Binhuan Wang, Yilong Zhang, Will Wei Sun, Yixin Fang

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Outline

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- Theoretical Analysis
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Cluster Analysis

Cluster analysis aims to assign observations into a number of clusters such that observations in the same group are similar to each other.

Traditional clustering methods:

- K-means.
- Hierarchical clustering.
- Gaussian mixture models.

However, these methods suffer from instabilities due to their non-convex optimization formulations.

Convex Clustering

Convex Clustering [Lindsten et al., 2011, Hocking et al., 2011]:

$$\min_{\mathbf{A} \in \mathbb{R}^{n \times p}} \frac{1}{2} \sum_{i=1}^{n} \left\| X_i - A_{i.} \right\|_2^2 + \gamma \sum_{i_1 < i_2} \left\| A_{i_1.} - A_{i_2.} \right\|_q$$

where $X \in \mathbb{R}^{n \times p}$, $A_{i.}$ is the *i*-th row of **A** and $\|\cdot\|_q$ is the L_q -norm of a vector with $q \in \{1, 2, \infty\}$.

- K-means clustering and hierarchical clustering consider L_0 -norm in the second term, which leads to a non-convex optimization problem.
- Small γ (e.g. $\gamma = 0$) makes each observation by itself is a cluster.
- Large γ (e.g. $\gamma = \infty$) makes all the row of \hat{A} be identical.

Convex Clustering

- In recent years, much effort has been spent on developing algorithms and theory for convex clustering [Chi and Lange, 2015, Tan and Witten, 2015].
- When the number of features becomes large, many of them may contain no information. Thus the performance of these methods can be severely deteriorated.
- To overcome this problem, an algorithm that can simultaneously perform cluster analysis and select informative variables is in demand.

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$$\min_{\mathbf{A} \in \mathbb{R}^{n \times p}} \frac{1}{2} \sum_{i=1}^{n} \|X_{i.} - A_{i.}\|_{2}^{2} + \gamma \sum_{i_{1} < i_{2}} w_{i_{1}, i_{2}} \|A_{i_{1}.} - A_{i_{2}.}\|_{q}$$

$$\tag{1}$$

where the weight $w_{i_1,i_2} \ge 0$.

- [Hocking et al., 2011] considered a pairwise affinity weight $w_{i_1,i_2} = \exp\left(-\phi \left\|X_{i_1} X_{i_2}\right\|_2^2\right)$.
- [Chi and Lange, 2015] suggested $w_{i_1,i_2} = \iota_{i_1,i_2}^m \exp\left(-\phi \left\|X_{i_1} X_{i_2}\right\|_2^2\right)$ where ι_{i_1,i_2}^m is 1 if observation i_2 is among i_1 's m nearest neighbors or vice verse, and 0 otherwise.

Write the data matrix X in feature-level as column vector $\mathbf{X} = (\mathbf{x}_1, \cdots, \mathbf{x}_p)$, where $\mathbf{x}_j = \begin{pmatrix} X_{1j}, \cdots, X_{nj} \end{pmatrix}^T$, $j = 1, \ldots, p$ and denote \mathbf{A} in feature level as column vector $\mathbf{A} = (\mathbf{a}_1, \cdots, \mathbf{a}_p)$. Simple algebra implies that (1) can be reformulated as

$$\min_{\mathbf{A} \in \mathbb{R}^{n \times p}} \frac{1}{2} \sum_{j=1}^{p} \|\mathbf{x}_{j} - \mathbf{a}_{j}\|_{2}^{2} + \gamma \sum_{l \in \mathscr{E}} w_{l} \|A_{i_{1}} - A_{i_{2}}\|_{q}$$
 (2)

where $\mathscr{E} = \{l = (i_1, i_2) : 1 \le i_1 < i_2 \le n\}$.

- Without loss of generality, we assume the feature vectors are centered, i.e., $\sum_{i=1}^{n} X_{ij} = 0$ for each j = 1, ..., p.
- When $\hat{\mathbf{a}}_j$ are identical, when the corresponding feature j is not informative for clustering, i.e., $\|\hat{\mathbf{a}}_j\|_2^2 = 0$.

