Projective Low-rank Subspace Clustering via Learning Deep Encoder

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Abstract

Low-rank subspace clustering (LRSC) has been considered as the state-of-the-art method on smal-1 datasets. LRSC constructs a desired similarity graph by low-rank representation (LRR), and employs a spectral clustering to segment the data samples. However, effectively applying LRSC into clustering big data becomes a challenge because both LRR and spectral clustering suffer from high computational cost. To address this challenge, we create a projective low-rank subspace clustering (PLrSC) scheme for large scale clustering problem. First, a small dataset is randomly sampled from big dataset. Second, our proposed predictive low-rank decomposition (PLD) is applied to train a deep encoder by using the small dataset, and the deep encoder is used to fast compute the low-rank representations of all data samples. Third, fast spectral clustering is employed to segment the representations. As a non-trivial contribution, we theoretically prove the deep encoder can universally approximate to the exact (or bounded) recovery of the row space. Experiments verify that our scheme outperforms the related methods on large scale datasets in a small amount of time. We achieve the state-of-art clustering accuracy by 95.8% on MNIST using scattering convolution features.

1 Introduction

Low-rank subspace clustering (LRSC) has become an important topic because of its impressive performance in many machine learning and computer vision applications, for example, image clustering [Zhang et al., 2015], motion segmentation [Wang et al., 2015; Zhao and Fu, 2015], and dictionary learning [Ding et al., 2016]. The underlying assumption is that data points sampled from multiple high-dimensional subspaces can be well represented by a union of low-dimensional subspaces. It leads to a useful self-expressiveness property that each data point can be efficiently reconstructed by a combination of other data points [Elhamifar and Vidal, 2013; Ding et al., 2015]. According to this property, LRSC is to learn low-rank representations for dividing the data samples into their respective subspaces. Hearteningly, many LRSC

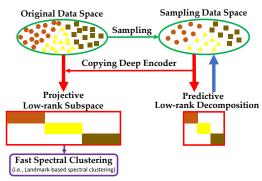


Figure 1: Projective low-rank subspace clustering by predictive low-rank decomposition.

methods [Li and Fu, 2015; Shen *et al.*, 2016] have been used to achieve the state-of-the-art results on small datasets. While as the data size grows, a critical problem comes, i.e. how to effectively apply LRSC to cluster big data.

Recall that the classical paradigm, LRSC consists of two steps. In the first step, the desired similarity graph is constructed by learning the lowest-rank representations of all data. The representations are obtained by using the low-rank representation (LRR) methods, such as robust principal component analysis (RPCA) [Candés et al., 2011], latent LRR (latLR-R) [Liu and Yan, 2011], and distributed LRR [Talwalkar et al., 2013]. Based on the self-expressiveness property, LRR and its variants usually select all data matrix as a dictionary. When facing a big dataset, this leads that LRR fails to work on a single machine with limited resource because of the high time and space complexity of the pseudoinverse and singular value decomposition (SVD). In order to reduce the number size of the dictionary, a small dataset sampled from the big dataset is regarded as a small dictionary [Wang et al., 2014; Peng et al., 2016]. However, it still takes more time to compute the codes (or representations) [Lin et al., 2011].

In the second step, spectral clustering (such as Normalized Cuts (NCut) [Shi and Malik, 2000]) is applied to the similarity graph for segmenting the data samples. However, it is a difficult problem to apply spectral clustering to big data tasks because of its computational complexity of $\mathcal{O}(n^3)$, where n is the number of samples [Cai and Chen, 2015]. Fortunately, fast spectral clustering (FSC) [Cai and Chen, 2015] can carry out the spectral clustering in linear time by sparsifying the similarity matrix. Scalable LRR (SLRR) [Peng et al., 2016]

and selection+LRR (selLRR) [Wang et al., 2014] select a small dataset to preform clustering to build a collaborative representation based classifier (CRC) or a linear classifier, respectively. Although these methods can deal with the big data clustering problem, they directly use the original data as the representations to lead to poor clustering results.

In light of the above arguments, we provide an efficient strategy to solve the big data clustering problem. To reduce the expensive computing time, we first select a small dataset from the large dataset for building a small LRR model, and then use the small dataset to learn a deep encoder to quickly approximate the low-rank representations of this small LRR model. After learning, the deep encoder is easily used to fast compute the low-rank representations of large dataset in the first step. Next, fast spectral clustering is employed to cluster the low-rank representations for avoiding the classification problem in the second step. The whole process is shown in Fig. 1. **Our contributions** are summarized as:

- We propose a predictive low-rank decomposition (PLD).
 PLD replaces the costly and highly non-linear SVD operations by a non-iterative deep encoder for quickly calculating the low-rank coding subspaces. We theoretically prove the deep encoder can universally approximate to the exact (or bounded) recovery of the row space.
- We create a projective low-rank subspace clustering (PLrSC) scheme to large scale clustering problem by a manner of "sampling, PLD, fast spectral clustering".
- Experimental results verify that our scheme outperforms the state-of-art baseline methods on large scale image datasets. It is important to note that we reach the best clustering accuracy 95.8% on MNIST-SC.

