Online Convex Matrix Factorization with Representative Regions

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Key Proof Ideas

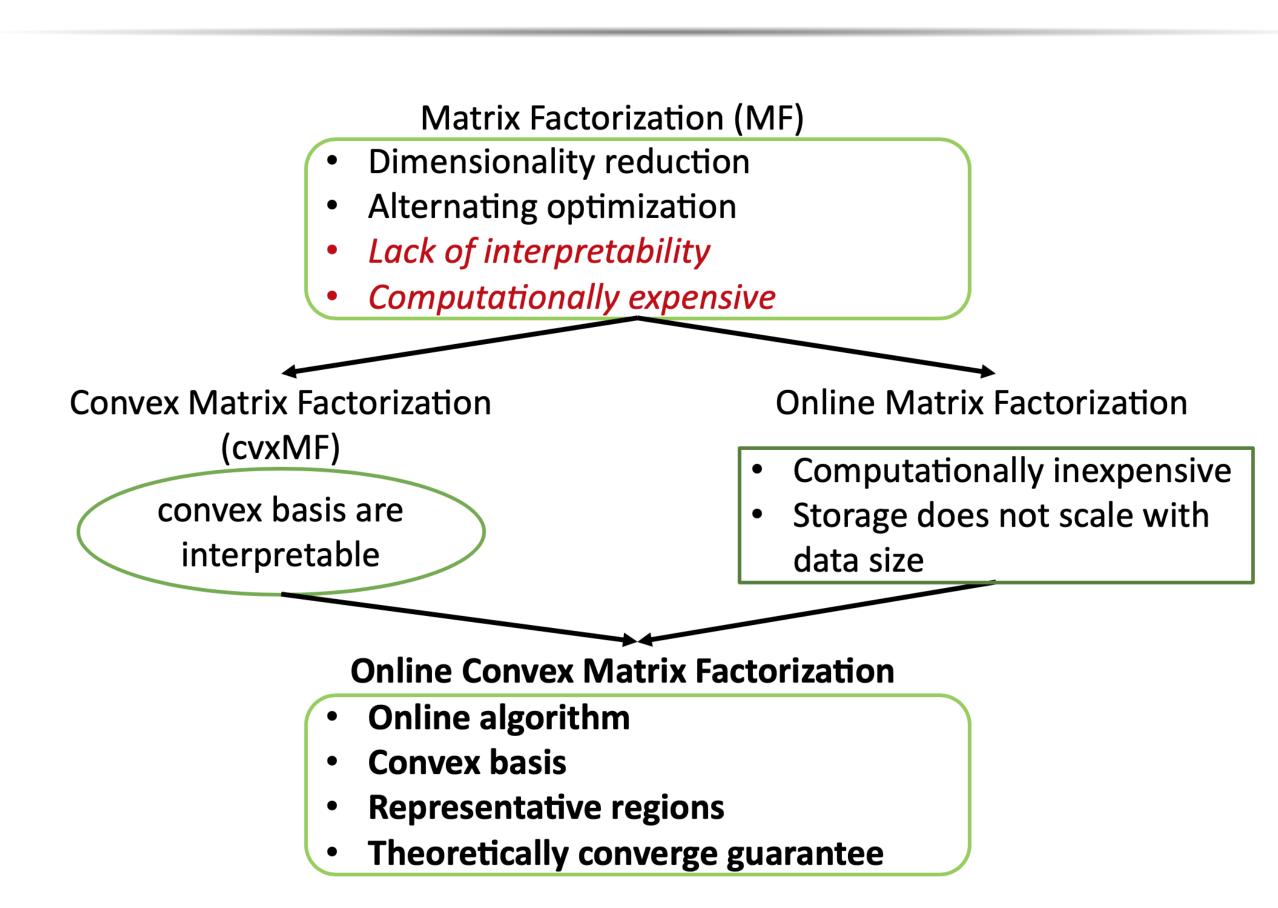
<u>Lemma 4</u>

 $\hat{g}_t(\boldsymbol{D}_t) - \hat{g}_t(\boldsymbol{D}_t^{\star}) \stackrel{a.s.}{\longrightarrow} 0$

 $\hat{g}_t(\boldsymbol{D}_t^{\star}) - g(\boldsymbol{D}_t^{\star}) \xrightarrow{a.s.} 0$

 $g(\boldsymbol{D}_t^{\star})$ converges a.s.