Sparse convex clustering solves

$$\min_{\mathbf{A} \in \mathbb{R}^{n \times p}} \frac{1}{2} \sum_{j=1}^{p} \|\mathbf{x}_{j} - \mathbf{a}_{j}\|_{2}^{2} + \gamma_{1} \sum_{l \in \mathcal{E}} w_{l} \|A_{i_{1}} \cdot -A_{i_{2}} \cdot \|_{q} + \gamma_{2} \sum_{j=1}^{p} u_{j} \|\mathbf{a}_{j}\|_{2}$$
(3)

where tuning parameter γ_1 controls the cluster size and tuning parameter γ_2 controls the number of informative features.

• In the group-lasso penalty, the weight u_j plays an important role to adaptively penalize the features.

Algorithm

Two optimization approaches similar to [Chi and Lange, 2015].

- Sparse alternating direction method of multipliers (S-ADMM).
- Sparse alternating minimization algorithm (S-AMA).

Equivalent Form

This is equivalent to minimize the augmented Lagrangian function,

$$L_{\mathbf{v}}(\mathbf{A}, \mathbf{V}, \mathbf{\Lambda}) = \frac{1}{2} \sum_{j=1}^{p} \|\mathbf{x}_{j} - \mathbf{a}_{j}\|_{2}^{2} + \gamma_{1} \sum_{l \in \mathscr{E}} w_{l} \|\mathbf{v}_{l}\|_{q} + \gamma_{2} \sum_{j=1}^{p} u_{i} \|\mathbf{a}_{j}\|_{2}$$
$$+ \sum_{l \in \mathscr{E}} \langle \lambda_{l}, \mathbf{v}_{l} - A_{i_{1}} \cdot + A_{i_{2}} \rangle + \frac{v}{2} \sum_{l \in \mathscr{E}} \|\mathbf{v}_{l} - A_{i_{1}} \cdot + A_{i_{2}}\|_{2}^{2}$$

where v is a small constant, $\mathbf{V} = \left(\mathbf{v}_1, \dots, \mathbf{v}_{|\mathscr{E}|}\right)$, and $\mathbf{\Lambda} = \left(\lambda_1, \dots, \lambda_{|\mathscr{E}|}\right)$.

S-ADMM

S-ADMM solves

$$\begin{split} \mathbf{A}^{m+1} &= \underset{\mathbf{A}}{\operatorname{argmin}} L_{\mathcal{V}}\left(\mathbf{A}, \mathbf{V}^{m}, \boldsymbol{\Lambda}^{m}\right), \\ \mathbf{V}^{m+1} &= \underset{\mathbf{V}}{\operatorname{argmin}} L_{\mathcal{V}}\left(\mathbf{A}^{m+1}, \mathbf{V}, \boldsymbol{\Lambda}^{m}\right), \\ \mathbf{\lambda}_{l}^{m+1} &= \mathbf{\lambda}_{l}^{m} + \mathcal{V}\left(\mathbf{v}_{l}^{m+1} - A_{l_{1}\cdot}^{m+1} + A_{l_{2}\cdot}^{m+1}\right), l \in \mathcal{E}. \end{split}$$

Step 1 : Update *A* Denote $\tilde{\mathbf{v}}_l = \mathbf{v}_l + \frac{1}{v} \boldsymbol{\lambda}_l$. Updating A is equivalent to minimizing

$$f(\mathbf{A}) = \frac{1}{2} \sum_{j=1}^{p} \|\mathbf{x}_{j} - \mathbf{a}_{j}\|_{2}^{2} + \frac{\mathbf{v}}{2} \sum_{l \in \mathscr{E}} \|\widetilde{\mathbf{v}}_{l} - A_{i_{1}.} + A_{i_{2}.}\|_{2}^{2} + \gamma_{2} \sum_{j=1}^{p} u_{j} \|\mathbf{a}_{j}\|_{2}$$
(4)

This optimization problem is challenging because the objective function involves both rows and columns of the matrix A.