Notations: For a matrix $\mathbf{A} \in \mathbb{R}^{d \times n}$, the nuclear norm, ℓ_1 norm, $\ell_{2,1}$ norm, F-norm, and square of F-norm are denoted by $\|\mathbf{A}\|_* = \sum_i \sigma_i(\mathbf{A})$ (the sum of the singular values of \mathbf{A}), $\|\mathbf{A}\|_1 = \sum_{ij} |\mathbf{A}_{ij}|$, $\|\mathbf{A}\|_{2,1} = \sum_j \|\mathbf{A}_{:j}\|_2$, $\|\mathbf{A}\|_F = \sqrt{\sum_{ij} (\mathbf{A}_{ij})^2}$, and $\|\mathbf{A}\|_F^2 = \sum_{ij} (\mathbf{A}_{ij})^2$.

2 Related Works

We will review fast coding models, and fast spectral clustering. Fast Coding were presented to fast infer the codes by learning a (linear) encoder, such as projective low-rank representation (PLR) [Li et al., 2016; Li et al., 2014; Ding et al., 2016], RPCA encoder (RPCAec) [Sprechmann et al., 2015], and latLRR [Liu and Yan, 2011]. However, PLR was a supervised model for classification tasks, not subspace clustering. RPCAec assumed that the underlying data structure was a single low-rank subspace, while our model (PLD) could better handle the multiple subspaces data. Moreover, although latLRR could quickly calculate the codes by the linear encoder, it essentially resolved the insufficient sampling data [Liu and Yan, 2011], and the linear encoder was difficult to capture the complex data structure [Bengio et al., 2013; Li et al., 2017b]. Fortunately, PLD was to learn a deep encoder, which had power to draw the high-level features.

Fast Spectral Clustering (FSC) tried to speed up the spectral clustering algorithms due to the high computational complexity of the traditional spectral clustering, for example,

Nyström [Chen et al., 2011] and landmark-based spectral clustering (LSC) [Cai and Chen, 2015]. Nyström was to reduce the computational costs of eigen-decomposition over the whole similarity matrix [Fowlkes et al., 2004]. Moreover, it was parallelized both memory use and computation on distributed systems [Chen et al., 2011]. LSC [Cai and Chen, 2015] selected small data points to sparsely represent other data points for computing the spectral embedding of the data. Unfortunately, LSC and Nyström result in a poor result with the original input data. Our method is to learn "good" features by deep encoder, and employ LSC to cluster the features.

3 Predictive Low-rank Decomposition (PLD)

Before building our projective low-rank subspace clustering scheme, we firstly propose a predictive low-rank decomposition (PLD) model to fast calculate the projective low-rank representations in this section. Given a dataset $\mathbf{Y} = [\mathbf{Y}^1, \cdots, \mathbf{Y}^i, \cdots, \mathbf{Y}^k] \in \mathbb{R}^{d \times n}$, where each column is a data point, k is the number of subspaces, d is the number of dimensions, and n is the number of data points, we construct a formulation of PLD, and solve PLD by combining an alternating direction method (ADM) [Liu $et\ al.$, 2013] and a gradient descent algorithm [Li $et\ al.$, 2015].

3.1 Problem Formulation.

Large scale LRSC is restrained by solving LRR due to the high computational complexity of Singular Value Thresholding (SVT) operator [Cai et al., 2010]. Actually, sparse coding also faces the expensive inference problem [Elhamifar and Vidal, 2013]. In order to avoid the expensive inference, predictive sparse decomposition (PSD) [Gregor and LeCun, 2010; Li et al., 2017a] is to fast approximate the codes or representations learned from the sparse coding model by training a feedforward neural network. Inspired by PSD, we learn a non-iterative deep encoder $f_{de}(\mathbf{Y};\theta)$ (θ is the learning parameter) to approximate the low-rank representations. The deep encoder is used to quickly calculating the low-rank codes for replacing many costly non-linear SVT operations. Therefore, we propose a predictive low-rank decomposition (PLD), which is written as a following nonconvex optimization:

$$\min_{\mathbf{E},\theta} \|\mathbf{Z}\|_* + \lambda \|\mathbf{E}\|_{2,1} + \gamma \|\mathbf{Z} - f_{de}(\mathbf{Y};\theta)\|_{\mathrm{F}}^2$$
(1)
$$s.t. \ \mathbf{Y} = \mathbf{YZ} + \mathbf{E},$$

where $\mathbf{Y} \in \mathbb{R}^{d \times n}$ is the input data, \mathbf{Y} is chosen as the dictionary, $\mathbf{E} \in \mathbb{R}^{d \times n}$ is the sparse noise matrix, λ is the regularization parameter of the sparse noise term, $\|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{\mathrm{F}}^2$ is an approximation term, γ is a regularization parameter of the approximation term (usually, $\gamma = 1$), $f_{de}(\mathbf{Y}; \theta) = g(\mathbf{W}^L \cdots g(\mathbf{W}^i \cdots g(\mathbf{W}^2 \mathbf{Y}))$) is a deep encoder with L layers, g is an activation function (such as sigmoid, g tanh and ReLU), $g = \{\mathbf{W}^2, \cdots, \mathbf{W}^L\} \in \mathbb{R} = \{\mathbb{R}^{\ell_2 \times \ell_1}, \cdots, \mathbb{R}^{\ell_L \times \ell_{L-1}}\}$ is a learning parameter set, and ℓ_i is the number of units in the i-th layer $(\ell_1 = d$ and $\ell_L = n)$. For example, a deep encoder with L = 3 layers is $f_{de}(\mathbf{Y}; \theta) = g(\mathbf{W}^3 g(\mathbf{W}^2 \mathbf{Y}))$.