Matrix Factorization Problem



Stochastic Loss function:

$$g(\mathbf{D}) \triangleq \mathbb{E} \left| \min_{\alpha} \|X - \mathbf{D}\alpha\|^2 + \lambda \|\alpha\|_1 \right|$$

 $X \in \Omega \subset \mathbb{R}^m$: Random Variable input

 $\mathbf{D} \in \mathbb{R}^{m \times k}$: Basis Vector matrix

 $\alpha \in \mathbb{R}^k$: Coefficient vector corresponding to input X

 $\lambda \in \mathbb{R}$: Sparsity parameter

Approach and Challenges

Let $\ell(x_t, \mathbf{D}, \alpha) \triangleq ||x_t - \mathbf{D}\alpha||^2 + \lambda ||\alpha||_1$ be the loss for a single input x_t . Then, $g(\mathbf{D}) = \mathbb{E}[\min_{\alpha} \ell(X, \mathbf{D}, \alpha)] = \mathbb{E}[\ell(X, D)]$ An empirical approximation of g(D) can be defined as:

$$g_t(\mathbf{D}) = \sum_{n \le t} \min_{\alpha} ||x_t - \mathbf{D}\alpha||^2 + \lambda ||\alpha||_1, \mathbf{D} \in \Omega$$

Minimizing $g_t(D)$ using an online algorithm is infeasible without keeping all the history data.

• We choose a surrogate function

$$\hat{g}_t(\mathbf{D}) = \sum_{n \le t} ||x_t - \mathbf{D}\alpha_n||^2 + \lambda ||\alpha_n||_1, \mathbf{D} \in \Omega$$

- Even optimizing the surrogate function $\hat{g}_t(\mathbf{D})$ for $\mathbf{D} \in \Omega$ is infeasible for an online algorithm, since Ω is unknown
- Instead we compute $\min_{\mathbf{D} \in \text{cvx}(\hat{X}_t)} \hat{g}_t(\mathbf{D})$ in each step, while updating the region $\operatorname{cvx}(\hat{X}_t)$ in each iteration
- New objective: $\min_{\mathbf{D} \in \text{cvx}(\hat{X}_t)} \hat{g}_t(\mathbf{D})$

We prove that the results are asymptotically the same as optimizing for $g(D), \mathbf{D} \in \Omega$

Algorithm Setting

We consider two versions of the algorithm

- $\mathcal{R}_u : \mathbf{D}_j \in \Omega, \forall j \in [k]$
- $\mathcal{R}_r: \mathbf{D}_j \in \Omega_j, \forall j \in [k]$ where Ω_j corresponds to the cluster j in the data