Lemma 1

Let \mathbf{I}_n be an $n \times n$ identity matrix, $\mathbf{1}_n \in \mathbb{R}^n$ be a vector with each component being 1, and \mathbf{e}_i be a vector with each component being 0 but its i-th component being 1. Define $\mathbf{N}^{-1} = (1+nv)^{-1/2} \left[\mathbf{I}_n + n^{-1} (\sqrt{1+nv} - 1) \mathbf{1}_n \mathbf{1}_n^T \right]$ and denote $\mathbf{y}_j = \mathbf{N}^{-1} \left[\mathbf{x}_j + \ v \sum_{l \in \mathscr{E}} \widetilde{v}_{jl} \left(\mathbf{e}_{i_1} - \mathbf{e}_{i_2} \right) \right]$ with \widetilde{v}_{jl} the j-th element of $\widetilde{\mathbf{v}}_l$. Then, minimizing (4) is equivalent to

$$\min_{\mathbf{a}_{j}} \frac{1}{2} \|\mathbf{y}_{j} - \mathbf{N}\mathbf{a}_{j}\|_{2}^{2} + \gamma_{2}u_{j} \|\mathbf{a}_{j}\|_{2}, \text{ for each } j = 1, \dots, p$$

remark: Based on this property, we are able to solve the minimization of f(A) by p separate sub-optimization problems.

S-ADMM

Step 2 : Update V For any $\sigma > 0$ and norm $\Omega(\cdot)$, we define a proximal map,

$$\text{prox}_{\sigma\Omega}(\mathbf{u}) = \underset{\mathbf{v}}{\text{argmin}} \left[\sigma\Omega(\mathbf{v}) + \frac{1}{2}\|\mathbf{u} - \mathbf{v}\|_2^2 \right]$$

In S-ADMM, $\Omega(\cdot)$ is a q-norm $\|\cdot\|_q$ with q=1,2, or ∞ , and $\sigma=\gamma_1w_l/\nu$. Because vectors \mathbf{v}_l are separable, they can be solved via proximal maps, that is

$$\mathbf{v}_{l} = \underset{\mathbf{v}_{l}}{\operatorname{argmin}} \frac{1}{2} \left\| \mathbf{v}_{l} - \left(A_{i_{1}} - A_{i_{2}} - \mathbf{v}^{-1} \boldsymbol{\lambda}_{l} \right) \right\|_{2}^{2} + \frac{\gamma_{1} w_{l}}{\mathbf{v}} \left\| \mathbf{v}_{l} \right\|_{q}$$
$$= \operatorname{prox}_{\sigma_{l} \left\| \cdot \right\|_{q}} \left(A_{i_{1}} - A_{i_{2}} - \mathbf{v}^{-1} \boldsymbol{\lambda}_{l} \right)$$

Step 2 : Update $\Lambda \lambda_l$ can be updated by $\lambda_l = \lambda_l + v(\mathbf{v}_l - A_{i_1} + A_{i_2})$.

S-ADMM

- 1 Initialize V^0 and Λ^0 . For m = 1, 2, ...
- **2** For j = 1, ..., p, do

$$\begin{split} \widetilde{\mathbf{v}}_{l}^{m-1} &= \mathbf{v}_{l}^{m-1} + \frac{1}{\nu} \boldsymbol{\lambda}_{l}^{m-1}, l \in \mathscr{E} \\ \mathbf{y}_{j}^{m-1} &= \mathbf{N}^{-1} \left(\mathbf{x}_{j} + \nu \sum_{l \in \mathscr{E}} \widetilde{\nu}_{lj}^{m-1} \left(\mathbf{e}_{i_{1}} - \mathbf{e}_{i_{2}} \right) \right) \\ \mathbf{a}_{j}^{m} &= \operatorname*{argmin}_{\mathbf{a}_{j}} \frac{1}{2} \left\| \mathbf{y}_{j}^{m-1} - \mathbf{N} \mathbf{a}_{j} \right\|_{2}^{2} + \gamma_{2} u_{j} \left\| \mathbf{a}_{j} \right\|_{2} \\ \mathbf{a}_{j}^{m} &= \mathbf{a}_{j}^{m} - \overline{\mathbf{a}}_{j}^{m} \mathbf{1}_{n}, \text{ where } \overline{\mathbf{a}}_{j}^{m} = \mathbf{1}_{n}^{T} \mathbf{a}_{j}^{m} / n \end{split}$$

3 For $l \in \mathcal{E}$, do

$$\mathbf{v}_l^m = \operatorname{prox}_{\sigma_l \| \cdot \|_q} \left(A_{i_1.}^m - A_{i_2.}^m - \mathbf{v}^{-1} \mathbf{\lambda}_l^{m-1} \right)$$

4 For $l \in \mathcal{E}$, do

$$\boldsymbol{\lambda}_{l}^{m} = \boldsymbol{\lambda}_{l}^{m-1} + v \left(\mathbf{v}_{l}^{m} - A_{i_{1}}^{m} + A_{i_{2}}^{m} \right)$$

5 Repeat Steps 2-4 until convergence.



S-AMA

S-AMA aims to increase the computational efficiency of S-ADMM.