3.2 Optimization.

The problem (1) is non-convex because of the non-convex deep encoder. Clearly, it is an important challenge. We employ an alternating direction method (ADM) [Liu *et al.*, 2013] and a gradient descent algorithm [Li *et al.*, 2015] to solve (1). For efficiency, the problem (1) is first transformed into the following equivalent problem:

$$\min_{\mathbf{E}, \mathbf{J}, \mathbf{Z}, \theta} \|\mathbf{J}\|_* + \lambda \|\mathbf{E}\|_{2,1} + \gamma \|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{\mathrm{F}}^2$$

$$s.t. \mathbf{Y} = \mathbf{YZ} + \mathbf{E}. \mathbf{Z} = \mathbf{J}.$$
(2)

Then the problem (2) can be solved by minimizing the following augmented Lagrange function:

$$\mathcal{L} = \|\mathbf{J}\|_{*} + \lambda \|\mathbf{E}\|_{2,1} + \gamma \|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{F}^{2} + \mu(\|\mathbf{Y} - \mathbf{Y}\mathbf{Z} - \mathbf{E} + \mathbf{Q}_{1}/\mu\|_{F}^{2} + \|\mathbf{Z} - \mathbf{J} + \mathbf{Q}_{2}/\mu\|_{F}^{2})/2 - (\|\mathbf{Q}_{1}\|_{F}^{2} + \|\mathbf{Q}_{2}\|_{F}^{2})/(2\mu),$$
(3)

where \mathbf{Q}_1 and \mathbf{Q}_2 are the Lagrange multipliers, and μ is a penalty parameter. To tackle this unconstrained problem, it can be minimized by ADM, which is to iteratively update the variables $\{\mathbf{E}, \mathbf{J}, \mathbf{Z}, \theta\}$, the Lagrange multipliers $\{\mathbf{Q}_1, \mathbf{Q}_2\}$ and the penalty parameter μ until convergence. The iteratively key steps are as following:

Updating θ : θ is calculated by minimizing \mathcal{L} with respect to the weight parameters θ , while others are fixed. The optimal solution is to solve a subproblem¹: $\mathcal{L}_{\theta} = \|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{\mathrm{F}}^2$. By using a gradient descent (GD) algorithm [Li *et al.*, 2015] to minimize \mathcal{L}_{θ} , the updating form is

$$\theta = \theta - \zeta \partial \mathcal{L}_{\theta} / \partial \theta, \tag{4}$$

where ζ is a learning rate, and $\partial \mathcal{L}_{\theta}/\partial \theta$ is the gradient of \mathcal{L}_{θ} . Updating J: J is calculated by minimizing \mathcal{L} with respect to J, while others are fixed. The optimal solution is to solve a subproblem: $\mathcal{L}_{J} = \arg\min_{J} \frac{1}{\mu} \|\mathbf{J}\|_{*} + \frac{1}{2} \|\mathbf{J} - (\mathbf{Z} + \mathbf{Q}_{2}/\mu)\|_{F}^{2}$. This subproblem \mathcal{L}_{J} has a closed-form solution given by

$$\mathbf{J} = \Upsilon_{\frac{1}{\mu}}(\mathbf{Z} + \mathbf{Q}_2/\mu),\tag{5}$$

where $\Upsilon_{\alpha}(\mathbf{C}) = \mathbf{U}\mathcal{S}_{\alpha}\left[\Sigma\right]\mathbf{V}^{T}$ is the SVT operator [Cai *et al.*, 2010], $\mathbf{U}\Sigma\mathbf{V}^{T}$ is the singular value decomposition (SVD) of \mathbf{C} , and $\mathcal{S}_{\epsilon}[\cdot]$ is the shrinkage-thresholding operator defined as $\mathcal{S}_{\epsilon}[x] = \max\{|x| - \epsilon, 0\}\operatorname{sgn}(x)$.

Updating Z: Z is calculated by minimizing \mathcal{L} with respect to **Z**, while others are fixed. The optimal solution is to solve a subproblem: $\mathcal{L}_{\mathbf{Z}} = \arg\min_{\mathbf{Z}} \gamma \|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{\mathrm{F}}^2 + \mu(\|\mathbf{Y} - \mathbf{YZ} - \mathbf{E} + \mathbf{Q}_1/\mu\|_{\mathrm{F}}^2 + \|\mathbf{Z} - \mathbf{J} + \mathbf{Q}_2/\mu\|_{\mathrm{F}}^2)/2$. This subproblem $\mathcal{L}_{\mathbf{Z}}$ has a closed-form solution

$$\mathbf{Z} = (2\gamma \mathbf{I}/\mu + \mathbf{Y}^T \mathbf{Y})^{-1} (\mathbf{Y}^T (\mathbf{Y} - \mathbf{E}) + \mathbf{J} + (2\gamma f_{de}(\mathbf{Y}; \theta) + \mathbf{Y}^T \mathbf{Q}_1 - \mathbf{Q}_2)/\mu).$$
(6)