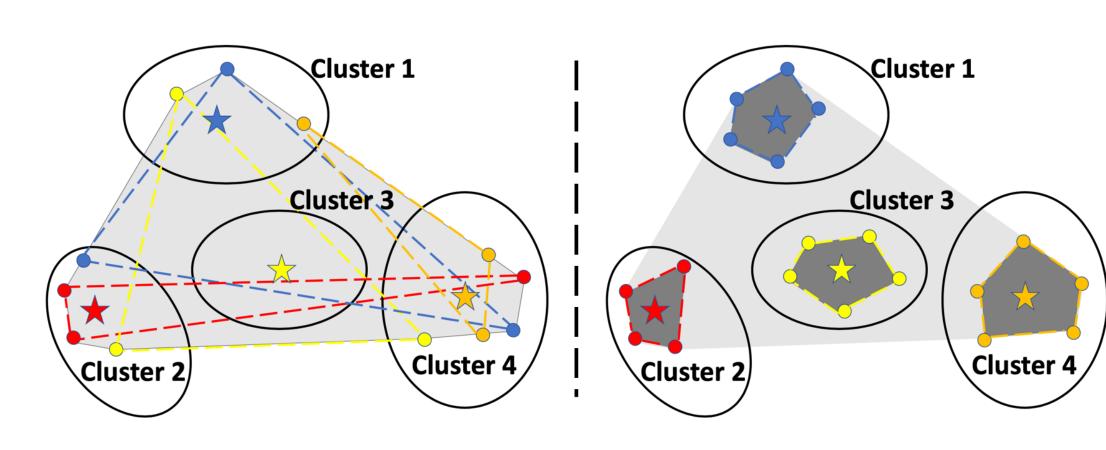


Figure 1:A multi-cluster dataset: Stars represent the learned bases, while circles denote representative samples for the basis of the same color. Left: The representative sets for the individual basis elements are unrestricted. Right: The representative sets are restricted to lie within their corresponding clusters.