• S-AMA solves A by treating v=0, i.e., $\mathbf{A}^{m+1}=\operatorname{argmin}_{\mathbf{A}}L_0(\mathbf{A},\mathbf{V}^m,\mathbf{\Lambda}^m)$. When v=0, we have $N=\mathbf{I_n}$ and $y_j=x_j$. According to Lemma 1, updating A requires to solve p group-lasso problems:

$$\min_{\mathbf{a}_{j}} \frac{1}{2} \|\mathbf{x}_{j} - \mathbf{a}_{j}\|_{2}^{2} + \gamma_{2} u_{j} \|\mathbf{a}_{j}\|_{2}, j = 1, \dots, p$$
 (5)

By Karush-Kuhn-Tucker (KKT) conditions of the group lasso problem [Yuan and Lin, 2006], the solution to (5) has a closed form as

$$\widehat{\mathbf{a}}_{j} = \left(1 - \frac{\gamma_{2} u_{j}}{\|\mathbf{z}_{j}\|_{2}}\right)_{+} \mathbf{z}_{j}$$

where $\mathbf{z}_j = \mathbf{x}_j + \sum_{l \in \mathscr{E}} \lambda_{jl} (\mathbf{e}_{i_1} - \mathbf{e}_{i_2})$ and $(z)_+ = \max\{0, z\}$.

• S-AMA does not need to update *V*.



S-AMA

- 1 Initialize Λ^0 . For m = 1, 2, ...
- **2** For j = 1, ..., p, do

$$\begin{split} \mathbf{z}_{j}^{m} &= \mathbf{x}_{j} + \sum_{l \in \mathscr{E}} \lambda_{lj}^{m-1} \left(\mathbf{e}_{i_{1}} - \mathbf{e}_{i_{2}} \right) \\ \mathbf{a}_{j}^{m} &= \left(1 - \frac{\gamma_{2} u_{i}}{\left\| \mathbf{z}_{i}^{m} \right\|_{2}} \right)_{+} \mathbf{z}_{j}^{m} \\ \mathbf{a}_{j}^{m} &= \mathbf{a}_{j}^{m} - \overline{\mathbf{a}}_{j}^{m} \mathbf{1}_{n}, \text{ where } \overline{\mathbf{a}}_{j}^{m} = \mathbf{1}_{n}^{T} \mathbf{a}_{j}^{m} / n \end{split}$$

3 For $l \in \mathscr{E}$, do

$$\boldsymbol{\lambda}_{l}^{m} = P_{C_{l}} \left[\boldsymbol{\lambda}_{l}^{m-1} - v \left(A_{i_{1}}^{m} - A_{i_{2}}^{m} \right) \right]$$

where
$$C_l = \left\{ \boldsymbol{\lambda}_l : \|\boldsymbol{\lambda}_l\|_{\dagger} \leqslant \gamma_1 w_l \right\}$$

4 Repeat Steps 2-3 until convergence.

remark: $P_{C_l}(\cdot)$ denotes projection onto C_l , and $\|\cdot\|_{\dagger}$ denotes the dual norm.

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Some Notations

- Assume $\mathbf{x} = \mathbf{a}_0 + \boldsymbol{\varepsilon}$, where $\boldsymbol{\varepsilon} \in \mathbb{R}^{np}$ is a vector of independent sub-Gaussian noise terms with mean zero and variance σ^2 , and $\mathbf{a}_0 = \left(\mathbf{a}_{01}^{\mathrm{T}}, \dots, \mathbf{a}_{0p}^{\mathrm{T}}\right)^{\mathrm{T}}$ is a np-dimensional mean vector.
- Assume that only the first $p_0 < p$ features are informative, i.e., $\|\mathbf{a}_{0j}\|_2 \neq 0$ for $j \leq p_0$ and $\|\mathbf{a}_{0j}\|_2 = 0$ for $j > p_0$. The informative feature set is denoted as $A = \{1, \ldots, p_0\}$ and the noninformative feature set is $A^c = \{p_0 + 1, \ldots, p\}$. For simplicity, we consider the case with $w_l = 1$.
- Sparse convex clustering in (3) can be reformulated as the following problem:

$$\widehat{\mathbf{a}} = \underset{\mathbf{a} \in \mathbb{R}^{np}}{\operatorname{argmin}} \frac{1}{2} \|\mathbf{x} - \mathbf{a}\|_{2}^{2} + \gamma_{1} \sum_{l \in \mathscr{E}} \|\mathbf{C}_{l} \mathbf{a}\|_{q} + \gamma_{2} \sum_{j=1}^{p} u_{j} \|\mathbf{a}_{j}\|_{2}$$
(6)

where $\mathbf{C}_l = \mathbf{I}_p \otimes (\mathbf{e}_{i_1} - \mathbf{e}_{i_2})^{\mathrm{T}}$ and hence $\mathbf{C}_l \mathbf{a} = A_{i_1} - A_{i_2..}$ Define $\mathbf{C} = \left(\mathbf{C}_1^{\mathrm{T}}, \dots, \mathbf{C}_{|\mathscr{E}|}^{\mathrm{T}}\right)^{\mathrm{T}}$ and denote $\mathbf{u} = (u_1, \dots, u_p)^{\mathrm{T}}$.



Prediction error for q = 1

Theorem 1

Let $\hat{\mathbf{a}}$ be the estimate of (6) with q=1. If $\gamma_1>4\sigma\sqrt{\frac{\log\left(p\cdot\left(\begin{array}{c}n\\2\end{array}\right)\right)}{n}}$, then

$$\frac{1 - \gamma_2}{2np} \|\widehat{\mathbf{a}} - \mathbf{a}_0\|_2^2 \leqslant \frac{3\gamma_1}{2np} \|\mathbf{C}\mathbf{a}_0\|_1 + \frac{\gamma_2 \|\mathbf{u}\|_2^2}{2np} + \sigma^2 \left[\frac{1}{n} + \sqrt{\frac{\log(np)}{n^2p}} \right] + \frac{1}{np}$$

holds with probability at least $1-c_3$, where

$$c_3 = \frac{2}{p \cdot \binom{n}{2}} + \exp\left\{-\min\left(c_1\log(np), c_2\sqrt{p\log(np)}\right)\right\} + 2\exp\left(-\frac{np}{\left(2\sigma^2\gamma_2^2\|\mathbf{u}\|_1^2\right)}\right)$$

for some positive constants c_1 and c_2 defined in Lemma <u>S.1</u>

Prediction error for q = 2

Theorem 2

Let $\widehat{\mathbf{a}}$ be the estimate of (6) with q=2. If $\gamma_1>4\sigma\sqrt{\frac{\log\left(p\cdot\left(\begin{array}{c}n\\2\end{array}\right)\right)}{n}}$, then

$$\frac{1 - \gamma_2}{2np} \|\widehat{\mathbf{a}} - \mathbf{a}_0\|_2^2 \leqslant \frac{3\gamma_1}{2np} \sum_{l \in \mathscr{E}} \|\mathbf{C}_l \mathbf{a}_0\|_2 + \frac{\gamma_2 \|\mathbf{u}\|_2^2}{2np} + \sigma^2 \left[\frac{1}{n} + \sqrt{\frac{\log(np)}{n^2p}} \right] + \frac{1}{np}$$

holds with probability at least $1-c_3$, where c_3 is defined in Theorem 1.

Theorem 3

Theorem 3

If
$$\gamma_1 > 4\sigma \sqrt{\log\left(p\cdot \left(\begin{array}{c} n\\ 2 \end{array}\right)\right)/n}$$
, $\gamma_1 \left\|\operatorname{Ca}_0\right\|_1/(2np) = o(1)$, $\gamma_2 \to 0$ and $\gamma_2 \left\|\mathbf{u}\right\|_1^2/(np) \to 0$ as $n,p\to\infty$, then $P\left(\left\|\widehat{\mathbf{a}}_j\right\|_2 = 0\right) \to 1$ for any $j\in A^c$, with the solution $\widehat{\mathbf{a}}$ to (6) with either $q=1$ or $q=2$.