Due to the nonlinear deep encoder, the term $\gamma \| \mathbf{Z} - f_{de}(\mathbf{Y}; \theta) \|_{\mathrm{F}}^2$ will increase the number of iterations as $\mathcal{L}_{\mathbf{Z}}$ is difficult to guarantee convergence. For fast obtaining local convergence, we can remove this term, and $\mathcal{L}_{\mathbf{Z}}$ also has another closed-form solution \mathbf{Z} , which is as follow:

$$\mathbf{Z} = (\mathbf{I} + \mathbf{Y}^T \mathbf{Y})^{-1}$$
$$(\mathbf{Y}^T (\mathbf{Y} - \mathbf{E}) + \mathbf{J} + (\mathbf{Y}^T \mathbf{Q}_1 - \mathbf{Q}_2)/\mu). \tag{7}$$

Algorithm 1 PLD via ADM and gradient descent.

1: **Input:** A small data matrix **Y**.

```
2: Initialize: \theta is randomly initialized, \mathbf{Z} = 0, \mathbf{E} = 0, \mathbf{Q}_1 = 0, \mathbf{Q}_2 = 0, \zeta = 0.01, \mu = 10^{-5}, \mu_{max} = 10^{6}, \rho = 1.2, \epsilon = 10^{-4}.
  3: While not converged do
  4:
               repeat
  5:
                             update \theta by Eq. (4),
               until \|\mathbf{Z} - f_{de}(\mathbf{Y}; \theta)\|_{\mathrm{F}}^2 < \epsilon,
  6:
  7:
               update \mathbf{J} by Eq. (5),
               update Z by Eq. (6) or Eq. (7),
  8:
  9:
               update E by Eq. (9),
10:
               update \mathbf{Q}_1 and \mathbf{Q}_2 by Eq. (10) and Eq. (11).
               update the parameter \mu = \min\{\rho\mu, \mu_{max}\}, check: \|\mathbf{Y} - \mathbf{YZ} - \mathbf{E}\|_{\mathrm{F}}^2 < \epsilon and \|\mathbf{Z} - \mathbf{J}\|_{\mathrm{F}}^2 < \epsilon.
11:
12:
13: End
14: Return solutions \theta, Z and E.
```

Updating E: E is calculated by minimizing \mathcal{L} with respect to E, while others are fixed. The optimal solution is to solve the following subproblem \mathcal{L}_E , which is as follow:

$$\arg\min_{\mathbf{E}} \ \frac{\lambda}{\mu} \|\mathbf{E}\|_{2,1} + \frac{1}{2} \|\mathbf{E} - (\mathbf{Y} - \mathbf{YZ} + \mathbf{Q}_1/\mu)\|_{\mathrm{F}}^2. \tag{8}$$

According to the Lemma 4.1 [Liu *et al.*, 2013; Yang *et al.*, 2009], the solution of $\mathcal{L}_{\mathbf{E}}$ is $\mathbf{E} = \Psi_{\frac{\lambda}{\mu}}(\mathbf{Y} - \mathbf{YZ} + \mathbf{Q}_1/\mu)$, where $\Psi_{\alpha}(\mathbf{C})$ as follow:

$$\mathbf{E}_{:j} = \begin{cases} \frac{\|\mathbf{C}_{:j}\|_2 - \alpha}{\|\mathbf{C}_{:j}\|_2} \mathbf{C}_{:j}, & \text{if } \|\mathbf{C}_{:j}\|_2 > \alpha, \\ 0, & \text{otherwise.} \end{cases}$$
 (9)

Updating Q₁ and **Q**₂: Lagrange multipliers are updated by performing a gradient ascent as follow:

$$\mathbf{Q}_1 = \mathbf{Q}_1 + \mu(\mathbf{Y} - \mathbf{YZ} - \mathbf{E}), \tag{10}$$

$$\mathbf{Q}_2 = \mathbf{Q}_2 + \mu(\mathbf{Z} - \mathbf{J}). \tag{11}$$

The procedure iterates five steps and the penalty parameter until a stopping criteria is met. The whole optimization procedure is summarized in **Algorithm 1**. After training the deep encoder, the low-rank representations can be computed by this deep encoder in *linear* time to n.

3.3 Complexity and Convergence.

The major computation of Algorithm 1 are Steps 4 and 6. The complexity of Step 4 is $\mathcal{O}(n^3)$ as we use the GD algorithm [Li et al., 2015]. Based on LRR [Liu et al., 2013], the complexity of Step 6 is $\mathcal{O}(nr_Y^2 + r_Y^3)$, where r_Y is the rank of Y. The whole complexity of Algorithm 1 is $\mathcal{O}(Tn^3 + nr_Y^2 + r_Y^3)$, where T is the training epoch of training deep encoder, and n is the number of sampling data points. T is less than 5 epochs.

