

Adaptive Graph Convolutional Subspace Clustering

Lai Wei, Zhengwei Chen, Jun Yin, Changming Zhu, Rigui Zhou, Jin Liu
Shanghai Maritime University
Haigang Avenue 1550, Shanghai, China

weilai@shmtu.edu.cn, 965976272@qq.com, junyin@shmtu.edu.cn, cmzhu@shmtu.edu.cn
rgzhou@shmtu.edu.cn, jinliu@shmtu.edu.cn

Abstract

Spectral-type subspace clustering algorithms have shown excellent performance in many subspace clustering applications. The existing spectral-type subspace clustering algorithms either focus on designing constraints for the reconstruction coefficient matrix or feature extraction methods for finding latent features of original data samples. In this paper, inspired by graph convolutional networks, we use the graph convolution technique to develop a feature extraction method and a coefficient matrix constraint simultaneously. And the graph-convolutional operator is updated iteratively and adaptively in our proposed algorithm. Hence, we call the proposed method adaptive graph convolutional subspace clustering (AGCSC). We claim that, by using AGCSC, the aggregated feature representation of original data samples is suitable for subspace clustering, and the coefficient matrix could reveal the subspace structure of the original data set more faithfully. Finally, plenty of subspace clustering experiments prove our conclusions and show that AGCSC¹ outperforms some related methods as well as some deep models.

1. Introduction

Subspace clustering has become an attractive topic in machine learning and computer vision fields due to its success in a variety of applications, such as image processing [14, 48], motion segmentation [6, 17], and face clustering [27]. The goal of subspace clustering is to arrange the high-dimensional data samples into a union of linear subspaces where they are generated [1, 25, 36]. In the past decades, different types of subspace clustering algorithms have been proposed [3, 11, 23, 37, 44]. Among them, spectral-type subspace clustering methods have shown promising performance in many real-world tasks.

¹We present the codes of AGCSC and the evaluated algorithms on <https://github.com/weilyshmtu/AGCSC>.

Suppose a data matrix $\mathbf{X} = [\mathbf{x}_1; \mathbf{x}_2; \dots; \mathbf{x}_n] \in \mathcal{R}^{n \times d}$ contains n data samples drawn from k subspaces, and d is the number of features. The general formulation of a spectral-type subspace clustering algorithm could be expressed as follows:

$$\min_{\mathbf{C}} \Omega(\Phi(\mathbf{X}) - \mathbf{C}\Phi(\mathbf{X})) + \lambda\Psi(\mathbf{C}), \quad (1)$$

where $\Phi(\cdot)$ is a function that is used to find the meaningful latent features for original data samples. It could be either a linear or a non-linear feature extraction method [26, 42, 45], or even a deep neural network [27]. $\mathbf{C} \in \mathcal{R}^{n \times n}$ is the reconstruction coefficient matrix and $\Psi(\mathbf{C})$ is usually some kind of constraint of \mathbf{C} . In addition, $\Omega(\cdot)$ is a function to measure the reconstruction residual, and λ is a hyper-parameter. After \mathbf{C} is obtained, an affinity graph \mathbf{A} is defined as $\mathbf{A} = (|\mathbf{C}| + |\mathbf{C}^\top|)/2$, where \mathbf{C}^\top is the transpose of \mathbf{C} . Then a certain spectral clustering, e.g. Normalized cuts (Ncuts) [33] is used to produce the final clustering results.

Classical spectral-type subspace clustering algorithms mainly focus on designing $\Psi(\mathbf{C})$ to help \mathbf{C} to carry certain characteristics and hope \mathbf{C} can reveal the intrinsic structure of the original data set. For example, sparse subspace clustering (SSC) [6] lets $\Psi(\mathbf{C}) = \|\mathbf{C}\|_1$, which makes \mathbf{C} a sparse matrix. In low-rank representation (LRR) [17], $\Psi(\mathbf{C})$ is the nuclear norm of \mathbf{C} which helps discover the global structure of a data set. Least square regression (LSR) [20] aims to find a dense reconstruction coefficient matrix by setting $\Psi(\mathbf{C}) = \|\mathbf{C}\|_F^2$. Block diagonal representation (BDR) [19] makes $\Psi(\mathbf{C})$ a k -block diagonal regularizer to pursue a k -block diagonal coefficient matrix.

Recently, deep subspace clustering algorithms (DSCs) reported much better results than the classical spectral-type subspace clustering algorithms. The main difference between (DSCs) and the classical spectral-type subspace clustering algorithms is that DSCs use deep auto-encoders to extract latent features from original data [21, 27, 29]. But it is pointed out that the success of DSCs may be attributed to the usage of an ad-hoc post-processing strategy [8]. Though the rationality of the existing DSCs is required further dis-

discussion, some deep learning techniques are still worthy of being introduced into spectral-type subspace clustering algorithms.

In this paper, inspired by graph convolutional networks [4, 12, 40], we explore the problem of using graph convolutional techniques to design the feature extraction function $\Theta(\cdot)$ and the constraint function $\Psi(\cdot)$ simultaneously. Different from the existing graph convolution methods which need a predefined affinity graph, we apply the required coefficient matrix \mathbf{C} to construct a graph convolutional operator. So the graph convolutional operator will be updated adaptively and iteratively in the proposed subspace clustering algorithm. Consequently, on one hand, by applying the adaptive graph convolutional operator, the aggregated feature representations of original data samples in the same subspace will be gathered closer, and those in different subspaces will be separated further. On the other hand, in the obtained coefficient matrix, the coefficients corresponding to different data samples will also have a similar characteristic to the samples' feature representations, so it could reveal the intrinsic structure of data sets more accurately. The overview pipeline of the proposed method is illustrated in Fig. 1.