Algorithm

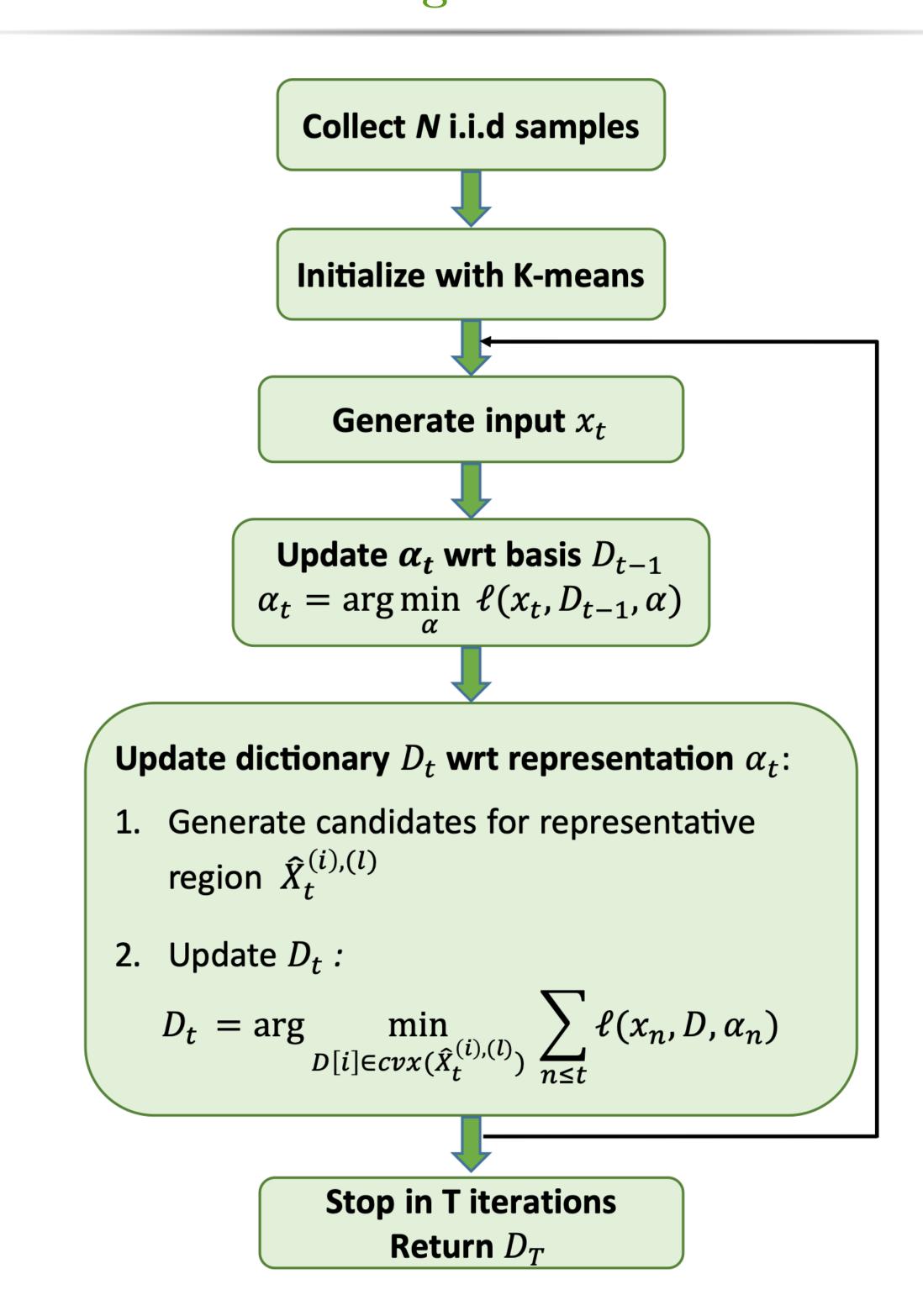


Figure 2:Flowchart of online cvxMF

Algorithm (cont.)

Figure 3 also shows one step in the online cvxMF algorithms for a three clusters toy dataset.

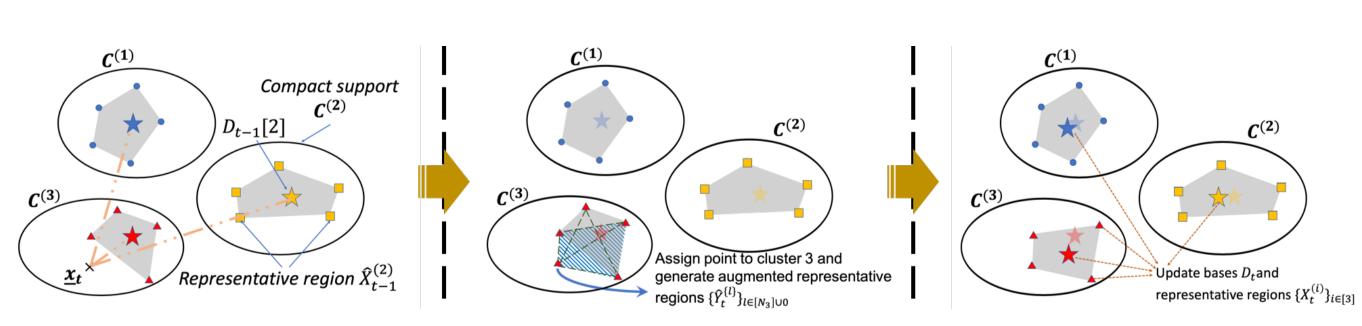


Figure 3:One step of the online cvxMF algorithm with multiple-representative regions.

Assumptions

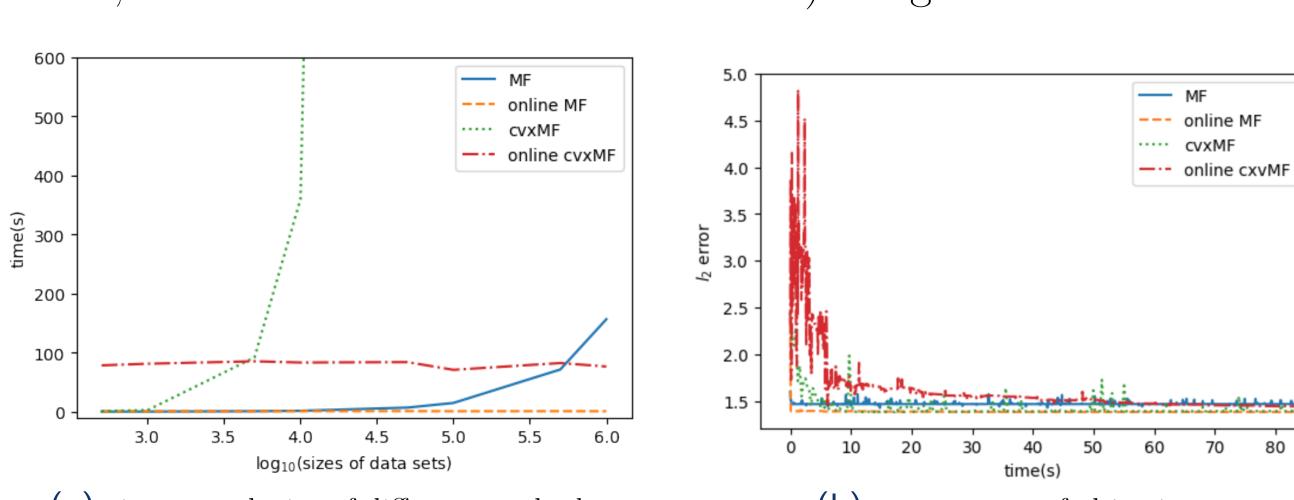
(A.1) The data distribution on a compact support set C has bounded "skewnes". The bounded skewness assumption for the distribution of X reads as