Due to the non-smooth function and the nonlinear deep encoder, it is difficult to theoretically prove the convergence. Fortunately, there actually exist three sufficient conditions for ensuring the convergence of Algorithm 1. The first two conditions are that the dictionary matrix **Y** is of full column rank, and the optimality gap produced in each iteration step is monotonically decreasing [Liu *et al.*, 2013]. The third

¹Here, we do not show the regularization technique in deep encoder. Generally, the Frobenius norm or ℓ_2 norm can be also used to prevent the over-fitting of weights.

condition is that any continuous activation function can be approximated uniformly on compact by an over-three layers deep encoder with enough hidden units [Ripley, 1996]. This condition results in that the deep encoder can approximate the low-rank representation by using the gradient descent algorithm [Haykin, 2009]. Notation that while using (7) to solve $\mathcal{L}_{\mathbf{Z}}$, we can move the steps 3-5 to end in Algorithm 1 for saving the training iteration time.

4 Projective Low-rank Subspace Clustering

Suppose a big dataset $\mathbf{X} = [\mathbf{X}^1, \cdots, \mathbf{X}^i, \cdots, \mathbf{X}^k] \in \mathbb{R}^{d \times m}$ is over-sufficiently drawn from a union of k subspaces \mathcal{S}_i , where $r \ll m, m = \sum_{i=1}^k m_i, r = \sum_{i=1}^k r_i = \sum_{i=1}^k rank(\mathbf{X}^i), m_i$ is the number of data points \mathbf{X}^i of \mathcal{S}_i , and r_i is the number of the bases of \mathcal{S}_i . In this section, we create a Projective Lowrank Subspace Clustering (PLrSC) scheme for clustering \mathbf{X} by using PLD, and give an universal approximation theorem to recover the row space with bounded errors. The whole scheme of clustering the big dataset \mathbf{X} shown in Fig. 1 consists of sampling, PLD, and fast spectral clustering (FSC).

Sampling is used to reduce the number of data points. We assume that a small dataset $\mathbf{Y} = [\mathbf{Y}^1, \cdots, \mathbf{Y}^i, \cdots, \mathbf{Y}^k] \in$ $\mathbb{R}^{d \times n}$ sampled from **X** is still sufficient, that is, $r_i \ll n_i < m_i$, where $r_i = rank(\mathbf{Y}^i) = rank(\mathbf{X}^i)$ and n_i is the number of data points \mathbf{Y}^i . It leads to $r \ll n < m$, where $n = \sum_{i=1}^k n_i$. In many cases, **X** repeatedly drawn from a union of subspaces. Thus, by using some sampling or clustering techniques, we can always obtain the sampled dataset Y, which satisfies that, for any $\mathbf{x} \in \mathbf{X}$ and any small number η , there is a $\mathbf{y} \in \mathbf{Y}$ such that $\|\mathbf{y} - \mathbf{x}\|_2 \le \eta$. When n = m, $\eta = 0$. For simplicity, we use the function randperm in matlab to implement two sampling ways. In the first way, a small dataset Y is randomly sampled from the whole dataset X if each subspace has the similar number of data points. The second way is to randomly choose \mathbf{Y}^i from the *i*-th subspace \mathbf{X}^i for ensuring that every subspace has the similar number in Y. Note that the sampling method is not studied in this paper since we focus on PLD.

PLD can learn a non-iterative deep encoder $f_{de}(\cdot;\theta)$ from **Y** in the above section 3. Projective low-rank representations of the big dataset **X** can be quickly computed by the deep encoder $f_{de}(\mathbf{X};\theta)$. Clearly, the complexity of non-iterations deep encoder is *linear* in data point number m, that is, $\mathcal{O}(m)$.

FSC can fast perform the spectral clustering. In this paper we employ the landmark-based spectral clustering (LSC) [Cai and Chen, 2015] to cluster the projective low-rank representations. The complexity of LSC is $\mathcal{O}(m)$. Algorithm 2 summarizes the whole procedure of performing our PLrSC scheme. The complexity of Algorithm 2 is $\mathcal{O}(m)$.

Theoretical Analysis. In order to ensure the effectiveness of PLD in Algorithm 2, we give a theorem to show that the deep encoder can recover the row spaces of \mathbf{Y} and \mathbf{X} . For better understanding the effectiveness, the big dataset \mathbf{X} is divided into m/n datasets $\overline{\mathbf{X}} \in \mathbb{R}^{d \times n}$. In fact, we can repeatedly select some data points in \mathbf{X} to make m/n an integer.

Theorem 1. Let the size of clear data Y_0 , noise data Y, new clear data \overline{X}_0 and new noise data \overline{X} be $d \times n$, and the rank of Y_0 be r_0 . Suppose the skinny SVDs of Y_0 and \overline{X}_0 are

Algorithm 2 PLrSC via PLD

- 1: **Initialize:** Large data matrix \mathbf{X} , number m of subspaces.
- 2: randomly select Y from X,
- 3: learn a deep encoder $f_{de}(\cdot; \theta)$ by PLD,
- 4: fast calculate $\mathbf{Z}_{\mathbf{X}}$ by $f_{de}(\mathbf{X}; \theta)$,
- 5: perform LSC on $\mathbf{Z}_{\mathbf{X}}$ to segment \mathbf{X} into m clusters.

Table 1: Datasets.