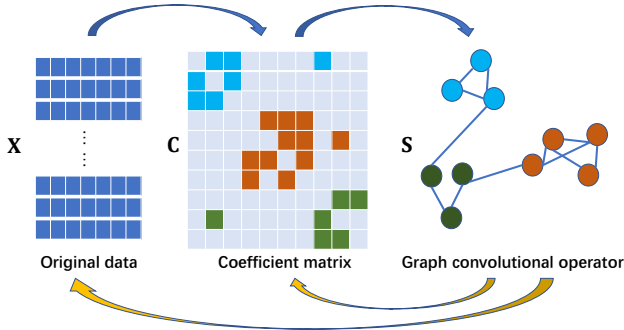


Figure 1. The overview of the proposed method. The graph convolutional operator \mathbf{S} will be updated iteratively based on \mathbf{C} . And the updated \mathbf{S} will in turn affect the computation of \mathbf{C} and feature aggregation.

2. Related Work

2.1. Graph convolutional networks (GCNs)

GCNs [12] learn new feature representations for a group of feature vectors \mathbf{X} by a multi-layer network. At the l -th layer of a GCN model, the features \mathbf{H}^{l-1} ($\mathbf{H}^0 = \mathbf{X}$) of each node are averaged with the feature vectors in its local neighborhood first. Then the aggregated features will be transformed linearly. Finally, a nonlinear activation (e.g. ReLU) is applied to output new feature representations \mathbf{H}^l . In summary, the updating function could be expressed as

follows:

$$\mathbf{H}_l \leftarrow \sigma(\mathbf{S}\mathbf{H}_{l-1}\mathbf{W}_{l-1}), \quad (2)$$

where $\mathbf{S} = \tilde{\mathbf{D}}^{-\frac{1}{2}}\tilde{\mathbf{A}}\tilde{\mathbf{D}}^{-\frac{1}{2}}$, $\tilde{\mathbf{A}} = \mathbf{A} + \mathbf{I}$, $\tilde{\mathbf{D}}$ is the degree matrix of $\tilde{\mathbf{A}}$. We here call \mathbf{S} a **graph convolutional operator**, (**GCO**). In addition, $\mathbf{A} \in \mathcal{R}^{n \times n}$ is a pre-defined adjacent matrix, $\mathbf{I} \in \mathcal{R}^{n \times n}$ is an identity matrix, \mathbf{W}_{l-1} is the linear transform matrix in the l -th layer, $\sigma(\cdot)$ represents the non-linear activate function.

SGC (simple graph convolution) claims that “the nonlinearity between GCN layers is not critical - but that the majority of the benefit arises from the local averaging” [39]. Therefore, SGC simplifies a graph convolutional layer as

$$\mathbf{H}_l = \mathbf{S}\mathbf{H}_{l-1}\mathbf{W}_{l-1}. \quad (3)$$

By integrating several graph convolutional layers, the final feature representations is

$$\mathbf{F} = \overbrace{\mathbf{S} \cdots \mathbf{S}}^M \mathbf{X} \mathbf{W}_1 \cdots \mathbf{W}_M = \mathbf{S}^M \mathbf{X} \mathbf{W} \quad (4)$$

where M is the number of graph convolutional layers, the $\mathbf{W} = \mathbf{W}_1 \cdots \mathbf{W}_M$ is also a linear transformation matrix. Compared to the traditional GCNs, SGC is much simple and achieves state-of-the-art results in many real-world applications.

2.2. Graph convolutional subspace clustering algorithms

Enlightened by SGC, Cai et al. devised a graph convolutional subspace clustering algorithm (GCSC) [2]. In GCSC, the new feature representations are directly obtained by 1-order GCO without linear transformation. Then GCSC defines an LSR-liked model. But different from LSR, the aggregated features obtained by a graph convolutional operator is used as the basis to reconstruct the original data samples.

Ma et al. proposed a graph filter LSR (FLSR) algorithm [24], which uses the coefficient matrix obtained by LSR to design the graph filter and then get the aggregated features. The two steps in FLSR are iteratively updated and the final obtained coefficient matrix will be used to get clustering results. In our opinion, both FLSR and GCSC focus on using graph convolutional techniques to devise the feature extraction function $\Phi(\cdot)$. Different from the two algorithms, our proposed method will use graph convolution techniques to design the $\Phi(\cdot)$ and $\Psi(\cdot)$ simultaneously.

3. Methodology

3.1. The proposed method

As described in Section 1, in spectral-type subspace clustering algorithms, the obtained reconstruction coefficient matrix \mathbf{C} is used to define the affinity matrix $\mathbf{A} =$

$(|\mathbf{C}| + |\mathbf{C}^\top|)/2$. If we impose some additional constraints on \mathbf{C} , such as $\mathbf{C} = \mathbf{C}^\top, \mathbf{C} \geq \mathbf{0}$, then $\mathbf{A} = \mathbf{C}$. Moreover, if \mathbf{C} also satisfies $\mathbf{C}\mathbf{1} = \mathbf{1}$ and $\text{diag}(\mathbf{C}) = \mathbf{0}$, then $\tilde{\mathbf{A}} = \mathbf{A} + \mathbf{I} = \mathbf{C} + \mathbf{I}$ and $\tilde{\mathbf{D}} = 2\mathbf{I}$. Here, $\mathbf{1} \in \mathcal{R}^{n \times 1} = [1, 1, \dots, 1]^\top$, $\text{diag}(\mathbf{C})$ is the diagonal vector of \mathbf{C} . Finally, we can deduce the graph convolutional operator $\mathbf{S} = \tilde{\mathbf{D}}^{-\frac{1}{2}} \tilde{\mathbf{A}} \tilde{\mathbf{D}}^{-\frac{1}{2}} = (\mathbf{C} + \mathbf{I})/2$.

After \mathbf{S} is defined, on one hand, we can get the new representation of the original data matrix as $\mathbf{F} = \mathbf{S}\mathbf{X} = \frac{1}{2}(\mathbf{C} + \mathbf{I})\mathbf{X} \implies 2\mathbf{F} = (\mathbf{C} + \mathbf{I})\mathbf{X}$. Suppose \mathbf{C} is a good reconstruction coefficient matrix², then the new feature representations of samples located in the same subspace will be more similar, while the new features of samples lie in a different subspace will be more different. Then similar to GCSC, we use \mathbf{F} to reconstruct \mathbf{X} and hope \mathbf{C} also be the reconstruction coefficient matrix, namely $\mathbf{X} = \mathbf{C}\mathbf{F}$.