$$P(||\mathbf{X} - p|| \le r \mid \mathbf{X} \in \mathcal{C}) \ge \kappa \operatorname{vol}(B(r, p)) / \operatorname{vol}(\mathcal{C})$$
 (1)

- (A.2) The quadratic surrogate functions \hat{g}_t are strictly convex, and have Hessians that are lower-bounded by a positive constant $\kappa_1 >$
- (A.3) The approximation-error function $\ell(x,D)$ is "well**behaved".** We assume that $\ell(x,D)$ is continuously differentiable and its expectation g(D) is continuously differentiable and Lipschitz on the compact set ${\cal C}$

Experimental Validation (cont.)

we present running times and times to convergence (or, if convergence is slow, the maximum number of iterations) in figure 5.



(a) time complexity of different methods (b) convergence of objective Figure 5:(a): Running times (s) vs. the log of the dataset sizes; (b) Running times (s) vs. the l_2 error.

The MNIST Dataset:

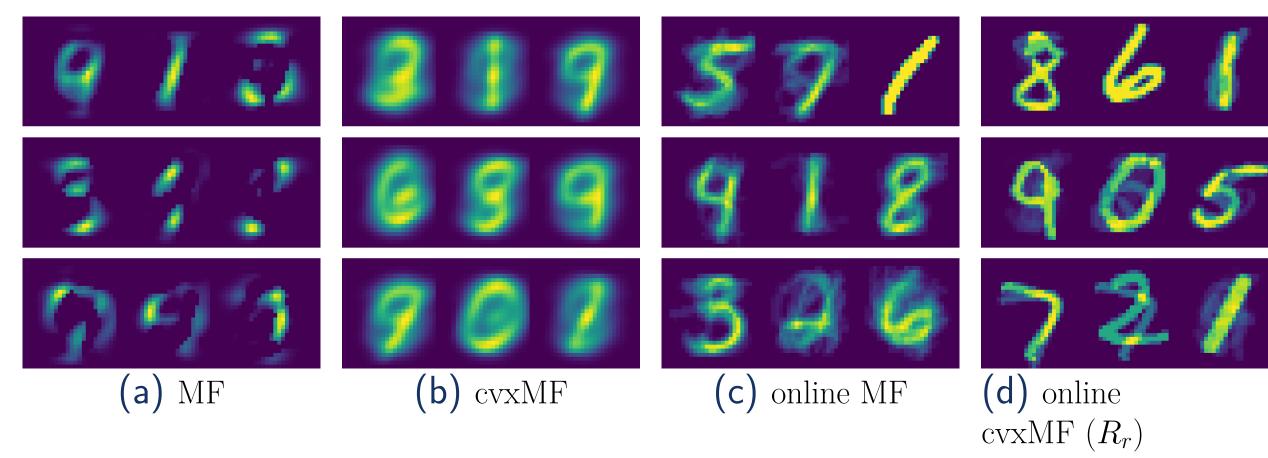


Figure 6:MNIST results (as the eigenimage set is overcomplete, clustering accuracy is omitted).