Data set	# data points	Dimension	# classes
MNIST	70,000	784	10
MNIST-SC	70,000	3,472	10
NORB	48,600	2,048	5

respectively $U_0\Sigma_0V_0^T$ and $\overline{U}_0\overline{\Sigma}_0\overline{V}_0^T$, $\|\overline{X}_0 - Y_0\|_F \le n\eta$ and $\|\overline{X} - Y\|_F \le n\eta$. Given any continuous function g (sigmoid, tanh or ReLU), $\lambda > 0$, and $\epsilon > 0$, there exist parameters $\theta = \{W^2, W^3\}$ such that

$$||f_{de}(\mathbf{Y}_0;\theta) - \mathbf{V}_0 \mathbf{V}_0^T||_{\mathbf{F}} < \epsilon, \tag{12}$$

$$||f_{de}(\mathbf{Y};\theta) - \mathbf{V}_0 \mathbf{V}_0^T||_{\mathbf{F}} < \epsilon + \min\{d, n\} + r_0, \tag{13}$$

$$||f_{de}(\overline{\mathbf{X}}_0; \theta) - \overline{\mathbf{V}}_0 \overline{\mathbf{V}}_0^T||_{\mathbf{F}} < \epsilon + n\alpha_0 \eta + O(\eta),$$
 (14)

$$||f_{de}(\overline{\mathbf{X}};\theta) - \overline{\mathbf{V}}_0 \overline{\mathbf{V}}_0^T||_{\mathsf{F}} < \epsilon + \min\{d, n\} + r_0 + n\alpha\eta + O(\eta), \tag{15}$$

where $\alpha_0 = \|Y_0^{\dagger}V\|_F + \|\frac{\partial f_{de}(Y_0;\theta)}{\partial Y_0}\|_F$, $\alpha = \|Y^{\dagger}V\|_F + \|\frac{\partial f_{de}(Y_0;\theta)}{\partial Y_0}\|_F$, Y_0^{\dagger} and Y^{\dagger} are the pseudoinverses of Y_0 and

 $Y, V = I - \overline{V}_0 \overline{V}_0^T$, and $O(\eta)$ is a higher-order infinitesimal. For \mathbf{Y}_0 (or \mathbf{Y}) as the input data in (1), (12) (or (13)) naturally implies that $f_{de}(\mathbf{Y}_0;\theta)$ (or $f_{de}(\mathbf{Y};\theta)$) universally approximates to the exact (or bounded) recovery of the row space identified by $\mathbf{V}_0 \mathbf{V}_0^T$. Moreover, when $\overline{\mathbf{X}}_0$ (or $\overline{\mathbf{X}}$) is close to \mathbf{Y}_0 (or \mathbf{Y}), (14) (or (15)) also implies that $f_{de}(\overline{\mathbf{X}}_0;\theta)$ (or $f_{de}(\overline{\mathbf{X}};\theta)$) universally approximates to the bounded recovery of the row space identified by $\overline{\mathbf{V}}_0 \overline{\mathbf{V}}_0^T$. Moreover, an experiment verifies the deep encoder effectively approximates to the low-rank recovery of row space in the right picture of Fig. 2.

5 Experiments

In this section, several experiments were conducted to verify the best results of our PLrSC. First, we described the three real-world datasets. Second, many state-of-art methods and evaluation metric were introduced. Finally, we verified the effectiveness of PLD and the clustering results.

Datasets: We conducted experiments on three large scale datasets shown in Table 1. A brief description of the datasets is listed below. $MNIST^2$ contains 70,000 training and testing examples with 28×28 pixel greyscale images of handwritten digits 0-9. MNIST-SC is a variant of MNIST. We follow the settings [You *et al.*, 2016], and compute the feature vectors of each image by using a scattering convolution network [Bruna and Mallat, 2013]. Each feature vector is of size 3,472, which is a concatenation of coefficients in each layer of the network and is dropped to 500 dimensions by PCA. $NORB^3$ contains

²http://yann.lecun.com/exdb/mnist/

³http://www.cs.nyu.edu/ ylclab/data/norb-v1.0-small/

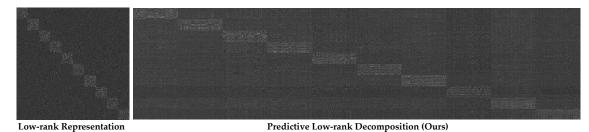


Figure 2: Illustrations of projective low-rank representations on MNIST-SC with 10 classes. Left: Representations produced by LRR on 500 samples. Right: Representations computed by deep encoder learned from PLD on 2,000 samples.

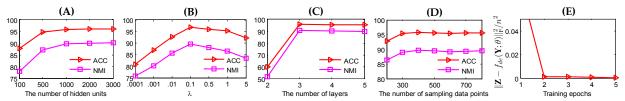


Figure 3: (A), (B), (C) and (D) show ACC (%) and NMI (%) with different hidden units, parameter λ , layers and sampling data points on MNIST-SC. (E) plots the convergence of approximated accuracy between low-rank representation and deep encoder.

images of 3D object toys belonging to 5 generic categories: four-legged animals, human figures, airplanes, trucks, and cars. There are 48,600 images combined the training samples with test samples. The original $2\times96\times96$ images were subsampled to $2\times32\times32$, and reduced to 500 by PCA.

5.1 Baseline Methods and Evaluation Metric.

We used the following algorithms as baseline methods.