On the other hand, in the subspace clustering domain, the reconstruction coefficient matrix is always seen as a representation of the original data matrix [17, 18]. Therefore, we could compute the new representation of \mathbf{C} as $\mathbf{S}\mathbf{C} = \frac{1}{2}(\mathbf{C} + \mathbf{I})\mathbf{C} = \frac{1}{2}(\mathbf{C}^2 + \mathbf{C})$. By applying the same assumption in the above paragraph, \mathbf{C} is hoped to faithfully reveal the subspace structure of data sets, then $\mathbf{S}\mathbf{C}$ should have a similar characteristic to \mathbf{C} . Therefore, we could define a constraint of \mathbf{C} as $\Psi(\mathbf{C}) = \|\mathbf{C} - \mathbf{S}\mathbf{C}\|_F^2 = \|\mathbf{C} - \frac{1}{2}(\mathbf{C}^2 + \mathbf{C})\|_F^2 = \|\frac{1}{2}\mathbf{C} - \frac{1}{2}\mathbf{C}^2\|_F^2 = \frac{1}{4}\|\mathbf{C} - \mathbf{C}^2\|_F^2$.

By collecting the above definitions and through some simple deductions, we could achieve the problem under the framework of Eq. (1) as follows:

$$\begin{aligned} \min_{\mathbf{F}, \mathbf{C}} \quad & \|2\mathbf{F} - (\mathbf{C} + \mathbf{I})\mathbf{X}\|_F^2 + \alpha\|\mathbf{X} - \mathbf{C}\mathbf{F}\|_F^2 \\ & + \beta\|\mathbf{C} - \mathbf{C}^2\|_F^2, \\ \text{s.t.} \quad & \mathbf{C} = \mathbf{C}^\top, \mathbf{C}\mathbf{1} = \mathbf{1}, \mathbf{C} \geq \mathbf{0}, \text{diag}(\mathbf{C}) = \mathbf{0}, \end{aligned} \quad (5)$$

where α and β are two positive parameters. In the proposed method, the graph convolutional operator $\mathbf{S} = \frac{1}{2}(\mathbf{C} + \mathbf{I})$ is updated adaptively and iteratively which is different from GCSC, hence we term Problem (5) adaptive graph convolutional subspace clustering (AGCSC).

3.2. Optimization

3.2.1 Optimization procedure

For the sake of solving Problem (5), we transfer it into the following equivalent problem:

$$\begin{aligned} \min_{\mathbf{F}, \mathbf{C}, \mathbf{Z}} \quad & \|2\mathbf{F} - (\mathbf{C} + \mathbf{I})\mathbf{X}\|_F^2 + \alpha\|\mathbf{X} - \mathbf{C}\mathbf{F}\|_F^2 \\ & + \beta\|\mathbf{C} - \mathbf{C}\mathbf{Z}\|_F^2, \\ \text{s.t.} \quad & \mathbf{C} = \mathbf{Z}, \mathbf{C}\mathbf{1} = \mathbf{1}, \\ & \mathbf{Z} = \mathbf{Z}^\top, \mathbf{Z} \geq \mathbf{0}, \text{diag}(\mathbf{Z}) = \mathbf{0}, \end{aligned} \quad (6)$$

where \mathbf{Z} is an auxiliary variable. The above problem could be solved by using ADMM (alternating direction method of

²Namely, its (i, j) -th element is zero when \mathbf{x}_i and \mathbf{x}_j come from the same subspace. Otherwise, the (i, j) -th element is non-zero.

multipliers method [16]). The augment Lagrangian function of Eq. (6) is:

$$\begin{aligned} \mathcal{L} = \quad & \|2\mathbf{F} - (\mathbf{C} + \mathbf{I})\mathbf{X}\|_F^2 + \alpha\|\mathbf{X} - \mathbf{C}\mathbf{F}\|_F^2 \\ & + \beta\|\mathbf{C} - \mathbf{C}\mathbf{Z}\|_F^2 + \langle \mathbf{\Gamma}, \mathbf{C} - \mathbf{Z} \rangle + \langle \mathbf{\Lambda}, \mathbf{C}\mathbf{1} - \mathbf{1} \rangle \\ & + \mu/2(\|\mathbf{C} - \mathbf{Z}\|_F^2 + \|\mathbf{C}\mathbf{1} - \mathbf{1}\|_F^2) \end{aligned} \quad (7)$$

where $\mathbf{\Gamma}, \mathbf{\Lambda}$ are two Lagrange multipliers, $\mu > 0$ is a parameter. By minimizing \mathcal{L} with respect to the variables $\mathbf{F}, \mathbf{C}, \mathbf{Z}$ one at a time while fixing the others at their latest values, the variables could be optimized alternately. Suppose t denotes the current iteration step, the precise updating schemes for the variables and Lagrange multipliers are described as follows:

$$\begin{cases} \mathbf{C}_{t+1} = (2\mathbf{X}\mathbf{X}^\top + 2\alpha\mathbf{F}_t\mathbf{X}^\top + 2\beta(\mathbf{I} - \mathbf{Z}_t)(\mathbf{I} - \mathbf{Z}_t)^\top \\ \quad + \mu_t(\mathbf{I} + \mathbf{1}\mathbf{1}^\top))(4\mathbf{F}_t\mathbf{X}^\top - 2\mathbf{X}\mathbf{X}^\top + 2\alpha\mathbf{X}\mathbf{F}_t^\top \\ \quad + \mu_t(\mathbf{Z}_t + \mathbf{1}\mathbf{1}^\top) - \mathbf{\Gamma}_t - \mathbf{\Lambda}_t\mathbf{1}^\top)^{-1}, \\ \mathbf{F}_{t+1} = (\alpha\mathbf{C}_{t+1}^\top\mathbf{C}_{t+1} + 2\mathbf{I})^{-1}((\mathbf{C}_{t+1} + \mathbf{I} \\ \quad + \alpha\mathbf{C}_{t+1}^\top)\mathbf{X}). \\ \mathbf{Z}_{t+1} = (2\beta\mathbf{C}_{t+1}^\top\mathbf{C}_{t+1} + \mu_t\mathbf{I})^{-1}(2\beta\mathbf{C}_{t+1}^\top\mathbf{C}_{t+1} \\ \quad + \mathbf{\Gamma}_t + \mu_t\mathbf{C}_{t+1}). \\ \mathbf{\Gamma}_{t+1} = \mathbf{\Gamma}_t + \mu_t(\mathbf{C}_{t+1} - \mathbf{Z}_{t+1}), \\ \mathbf{\Lambda}_{t+1} = \mathbf{\Lambda}_t + \mu_t(\mathbf{C}_{t+1}\mathbf{1} - \mathbf{1}), \\ \mu_{t+1} = \min(\mu_{max}, \rho\mu_t), \end{cases} \quad (8)$$