Remark: $\chi \|\mathbf{u}\|_1^2/(np) \to 0$ generally implies that the adaptive weights cannot be too large. For example, uniform weights satisfy this condition.

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Selection of weights

• Following [Chi and Lange, 2015], we choose weights by incorporating the m-nearest-neighbors methods with Gaussian kernel. In specific, the weight between the pair (i_1, i_2) is

$$w_{i_1,i_2} = \iota_{i_1,i_2}^m \exp\left(-\phi \|X_{i_1} \cdot -X_{i_2} \cdot\|_2^2\right),$$

where t_{i_1,i_2}^m equals 1 if observation i_2 is among observation i_1 's m nearest neighbors. In application, we set m=5 and $\phi=0.5$.

• μ_j can be chosen as $1/\|\hat{\mathbf{a}}_j^0\|_2$, where $\|\hat{\mathbf{a}}_j^0\|_2$ is the estimate of \mathbf{a}_j in (3) with $\gamma_2=0$.

Selection of Tuning Parameters

- γ₁ controls the number of estimated clusters.
- γ controls the number of selected informative features.
- Use stability selection to tune both γ_1 and γ_2 :
 - For any given γ_1 and γ_2 , based on two sets of bootstrapped samples, two clustering results can be produced by (3).
 - Compute the stability measurement [Fang and Wang, 2012] to measure the agreement between the two clustering result.
 - Repeat this procedure 50 times and then compute the averaged stability selection method.
- To speed up tunning process, stability path can be computed over of a coarse grid of γ₁ and a fine grid of γ₂.

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Case One

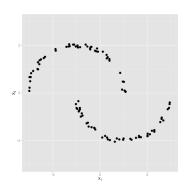
- Sample size n = 60 with the number of clusters either K = 2 or 4.
- The number of features either p = 150 or 500.
- For each i = 1, ..., n, cluster label Z_i is uniformly sampled from $\{1, ..., K\}$.
- The first 20 informative features are generated from $MVN_p(\mu_K(Z_i), \mathbf{I}_{20})$, where $\mu_K(Z_i)$ is defined as:
 - If K = 2, $\mu_2(Z_i) = \mu \mathbf{1}_{20} I(Z_i = 1) \mu \mathbf{1}_{20} I(Z_i = 2)$.
 - If K = 4, $\boldsymbol{\mu}_4(Z_i) = (\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}}, -\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}})^{\mathrm{T}} I(Z_i = 1) + (-\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}}, -\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}})^{\mathrm{T}} I(Z_i = 2) + (-\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}}, \boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}})^{\mathrm{T}} I(Z_i = 3) + (\boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}}, \boldsymbol{\mu} \mathbf{1}_{10}^{\mathrm{T}})^{\mathrm{T}} I(Z_i = 4)$.
- The rest p-20 noise features are generated from N(0,1).

Remark : μ controls the distance between cluster centers. A large μ indicates that clusters are well-separated, whereas a small μ indicates that clusters are overlapped.



Case Two

- n = 100, K = 2 and p = 40, where the first two features are informative, and the rest 38 noisy features are generated from N(0,0.5).
- The plot of the first two features for one example of two interlocking half moons.



Five Settings for Simulation

- Spherical settings
 - Setting 1 : $K = 2, n = 60, p = 150, \mu = 0.6$.
 - Setting 2: $K = 2, n = 60, p = 500, \mu = 0.7$.
 - Setting 3 : $K = 4, n = 60, p = 150, \mu = 0.9$.
 - Setting 4 : $K = 4, n = 60, p = 500, \mu = 1.2$.
- Non-spherical settings
 - Setting 5 : K = 2, n = 100, p = 40.

Evaluation Criteria

- RAND index : RAND index ranges between 0 and 1, and a higher value indicates better performance.
- False Negative Ratio (FNR).
- False Positive Ratio (FPR).

Due to the high computational burden for S-ADMM in high-dimensional settings, S-ADMM is not evaluated for p=500. Additionally, we run 200 repetitions for each setting.

