_i as well as c'_j at the same time. In our experiments, we use the normalized mutual information \overline{MI} as follows:

$$\overline{MI} = \frac{MI(C, C')}{\max(H(C), H(C'))}$$
(12)

where H(C) and H(C') are the entropies of *C* and *C'*, respectively. It is easy to check that \overline{MI} takes values between 0 and 1.

4.3. Clustering results

To demonstrate how our method improves the performance of document clustering, we compared it with the following five methods:

- 1. k-means on original term-document matrix (k-means),
- clustering based on Non-negative Matrix Factorization (NMF-NCW, [40]),

Table 3

Performance comparison on TDT2 corpus.

1	Accuracy											
	k-Means	LSI	NMF-NCW	LE	3%		5%		7%		9%	
					Semi-LE	CLE	Semi-LE	CLE	Semi-LE	CLE	Semi-LE	CLE
2	0.9328	0.9092	0.9827	0.9987	0.9987	0.9987	0.9988	0.9988	0.9988	0.9989	0.9989	0.9989
3	0.8953	0.8546	0.9379	0.9956	0.9957	0.9964	0.9958	0.9963	0.9962	0.9966	0.9959	0.9961
4	0.8547	0.7580	0.9268	0.9910	0.9951	0.9948	0.9951	0.9945	0.9955	0.9935	0.9959	0.9948
5	0.8392	0.7686	0.9229	0.9907	0.9841	0.9885	0.9874	0.9911	0.9887	0.9913	0.9890	0.9911
6	0.8303	0.7538	0.9407	0.9764	0.9786	0.9782	0.9768	0.9807	0.9804	0.9824	0.9749	0.9827
Ave.	0.8704 Mutual infe	0.8088 ormation	0.9422	0.9905	0.9904	0.9913	0.9908	0.9923	0.9919	0.9925	0.9909	0.9927
		ormation										
2	0.8487	0.7645	0.9451	0.9765	0.9771	0.9777	0.9780	0.9788	0.9785	0.9792	0.9796	0.9809
3	0.7980	0.7388	0.8695	0.9626	0.9638	0.9683	0.9649	0.9684	0.9677	0.9702	0.9660	0.9675
4	0.7809	0.6853	0.8819	0.9542	0.9695	0.9682	0.9704	0.9684	0.9713	0.9673	0.9731	0.9696
5	0.7803	0.7040	0.8777	0.9563	0.9496	0.9552	0.9569	0.9614	0.9584	0.9621	0.9599	0.9628
6	0.8008	0.7137	0.8937	0.9379	0.9402	0.9427	0.9423	0.9493	0.9451	0.9513	0.9403	0.9528
Ave.	0.8018	0.7213	0.8936	0.9575	0.9600	0.9624	0.9625	0.9652	0.9642	0.9660	0.9638	0.9667

Table 4

Performance comparison on Reuters corpus.

1	Accuracy											
	k-Means	LSI	NMF-NCW	LE	3%		5%		7%		9%	
					Semi-LE	CLE	Semi-LE	CLE	Semi-LE	CLE	Semi-LE	CLE
2 3 4 5 6 Ave.	0.7486 0.6434 0.6896 0.5328 0.4921 0.6213	0.7348 0.6297 0.6609 0.5151 0.4603 0.6002	0.8689 0.7247 0.7635 0.6539 0.6029 0.7228	0.8644 0.7996 0.7394 0.7203 0.6340 0.7515	0.8872 0.7996 0.7492 0.7495 0.6493 0.7669	0.9016 0.8024 0.7552 0.7493 0.6568 0.7731	0.9007 0.8173 0.7706 0.7697 0.6665 0.7850	0.9056 0.8272 0.7648 0.7763 0.7024 0.7952	0.9051 0.8293 0.7849 0.7912 0.6941 0.8009	0.9141 0.8447 0.7838 0.7982 0.7142 0.8110	0.9127 0.8456 0.8013 0.7985 0.7232 0.8163	0.9302 0.8595 0.8143 0.8297 0.7477 0.8363
	Mutual information											
2 3 4 5 6	0.3095 0.3867 0.4691 0.3726 0.3679	0.2728 0.3528 0.4485 0.3485 0.3365	0.5242 0.4266 0.5119 0.4225 0.4107	0.5240 0.5381 0.5102 0.5042 0.4498	0.5581 0.5397 0.5094 0.5171 0.4636	0.5865 0.5413 0.5045 0.5116 0.4618	0.5920 0.5643 0.5229 0.5395 0.4785	0.6023 0.5748 0.5257 0.5523 0.5087	0.5962 0.5859 0.5343 0.5573 0.4993	0.6143 0.6127 0.5482 0.5767 0.5267	0.6192 0.6087 0.5529 0.5627 0.5152	0.6613 0.6258 0.5841 0.6254 0.5576
Ave.	0.3811	0.3518	0.4592	0.5053	0.5176	0.5211	0.5395	0.5528	0.5546	0.5757	0.5718	0.6108

document space again. Similarly, when the percentage of labeled documents is 3% or more, CLE outperforms both LE and Semi-LE on average. But this time the improvement is much more obvious. With 9% labeled documents, the average *AC* of CLE is about 2% point higher than that of Semi-LE and 8.5% point higher than LE; the average \overline{MI} is about 4% point higher than of Semi-LE and 10.5% point higher than of LE.

Fig. 1 shows the accuracy of the six algorithms with different percentages of labeled documents and different cluster number on Reuters corpus. Apparently, the performance of Semi-LE and CLE becomes better as the percentage of labeled documents increases. More important, the improvement of CLE compared with Semi-LE becomes more obvious as the percentage of labeled documents e or the number of clusters r increases.

5. Conclusion and future work

In this paper, we propose a novel semi-supervised dimensionality reduction algorithm called constrained Laplacian Eigenmap (CLE). CLE aims to find the projection which respects the intrinsic geometrical structure inferred from all the data points, and also consists with labels. Document clustering experiments on TDT2 and Reuters-21578 corpora have shown that: when the label information is not scarce, CLE performs much better than the Semi-LE which incorporates label information through modifying the similarity matrix, and other unsupervised methods (*k*-means, LSI, NMF, LE). The advantage of CLE becomes more obvious as the percentage of labeled documents or the number of clusters increases.

Dimensionality reduction is widely used in face recognition. Eigenface [38], Fisherface [1], and Laplacianface [19,9] are three state-of-the-art face recognition techniques based on principal component analysis (PCA), linear discriminant analysis (LDA) and locality preserving projection (LPP), respectively. Compared with PCA and LDA, CLE can discover nonlinear manifolds. LPP seeks to preserve the intrinsic geometry of the data, but it is unsupervised. Previous studies have shown that face images are possibly reside on a nonlinear submanifold. Thus, CLE is very suitable for semisupervised face recognition, which will be investigated in our future work.



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