where μ_{max}, ρ are two given positive parameters. Moreover, after \mathbf{Z}_{t+1} is computed, we further let $\mathbf{Z}_{t+1} = \mathbf{Z}_{t+1} - \text{Diag}(\text{diag}(\mathbf{Z}_{t+1}))$, $\mathbf{Z}_{t+1} = \max(\mathbf{Z}_{t+1}, \mathbf{0})$, $\mathbf{Z}_{t+1} = (\mathbf{Z}_{t+1} + \mathbf{Z}_{t+1}^\top)/2$, where $\text{Diag}(\cdot)$ reformulates a vector to be the diagonal of a matrix. Then \mathbf{Z}_{t+1} satisfies the constraints w.r.t. \mathbf{Z} in Problem (6).

3.2.2 Algorithm

We summarize the algorithmic procedure for solving Problem (6) in Algorithm 1. After \mathbf{C} is obtained, an affinity matrix is defined as $\mathbf{A} = (|\mathbf{C}| + |\mathbf{C}^\top|)/2$. Then final clustering could be produced by applying Ncuts on \mathbf{A} .

We know that the convergence of ADMM for two variables has been well studied [16], however there are three variables in Problem (6). Fortunately, we could see that all the terms in the objective function of Problem (6) are strongly convex. Based on the **Theorem 4.1** presented in [9], it could be deduced the optimization procedure (Algorithm 1) is convergent.

Moreover, the computation time of Algorithm 1 is mainly rely on the updating of three variables \mathbf{C}, \mathbf{F} and \mathbf{Z} . We can see they all have closed form solutions in Eq.(e8). For updating each variable, it takes both $O(n^3)$ to compute the pseudo-inverse of an $n \times n$ matrix and the multiplication of two $n \times n$ matrices. Hence the time complexity of Algorithm 1 in each iteration taken together is $O(n)^3$. In

Algorithm 1 Adaptive graph convolutional subspace clustering

Input: The data matrix $\mathbf{X} = [\mathbf{x}_1; \mathbf{x}_2; \dots; \mathbf{x}_n]$, two parameters $\alpha, \beta > 0$, the maximal number of iteration *Maxiter*;

Output: The coefficient matrix \mathbf{C}_* and the new representation \mathbf{F}_* ;

- 1: Initialize the parameters, i.e., $t = 0$, $\mu_t = 10^{-6}$, $\mu_{max} = 10^{30}$, $\rho = 1.1$, $\varepsilon = 10^{-7}$ and $\mathbf{\Gamma}_t = \mathbf{Z}_t = \mathbf{0}$, $\mathbf{\Lambda}_t = \mathbf{0}$, $\mathbf{F}_t = \mathbf{X}$.
 - 2: **while** $\|\mathbf{C}_t - \mathbf{Z}_t\|_\infty > \varepsilon$, $\|\mathbf{C}_t \mathbf{1} - \mathbf{1}\|_\infty > \varepsilon$ and $t < \textit{Maxiter}$ **do**
 - 3: $t = t + 1$;
 - 4: Update variables $\mathbf{C}_t, \mathbf{F}_t, \mathbf{Z}_t$, Lagrange multipliers $\mathbf{\Gamma}_t, \mathbf{\Lambda}_t$ and parameter μ_t by using Eq. (8);
 - 5: **end while**
 - 6: **return** $\mathbf{C}_* = \mathbf{C}_t$ and $\mathbf{F}_* = \mathbf{F}_t$.
-

our experiments, the iterations of Algorithm 1 is always less than 500, hence its complexity is $O(n)^3$.

3.3. Further Discussion

We discuss the properties of the coefficient matrix obtained by Algorithm 1.

3.3.1 Block diagonal property

Minimizing $\Psi(\mathbf{C}) = \|\mathbf{C} - \mathbf{C}^2\|_F^2$ will lead \mathbf{C} to be an approximate idempotent matrix. $\mathbf{C} = \mathbf{I}$ and $\mathbf{C} = \mathbf{0}$ are two trivial solutions to Problem (5). Fortunately, constraints in Problem (5) will prevent \mathbf{C} to be the trivial solutions. Moreover, we have the following proposition, namely:

Proposition 1. *The coefficient matrix \mathbf{C} satisfies $\Psi(\mathbf{C})$ will be block diagonal.*

Firstly, in order to prove the proposition, we rewrite Problem (5) as follows:

$$\begin{aligned} \min_{\mathbf{C}} \quad & \|\mathbf{C} - \mathbf{C}^2\|_F^2 \\ \text{s.t.} \quad & 2\mathbf{F} = (\mathbf{C} + \mathbf{I})\mathbf{X}, \mathbf{X} = \mathbf{C}\mathbf{F}, \\ & \mathbf{C} = \mathbf{C}^\top, \mathbf{C}\mathbf{1} = \mathbf{1}, \mathbf{C} \geq \mathbf{0}, \text{diag}(\mathbf{C}) = \mathbf{0}. \end{aligned} \quad (9)$$