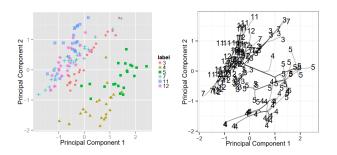
Results

		RAND		FNR		FPR	
Setting 1	k-means	0.95	0.06	0.00	0.00	1.00	0.00
	ADMM	0.53	0.39	0.00	0.00	1.00	0.00
	AMA	0.66	0.40	0.00	0.00	1.00	0.00
	S-ADMM	0.82	0.24	0.04	0.05	0.25	0.16
	S-AMA	0.96	0.06	0.03	0.07	0.30	0.21
Setting 2	k-means	0.95	0.11	0.00	0.00	1.00	0.00
	ADMM	0.14	0.20	0.00	0.00	1.00	0.00
	AMA	0.08	0.21	0.00	0.00	1.00	0.00
	S-AMA	0.97	0.07	0.07	0.09	0.11	0.10
Setting 3	k-means	0.83	0.15	0.00	0.00	1.00	0.00
	ADMM	0.56	0.22	0.00	0.00	1.00	0.00
	AMA	0.47	0.21	0.00	0.00	1.00	0.00
	S-ADMM	0.82	0.14	0.04	0.06	0.25	0.24
	S-AMA	0.84	0.13	0.02	0.04	0.11	0.18
Setting 4	k-means	0.89	0.14	0.00	0.00	1.00	0.00
_	ADMM	0.31	0.23	0.00	0.00	1.00	0.00
	AMA	0.31	0.20	0.00	0.00	1.00	0.00
	S-AMA	0.94	0.09	0.01	0.02	0.01	0.03
Setting 5	k-means	0.51	0.07	0.00	0.00	1.00	0.00
•	ADMM	0.54	0.08	0.00	0.00	1.00	0.00
	AMA	0.53	0.09	0.00	0.00	1.00	0.00
	S-AMA	0.57	0.07	0.00	0.00	0.34	0.27
	SPECC	0.52	0.08	0.00	0.00	1.00	0.00

- Convex clustering does not perform well when feature dimension is high.
- Sparse convex clustering selects informative features with great clustering accuracy.

Application: hand movement clustering

- Dataset contains 15 classes with each class referring to a hand movement type.
- Each class contains 24 observations and each observation has 90 features.



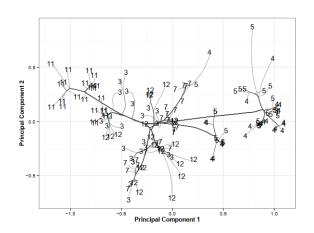
Convex clustering is only able to distinguish clusters 4 and 5 and treat the rest clusters as one class.

Results

Algorithm	# of clusters	# of features	RAND index
k-means	2	90	0.06
AMA	3	90	0.31
S-AMA	3	13	0.45

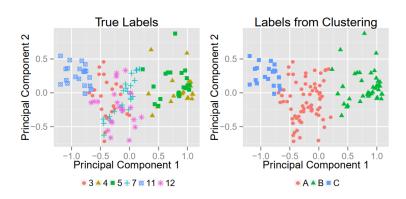
- Both convex clustering (AMA) and sparse convex clustering (S-AMA) perform better than k-means, which indicates that the performance of convex clustering or sparse convex clustering is less sensitive to the assumption of spherical clustering centers.
- By using only 13 informative features, our S-AMA is able to improve the clustering accuracy of convex clustering (AMA) by 45%. This indicates the importance of variable selection in high-dimensional clustering.

Clustering path of S-AMA



As tuning parameter γ_l increases, the clustering path of S-AMA tends to merge clusters 3, 7 and 12 into one big cluster, merge cluster 4 and 5 into another big cluster, and identify cluster 11 as the third cluster.

Results



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Conclusion and Future Work

- An extension of convex clustering, sparse convex clustering, is proposed to simultaneously cluster observations and conduct feature selection.
- The numerical results show that S-AMA is computationally faster and delivers better performance than S-ADMM.
- The numerical results show that the selection of tuning parameters in sparse convex clustering is important and the tuning method based on clustering stability performs well.
- Future work:
 - Extend convex bi-clustering [Chi et al., 2017] to sparse bi-clustering.
 - Use group L_0 penalty [Zhang et al., 2021] to replace the group lasso penalty applying on the feature level.



Any questions or comments?

References I



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