Single-Cell (sc) RNA Data:

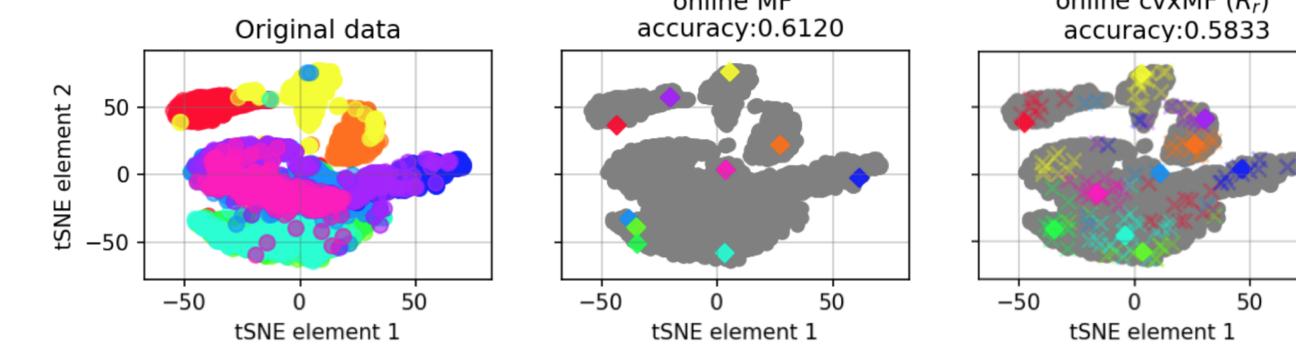
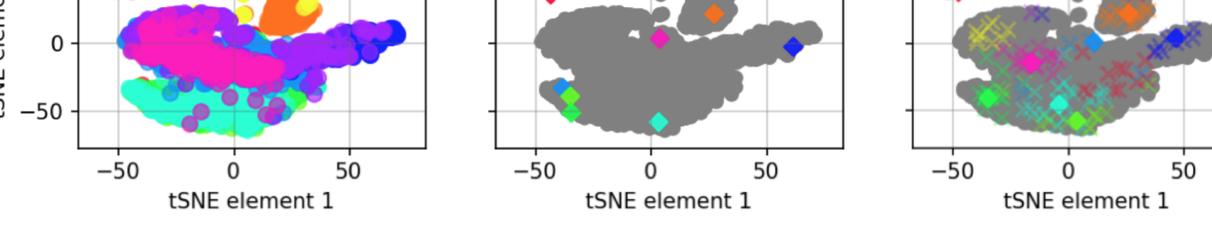


Figure 7:Results for the online methods executed on a blood-cell scRNA dataset.



Theorem 1 Under assumptions (A.1) to (A.3), the sequence $\{D_t\}_t$ converges almost surely to a stationary point of g(D)

Main Results

Lemma 2 Let $\Delta_t \triangleq \hat{g}_t(D_t) - \hat{g}_t(D_t^*)$. Then $\mathbb{E}[\Delta_t] \le O\left(\frac{1}{t}\right) + \mathbb{E}[\Delta_{t-1}] - V_m \,\mathbb{E}[\Delta_{t-1}]^{\frac{m+2}{2}}$

Lemma 2

Given $\Delta_t \triangleq \hat{g}_t(\boldsymbol{D}_t) - \hat{g}_t(\boldsymbol{D}_t^*)$. Then, for our algorithm:

 $\mathbb{E}[\Delta_t] \le O\left(\frac{1}{t}\right) + V_m \, \mathbb{E}[\Delta_{t-1}] - \mathbb{E}[\Delta_{t-1}]^{\frac{m+2}{2}}$

Where V_m is a coefficient depends on m, κ , the minimum cluster probability p_i , and the running average of $\alpha_t \alpha_t^T$. Here κ is the same constant used in assumption (A.1).

Lemma 3 $\sum_{t} \frac{||D_t - D_t^*||}{t+1}$ converges almost surely.