By combine the two equations in the second rows of Problem (9), we have $\mathbf{X} = \frac{1}{2}(\mathbf{C}^2 + \mathbf{C})\mathbf{X} \Rightarrow \mathbf{X} - \mathbf{C}\mathbf{X} + \frac{1}{2}\mathbf{C}\mathbf{X} - \frac{1}{2}\mathbf{C}^2\mathbf{X} = \mathbf{0} \Rightarrow (\frac{1}{2}\mathbf{C} + \mathbf{I})(\mathbf{X} - \mathbf{C}\mathbf{X}) = \mathbf{0}$. Because $(\frac{1}{2}\mathbf{C} + \mathbf{I})$ is invertible, hence $\mathbf{X} - \mathbf{C}\mathbf{X} = \mathbf{0}$. Therefore, Problem (5) could be regarded as a relax problem of the following problem:

$$\begin{aligned} \min_{\mathbf{C}} \quad & \|\mathbf{C} - \mathbf{C}^2\|_F^2 \\ \text{s.t.} \quad & \mathbf{X} = \mathbf{C}\mathbf{X}, \\ & \mathbf{C} = \mathbf{C}^\top, \mathbf{C}\mathbf{1} = \mathbf{1}, \mathbf{C} \geq \mathbf{0}, \text{diag}(\mathbf{C}) = \mathbf{0}. \end{aligned} \quad (10)$$

Problem (10) falls into the general subspace clustering formulation summarized in [19], namely

$$\min_{\mathbf{C}} f(\mathbf{X}, \mathbf{C}), \quad \text{s.t.} \quad \mathbf{X} = \mathbf{C}\mathbf{X}. \quad (11)$$

Secondly, for any permutation matrix \mathbf{P} , it is easily to verify that Problem (10) satisfies $f(\mathbf{X}, \mathbf{C}) = f(\mathbf{P}\mathbf{X}, \mathbf{P}\mathbf{C}\mathbf{P}^\top)$ which is the first EBD (enforced block diagonal) condition [19].

Thirdly, as described in Section 3.2.2, Algorithm 1 is convergent. Hence, Problem (5) has an unique solution $(\mathbf{F}^*, \mathbf{C}^*)$. For the same reason, Problem (10) also has an unique solution.

According to the above explanations and Theorem 3 in [19], we can conclude that the solution to Problem (10) will be block diagonal. Then as a relaxed problem, solving Problem (5) can also get an approximate block diagonal coefficient matrix.

3.3.2 Doubly stochastic property

The constraints $\mathbf{C} = \mathbf{C}^\top, \mathbf{C}\mathbf{1} = \mathbf{1}, \mathbf{C} \geq \mathbf{0}$ in Problem (5) restrict the reconstruction coefficient matrix \mathbf{C} to be a doubly stochastic matrix. Doubly stochastic matrices have a guarantee of a certain level of connectivity, which prohibits solutions with all-zero rows or columns (that can occur in other subspace clustering methods) [15]. And doubly stochastic normalization has been shown to significantly improve the performance of spectral clustering [38, 46, 47].

Moreover, due to the doubly stochastic property, for the i -th block of \mathbf{C} , we have $|\mathbf{C}_i]_{p,q} - \mathbf{C}_i]_{s,t}| \leq 1$, where $\mathbf{C}_i]_{p,q}$ and $\mathbf{C}_i]_{s,t}$ are the (p, q) -th and (s, t) -th elements in \mathbf{C}_i and $p, q, s, t \in 1, 2, \dots, n_i$, n_i denotes the number of sample in the i -th subspace and \mathbf{C}_i is the i -th block on \mathbf{C} . This means the differences in coefficients located in the same diagonal block will be small. This property can overcome the over-sparsity and get dense blocks [34] which is a benefit for obtaining good clustering results.

3.3.3 Post-processing strategy

As we mentioned in Section 1, post-processing could improve the performance of the existing subspace clustering algorithms [5, 8]. The frequently used post-processing strategy is to keep the m -largest values for each coefficient vector and discard the relatively small ones [24, 28]. The rationality of the post-processing strategy is because “the coefficients over intra-subspace data points are larger than those over inter-subspace data points.” [28]. The obtained coefficient matrices of AGCSC are approximate block-diagonal, this implies that the larger coefficients are computed over intra-subspace data points and small coefficients are computed over inter-subspace data points. Hence, we also could

Table 1. Detailed information of the seven benchmark datasets

Dataset	Samples	Classes	Size
ORL	400	40	32 × 32
YALEB	2432	38	48 × 42
Umist	480	20	32 × 32
COIL20	1440	20	32 × 32
COIL40	2880	40	32 × 32
MNIST	1000	10	28 × 28

use this kind of thresholding strategy to enhance the performance of AGCSC. In the following experiments, we call AGCSC with thresholding post-processing skill TAGCSC.

4. Experiments

In this section, we perform subspace clustering experiments to demonstrate the effectiveness of adaptive graph convolutional subspace clustering (AGCSC).

4.1. Dataset

We use several benchmark databases to verify the effectiveness of our proposed model. The data sets include three face image datasets (ORL, the extended Yale B and Umist), two object image datasets (COIL-20 and COIL-40), and one handwritten digit dataset (MNIST). ORL dataset [32] contains 400 face images with different poses and expressions of 40 persons. The extended Yale B (YALEB) [13] dataset has 2432 images of 38 persons with each person 64 near frontal images. Umist has 480 images with varied poses from 20 individuals. COIL20 dataset [31] is composed of 1440 images for 20 different objects. The images of each object are taken 5° apart as the object rotates on a turn table and each object has 72 images. COIL40 is similar to COIL20 which contains 2880 images for 40 different objects. MNIST dataset³ comprises 10 subjects, corresponding to 10 handwritten digits, namely “0”-“9”. For this database, we use the first 100 samples of each digit in the training data sets. The detailed information on these databases is summarized in Table 1.

In addition, the pixel value of each image belonging to some databases lies in [0, 255]. For efficient computation, we let each pixel value be divided by 255, so that the pixel value of each image fall into [0, 1]. This does not change the distribution of the original data sets.

4.2. Comparison methods and evaluation metrics

The representative and related subspace clustering models are compared in our experiments, including SSC [5], LRR [18], LSR [20], low rank subspace clustering (LRSC) [37], BDR [19], thresholding ridge regression (TRR) [28],

³<http://yann.lecun.com/exdb/mnist/>

TRR integrated with the graph filter method (FTRR), GCSC [2], FLSR [24]. AGCSC with thresholding post-processing skill (TAGCSC) is also evaluated.