LRR [Liu *et al.*, 2013] is to construct the similarity graph by learning low-rank representations and segment this graph by Normalized Cuts (NCut) [Shi and Malik, 2000]).

FSC is to fast build an similarity matrix for segmenting the data points, such as Nyström [Chen *et al.*, 2011], and landmark-based spectral clustering (LSC) [Cai and Chen, 2015]. Nyström was used to learn an approximate eigendecomposition of the whole similarity matrix, and NyströmO [Chen *et al.*, 2011] was constrained in the orthogonal eigenvectors. LSC [Cai and Chen, 2015] selected a few data points to build the similarity matrix. LSC-K used k-means to select the landmarks, while LSC-R was random.

Sampling-Clustering-Classification (S-C-C) is a scalable clustering strategy. Based on LRR, sparse subspace clustering (SSC) and least square regression (LSR), select+LRR (selLRR), select+SSC (selSSC) [Wang et al., 2014], scalable LRR (SLRR), scalable SSC (SSSC) and scalable LSR (SLSR) [Peng et al., 2016] are to sample a few data points to cluster the sampling data points, and then learn a linear classifier or CRC to segment the rest data.

PLrSC is our scalable clustering scheme in a pattern of "Sampling-PLD-FSC". We also sample a few data points to train a deep encoder by PLD for fast computing the low-rank representations, and employ LSC-K to segment the data points. Similarly, predictive sparse decomposition (PSD) [Gregor and LeCun, 2010], denoising autoencoder (DAE) [Vincent *et al.*, 2010], robust principal component analysis encoder (RPCAec) [Sprechmann *et al.*, 2015] and latent LRR (latLRR) [Liu and Yan, 2011] are used to learn sparse and low-rank codes.

Clustering results are measured by using Clustering Accu-

Table 2: Effectiveness of PLD compared with LRR on MNIST-SC (randomly selected 1200 samples).

	ACC		coding time (s)
PLrSC (ours)	88.6±3.96	81.4±3.50	0.025
LRR [Liu et al., 2013]	86.1±5.49	81.0±3.08	16.36

racy (ACC) [Cai and Chen, 2015] and Normalized Mutual Information (NMI) [Cai and Chen, 2015] between the clustering label and the ground truth label. ACC and NMI range from 0 to 1. A higher value reveals a better clustering result. All experiments were implemented in MATLAB R2015a and run on a Linux machine with 2.7 GHz CPU, 24GB memory. For each data set, the final results were reported as the average and variance of 10 times repetition.

5.2 Effectiveness and Parameter Analysis.

Before reporting clustering results, clustering time and parameter analysis, we first verify the effectiveness of our proposed PLD as PLrSC is built by PLD. PLD is to approximate low-rank codes learned from LRR by training a deep encoder. To verify this approximation, the first experiment is to compare our algorithm with LRR. Due to the limited space, we only select 1,200 training samples on MNIST-SC to conduct this experiment as LRR is nearly impossible to preform clustering on whole MNIST-SC. The results in Table 2 show that PLD has better ACC and NMI than LRR by 2.5% and 0.4% improvement. Furthermore, Fig. 3 (E) plots the convergence of approximated accuracy, and Fig. 2 illustrated the predictive low-rank representations calculated by the deep encoder.

The regularization parameter γ and the learning rate η are easy to set $\gamma=1$ and $\eta=0.001$ or 0.0001 in this paper. Particularly, we focus on the effects of λ , hidden units and layers, which are important to the proposed method, because λ controls the sparse term, and hidden units and layers restrict the power of deep encoder. Fig. 3 (A) shows PLrSC gets the best results when the number of hidden units was 2000 in the three-layer deep encoder in PLD. Fig. 3 (B) reveals that when $\lambda=0.1$ PLrSC reaches the best ACC and NMI. Fig. 3 (C) indicates that PLrSC achieves the good results when the

Table 3: ACC (%) and NMI (%) on large scale datasets. The randomly selected number of samples in MNIST-SC, MNIST and NORB are 500, 2000, 3000. The final results were reported as the average and variance of 10 times repetition.