Lemma 4 The following claims hold true: P.1) $\hat{g}_t(D_t)$ and $\hat{g}_t(D_t^*)$ converge almost surely $P.2) \hat{g}_t(D_t) - \hat{g}_t(D_t^*)$ converges almost surely to 0 P.3) $\hat{g}_t(D_t^*) - g(D_t^*)$ converges almost surely to 0 P.4) $g(D_t^*)$ converges almost surely

We compare the approximation error and running time of our proposed online cvxMF algorithm with non-negative MF (NMF), cvxMF [1] and online MF [2]

Experimental Validation

Main Result

Under assumptions A.1 to A.3:

 $\{\boldsymbol{D}_t\}_t$ converges almost surely (a.s.) to a stationary point of $g(\mathbf{D})$

Synthetic data: The synthetic datasets were generated by sampling from a 3σ -truncated Gaussian mixture model with 5 components, and with samples-sizes in $[10^3, 10^6]$.

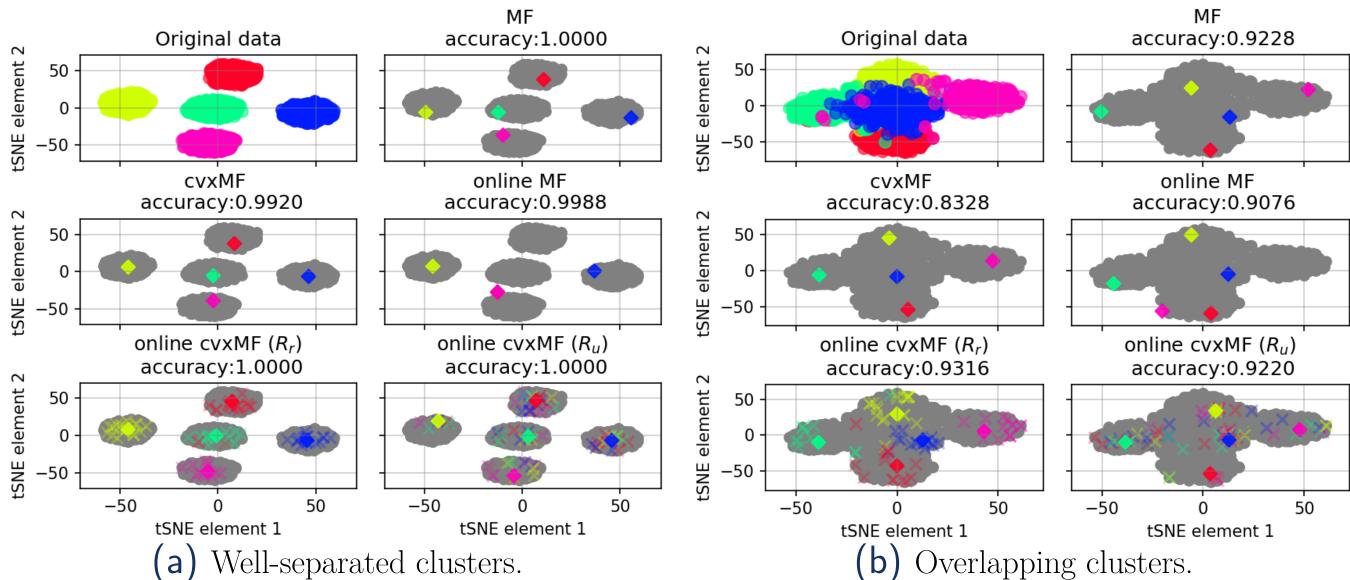


Figure 4:Results for Gaussian mixtures with color-coded clusters. Color-coded circles represent samples, diamonds represent basis vectors learned by the different algorithms, while crosses describe samples in the representative regions. The "interpretability property" can be easily observed visually.

References

[1] Chris HQ Ding, Tao Li, and Michael I Jordan. Convex and semi-nonnegative matrix factorizations. IEEE transactions on pattern analysis and machine intelligence, 32(1):45–55,

[2] Julien Mairal, Francis Bach, Jean Ponce, and Guillermo Sapiro. Online learning for matrix factorization and sparse coding. Journal of Machine Learning Research, 11(Jan):19–60, 2010.

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