Two popular metrics, i.e., clustering accuracy (ACC) and normalized mutual information (NMI), are applied to quantitatively evaluate the models’ performance.

ACC is defined as follows:

$$ACC = \frac{1}{n} \sum_{i=1}^n \delta(f(L_{pred}(i)), L_{true}(i)), \quad (12)$$

where L_{true} and L_{pred} denote the ground truth labels and the prediction labels of an algorithm. $L_{pred}(i)$ and $L_{true}(i)$ represent i -th points of L_{true} and L_{pred} respectively. $\delta(\cdot, \cdot)$ is the indicator function that satisfies $\delta(x, y) = 1$ if $x = y$, and $\delta(x, y) = 0$ otherwise, and $f(\cdot)$ is the best mapping function that permutes clustering labels to match the ground truth labels.

NMI is defined as

$$NMI = \frac{\mathbb{I}(L_{pred}, L_{true})}{\sqrt{\mathbb{H}(L_{pred})\mathbb{H}(L_{true})}}, \quad (13)$$

where $\mathbb{I}(\cdot, \cdot)$ is the mutual information that measures the information gain after knowing the partitions generated by an algorithm. $\mathbb{H}(L_{pred})$ and $\mathbb{H}(L_{true})$ is the entropy of L_{true} and L_{pred} respectively.

4.3. Parameter setting

Because parameters will influence the performance of the evaluated algorithms, hence we let the two parameters α and β in AGCSC vary in the set $\{1e - 5, 1e - 4, 1e - 3, 5e - 3, 0.01, 0.05, 0.1, 0.5\}$. And for the other evaluated methods, we will tune all the parameters by following the suggestions in the corresponding references. Especially, except BDR, the other compared models all have one hyper-parameter, we let the hyper-parameters in these algorithm change in the set $\{0.001, 0.01, 0.1, 1, 5, 10, 20, 50, 80, 100\}$. For BDR, the two parameters are selected in $\{0.1, 1, 10, 20, 50, 80\}$ and $\{0.001, 0.01, 0.1, 0.5, 1, 5, 10, 20, 50\}$ respectively. Additionally, the neighborhood size used in GCSC is fixed as 7. And for the thresholding skill-related methods including TRR, FTRR and TAGCSC, we let the threshold value m change from 4 to 10. Then in different experiments, the best performance of each algorithm will be recorded. The experiments are conducted on a Windows-based machine with an Intel i9-10900 CPU, 64-GB memory and MATLAB R2021b.

4.4. Clustering Results

We first summarize the clustering results in Table 2, where the best results are emphasized in bold and the second best results are denoted in bold and italics. From Table 2, we can get the following observations:

Table 2. Clustering results (in %) of various methods on the used benchmark data sets. The best results are emphasized in bold and the second best results are denoted in bold and italic.

Dataset	Metric	Method										
		SSC	LRR	LRSC	LSR	BDR	TRR	FLSR	FTRR	GCSC	AGCSC	TAGCSC
ORL	ACC	72.50	72.75	77.00	75.50	78.25	85.75	73.50	79.75	73.75	80.50	86.25
	NMI	84.52	83.26	85.02	84.97	88.46	91.49	83.84	87.60	83.98	88.51	92.84
YALEB	ACC	55.15	73.48	75.64	74.05	76.56	91.65	72.94	91.90	62.50	84.79	92.31
	NMI	55.71	77.11	78.32	78.13	80.34	93.03	76.57	93.18	68.04	87.37	94.04
Umist	ACC	52.92	64.79	63.33	64.17	64.92	74.38	60.62	69.37	79.58	81.04	90.83
	NMI	75.38	73.41	72.02	73.17	75.13	80.63	70.72	78.49	86.44	87.46	94.99
COIL20	ACC	68.61	70.14	71.81	69.17	71.71	85.97	69.93	86.53	79.79	88.75	98.96
	NMI	66.85	76.43	77.27	74.17	80.51	90.23	77.19	91.17	85.67	93.38	99.11
COIL40	ACC	63.13	60.42	58.23	56.88	57.25	65.00	62.88	71.39	73.72	78.12	92.60
	NMI	82.28	76.29	74.48	75.87	76.73	79.83	76.26	82.46	84.32	89.21	97.32
MNIST	ACC	63.70	64.60	64.30	62.80	61.30	67.70	65.10	66.40	67.70	71.40	72.80
	NMI	59.75	60.67	58.91	57.18	54.76	64.43	61.10	63.21	61.99	65.84	67.54

1. TAGCSC constantly achieves the best results and AGCSC also outperforms the other evaluated algorithms on all the data sets except ORL and YALEB. This shows that 1) the thresholding skill could improve the performance of AGCSC; 2) and it also implies that the obtained coefficient matrices obtained by AGCSC are block-diagonal, namely, the obtained coefficients by AGCSC over intra-subspace data points are larger than those over inter-subspace data points. Moreover, if we ignore those models using the thresholding skill (TRR, FTRR), we can see AGCSC outperforms the other algorithms. This means the coefficient matrices obtained by AGCSC could faithfully reveal the subspace structures of different data sets.

Take ORL as an example, we show the learned reconstruction coefficient matrices obtained by AGCSC, TAGCSC, and two other algorithms that get the competitive results (TRR and FTRR) in Fig. 2. For a clear comparison, we use the same color bar to denote the values in the four coefficient matrices. We can see the coefficient matrix obtained by AGCSC shows a more vital block diagonal characterization. For illustrating the effect of the thresholding skill, the partial coefficient matrices of AGCSC and TAGCSC corresponding to the samples from the first 10 classes are zoomed in Fig. 3. It can be seen that most coefficients that are eliminated are related to the samples existing in different subspaces.