	MNIST-	SC (500)	MNIST	(2000)	NORB	(3000)
	ACC	NMI	ACC	NMI	ACC	NMI
	Fast Spectral Clustering (FSC)					
Nyström [Chen et al., 2011]	70.8 ± 7.56	64.2±3.58	52.7±1.46	47.4 ± 0.38	37.1 ± 0.84	18.7±0.59
NyströmO [Chen et al., 2011]	71.0 ± 7.43	63.9 ± 3.65	52.1±3.48	46.9 ± 1.49	37.3 ± 0.88	18.9±0.25
LSC-R [Cai and Chen, 2015]	81.2 ± 1.06	73.9 ± 1.11	58.5±4.19	55.8±2.65	43.3±2.29	27.6±2.33
LSC-K [Cai and Chen, 2015]	85.9 ± 0.34	80.2±0.45	68.3±4.99	67.7±2.29	45.5±1.35	30.9 ± 0.52
	Sampling-Clustering-Classification (S-C-C)					
selSSC [Wang et al., 2014]	74.7±5.58	69.9±2.93	55.2±1.63	54.4±2.19	36.0±3.36	18.9±6.75
selLRR [Wang et al., 2014]	75.4 ± 4.31	69.2±2.27	55.4±5.11	52.1±2.50	40.6 ± 0.18	21.5 ± 0.21
SLSR [Peng et al., 2016]	75.4 ± 5.60	70.6 ± 2.91	54.1±1.56	48.1 ± 0.87	39.9 ± 0.31	20.6±0.15
SLRR [Peng et al., 2016]	79.7 ± 0.68	76.5 ± 1.44	50.0±3.87	49.1 ± 2.27	40.1 ± 0.74	21.6±0.81
SSSC [Peng et al., 2016]	78.9 ± 4.88	76.4±3.25	54.9±1.89	49.9±1.15	39.3±2.00	15.1±3.06
Sampling-Projective Coding-Fast Clustering (S-PC-FC)						
DAE [Vincent et al., 2010]	88.0±5.90	82.4±2.80	68.9±3.71	65.6±1.67	36.8±3.79	17.1±4.49
PSD [Gregor and LeCun, 2010]	79.0 ± 3.06	77.3 ± 2.10	50.1±1.79	43.7 ± 0.38	41.3±3.46	24.4±2.43
latLRR [Liu and Yan, 2011]	88.8±5.85	83.1±2.47	53.2±2.38	49.5 ± 1.21	39.0 ± 0.45	21.6±0.99
RPCAec [Sprechmann et al., 2015]	71.0 ± 5.44	64.6±4.06	54.6±5.95	52.2±4.16	45.3 ± 0.97	31.2±2.53
PLrSC (ours)	95.8±0.18	89.6±0.31	71.1±2.98	68.1±1.51	50.3±1.61	36.5±3.66

number of layers is 3 in PLD. Fig. 3 (D) shows ACC and NMI with different numbers of sampling data points, and there are similar results when the number is over 400.

5.3 Clustering Results.

All clustering results on MNIST-SC, MNIST, and JCNORB are shown in Table 3. From Table 3, we observe that our approach (PLrSC) performs best among the existing methods. Note that our result on MNIST-SC exceeded the state-of-art ACC 93.8% [You et al., 2016] by more than 2%, and 94.3% [Li et al., 2017a] by more than 1%. The former spends 1,680 seconds, while our model only takes 28 seconds. Moreover, the latter only uses 6,000 data points.

Compared to RPCAed, DAE PSD and latLRR, PLrSC performs best among all the fast coding methods although they have the similar inference times. The important reason is that PLrSC can effectively compute the predictive low-rank representation by deep encoder in Fig. 2. In the experiments, we use the best three-layer deep encoder in RPCAed, DAE and PSD, while latLRR is self-restricted to a linear encoder. In fact, PLrSC has higher ACC than the predictive methods by average 15.4% (RPCAed), 12.1% (latLRR), 15.6% (PSD) and 7.8% (DAE) improvement on the three datasets. PLrSC also is better in NMI than the predictive methods by at least 9.7% improvement on average.

Compared to selLRR, selSSC, SLRR, SSSC and SLSR, PLrSC's performance is much better, since it makes at least 14.7% ACC and 15.7% NMI improvement in Table 3. The important reason is that the ill clustering accuracy on the selected samples leads to poor classification results, while PLrSC quickly computes the low-rank codes and employs LSC-K to segment the data points. Compared to LSC-K, LSC-R, Nyström and NyströmO, fast coding is effective for FSC. Actually, Table 3 shows that PLrSC also registers at leat 5.8% ACC and 5.1% NMI improvement over FSC. This reveals that fast low-rank representations is very important to FSC.

We report the coding and clustering time costs in Tables 2 and 4, respectively. From Table 2, we see that PLD is at least 600 times faster than LRR in the coding time as it only

Table 4: Overall clustering time (s) compared to S-C-C.

	MNIST-SC	MNIST	NORB
selSSC [Wang et al., 2014]	310.62	739.58	759.32
selLRR [Wang et al., 2014]	549.21	1569.15	2078.52
SLSR [Peng et al., 2016]	289.91	601.23	620.39
SLRR [Peng et al., 2016]	283.25	638.76	667.57
SSSC [Peng et al., 2016]	291.53	613.01	651.16
PLrSC (ours)	27.12	93.53	43.61

computes a non-iterative deep encoder. Furthermore, Table 4 also shows that PLrSC was at least 6 times faster than S-C-C in the whole clustering time. Although PLrSC has the similar clustering time to the other S-PC-FC methods, it achieves the best results in Table 3. Compared to FCS, PLrSC needs a little more time cost as it uses the deep encoder to fast compute the low-rank codes. In fact, the coding time of PLrSC can be ignored when increasing the number of data points.

6 Conclusions

In this paper we have proposed a fast low-rank coding model, named predictive low-rank decomposition (PLD), which trained a deep encoder to approximate the low-rank codes. It was solved by an alternating direction method and a gradient descent algorithm. Based on PLD, we introduced a projective low-rank subspace clustering (PLrSC) scheme to segment big dataset. This scheme firstly sampled a few data points from big datasets, learned a deep encoder by PLD, computed low-rank codes of all data points by deep encoder, and employed fast spectral clustering to cluster the low-rank codes for segmenting big dataset. Experimental results have verified that PLrSC outperforms several state-of-the-art baseline methods.

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