2. On COIL20, COIL40 and Umist data sets, AGCSC and TAGCSC get excellent results. AGCSC even dominates the models with thresholding skills such as TRR and FTRR. Moreover, on the results obtained by AGCSC, TAGCSC increases the ACC and NMI by over 10% and 6% respectively. Specifically, on COIL40 data set, TAGCSC outperforms all the other methods including AGCSC by a large margin, the ACC and NMI are improved by over 18.5% and 9% respectively. Besides the effect of thresholding skill, we believe the good performance of AGCSC and TAGCSC also relies on the aggregated feature representation computed by AGCSC. We use t-SNE [35] to visualize the aggregated

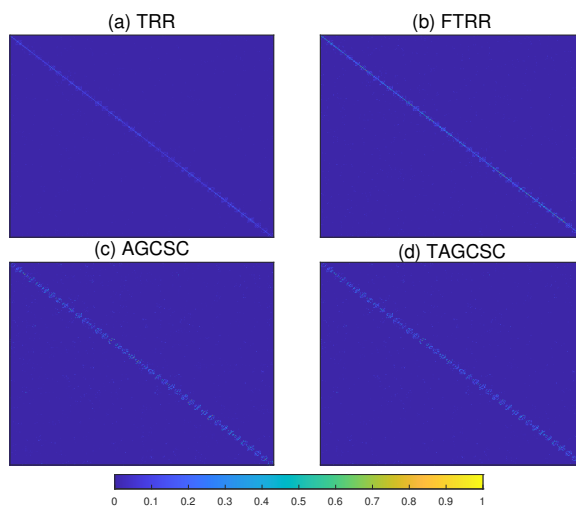


Figure 2. The obtained coefficient matrices obtained by (a) TRR, (b) FTRR, (c) AGCSC and (d) TAGCSC

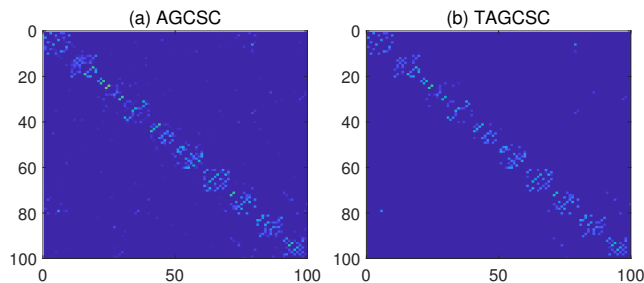


Figure 3. The partial coefficient matrices of (a) AGCSC and (b) TAGCSC corresponding to the samples from the first 10 classes.

gated feature representations from Umist data set obtained by AGCSC, GCSC, FLSR, and FTRR⁴ respectively in Fig. 4. As we can see the aggregated feature representations obtained by AGCSC display clear cluster structure, while for

⁴The rest algorithms use original feature representations of samples.

the other algorithms, the aggregated feature representations come different subspaces are overlap.

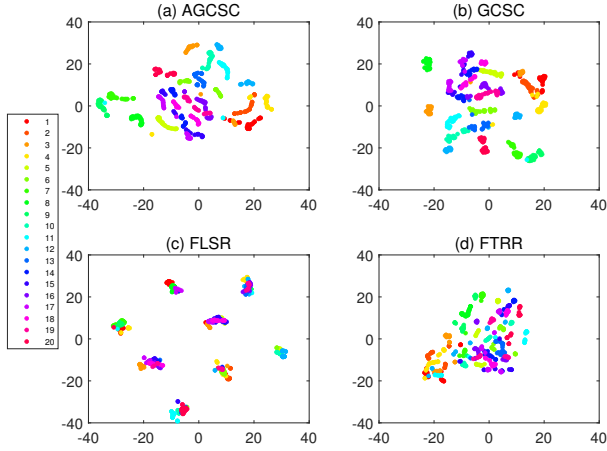


Figure 4. The visualizations of aggregated feature representations obtained by (a) AGCSC, (b) GCSC, (c) FLSR and (d) FTRR on Umist data set.

4.5. Parameter analysis

AGCSC has two parameters α and β . We show the effects of α and β on the clustering performance of AGCSC. Fig. 5 and Fig. 6 show the clustering accuracies and normalized mutual information obtained by AGCSC varied with the parameters respectively. It is clearly that on all the data sets, AGCSC achieves better results when α and β take relatively small values.

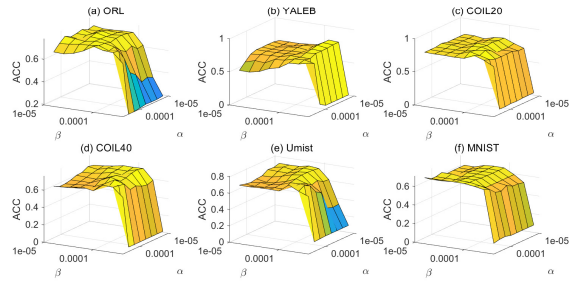


Figure 5. The influence of parameters α and β on clustering accuracy of AGCSC.

TAGCSC has an additional parameter, i.e., the thresholding value m . For a certain value of m , we record clustering results of TAGCSC on each pair of (α, β) . Then the best result could be determined for a fixed m . Then we let m vary from 4 to 15 and illustrate the clustering performance of TAGCSC in Fig. 7. It can be found that TAGCSC get better results when m is relatively small on all data sets except MNIST. And the performance of TAGCSC is much stable

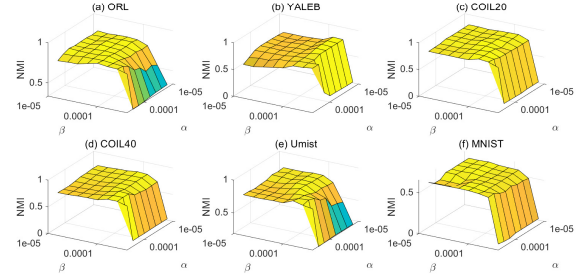


Figure 6. The influence of parameters α and β on normalized mutual information of AGCSC.

on MNIST dataset. Based on these results, we suggest m to be a relative small value (i.e., $m \leq 8$) for subspace clustering tasks.

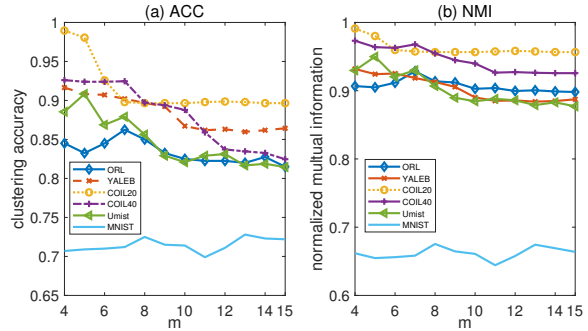


Figure 7. The influence of thresholding value m on the clustering performance of TAGCSC.

4.6. Convergence and complexity analysis

Firstly, we discuss the convergence of AGCSC. Take ORL as an example, we record the computed variables $\mathbf{F}, \mathbf{C}, \mathbf{Z}$ of Algorithm 1 in each iteration. Then the residuals of three variables (i.e., $\|\mathbf{F}_{t+1} - \mathbf{F}_t\|_F^2, \|\mathbf{C}_{t+1} - \mathbf{C}_t\|_F^2$ and $\|\mathbf{Z}_{t+1} - \mathbf{Z}_t\|_F^2$) could be obtained. The residuals versus the number of iterations are illustrated in Fig. 8. We can see that after about 250 iterations, Algorithm 1 converges into a stable solution.

Secondly, we compare the complexity of AGCSC with the other evaluated algorithms. Because LSR, LRSC, and TRR have closed-form solutions, we do not consider the consumption time of these algorithms. Then the average computation time (seconds) of the rest algorithms to run on different datasets is shown in Fig. 9. We can see that 1) the computation time of AGCSC is less than those of FLSR and FTRR on all the data sets except COIL40; 2) the complexity of AGCSC is in the same order as the time complexity of the other evaluated algorithms.

Table 3. Clustering results (in %) of AGCSC and TAGCSC compared with several deep clustering methods. The best results are emphasized in bold. Some results are ignored because the results are not found in corresponding literatures.

Dataset	Metric	Method								
		AE+SSC	DSC-L1	DSC-L2	DEC	DKM	DCCM	PSSC	AGCSC	TAGCSC
ORL	ACC	75.63	85.50	86.00	51.75	46.82	62.50	86.75	80.50	86.25
	NMI	85.55	90.23	90.34	74.49	73.32	79.06	93.49	88.51	92.84
YALEB	ACC	74.80	96.80	97.33	86.84	-	-	-	84.79	92.32
	NMI	78.33	96.87	97.03	92.40	-	-	-	87.34	94.04
Umist	ACC	70.42	72.42	73.12	55.21	51.06	54.48	79.17	81.04	90.83
	NMI	75.15	75.56	76.62	71.25	72.49	74.40	86.70	87.49	94.99
COIL20	ACC	87.11	93.14	93.68	72.15	66.51	80.21	97.22	88.75	98.96
	NMI	89.90	93.53	94.08	80.07	79.71	86.39	97.79	93.38	99.11
COIL40	ACC	73.91	80.03	80.75	48.72	58.12	76.91	83.58	78.12	92.60
	NMI	83.18	88.52	89.41	74.17	78.40	88.90	92.58	89.21	97.23
MNIST	ACC	48.40	72.80	75.00	61.20	53.32	40.20	84.30	71.40	72.80
	NMI	53.37	72.17	73.19	57.43	50.02	34.68	76.76	65.84	67.54

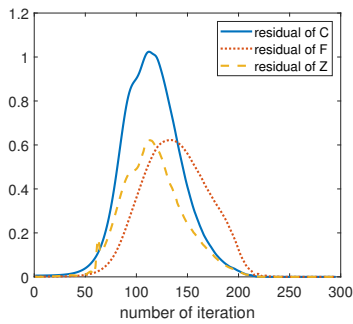


Figure 8. The residuals of variables F , C , Z versus the iterations on ORL database.

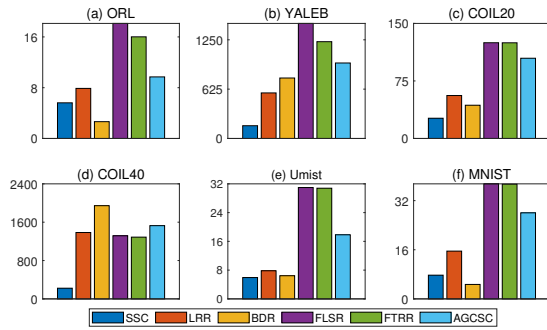


Figure 9. Computation time (seconds) for different algorithms to run on different datasets. The y-axis coordinates are in seconds.

4.7. Comparison with deep clustering methods

Due to the promising results achieved by deep clustering models, we here compare AGCSC and TAGCSC with some deep clustering algorithms including SSC with pre-trained convolutional auto-encoder features (AE+SSC), deep subspace clustering network (DSC) [10], deep embedding clustering (DEC) [43], deep K -means (DKM) [7], deep comprehensive correlation mining (DCCM) [41] and pseudo-supervised deep subspace clustering (PSSC) [22]. The ex-

perimental results of the deep models are cited from some latest literatures [10, 21, 24, 30]. Table 3 shows the comparison results.

We can observe from Table 3 that 1) TAGCSC outperforms the deep models on Umist, COIL20 and COIL40 datasets, and is close to DSC on MNIST datasets. This shows the effectiveness of TAGCSC. Considering the complexity and the unexplained characteristics of DNNs, our proposed algorithm is much appealing; 2) TAGCSC is inferior to DSC-L1 and DSC-L2 on YALEB dataset. However, as pointed in [8], the success of DSC-L1 and DSC-L2 may be attributed to post-processing method. Without post-processing, DSC can only achieve 59.09% clustering accuracy on YALEB. This result is much worse than those of AGCSC and TAGCSC. Hence, we can say that our proposed algorithm has a comparable performance to those of the deep models.

5. Conclusions

In this paper, we develop a new subspace clustering algorithm, named adaptive graph convolutional subspace clustering (AGCSC). In AGCSC, we propose to use the reconstruction coefficient matrix to design a graph convolutional operator. Then use the graph convolutional operator to smooth the feature representations and the coefficient matrix simultaneously. We demonstrate plenty of subspace clustering experiments to show the superiorities of feature representations and coefficient matrix obtained by AGCSC. And we claim that AGCSC and its extension (TAGCSC) are comparable to several deep clustering models.

6. Acknowledge

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