# Towards Evolutionary Nonnegative Matrix Factorization 

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#### Abstract

Nonnegative Matrix Factorization ( $N M F$ ) techniques has aroused considerable interests from the field of artificial intelligence in recent years because of its good interpretability and computational efficiency. However, in many real world applications, the data features usually evolve over time smoothly. In this case, it would be very expensive in both computation and storage to rerun the whole $N M F$ procedure after each time when the data feature changing. In this paper, we propose Evolutionary Nonnegative Matrix Factorization (eNMF), which aims to incrementally update the factorized matrices in a computation and space efficient manner with the variation of the data matrix. We devise such evolutionary procedure for both asymmetric and symmetric $N M F$. Finally we conduct experiments on several real world data sets to demonstrate the efficacy and efficiency of $e N M F$.


## Introductions

The recent years have witnessed a surge of interests on Nonnegative Matrix Factorization (NMF) from the artificial intelligence field (Lee and Seung 1999)(Lee and Seung 2001)(Lin 2007)(Kim and Park 2008). Differernt from traditional spectral decomposition methods such as Principal Component Analysis (PCA) and Singular Value Decomposition (SVD), NMF (1) is usually additive, which results in a better interpretation ability; (2) does not require the factorized latent spaces to be orthogonal, which allows more flexibility to adapt the representation to the data set. NMF has successfully been used in many real world applications, such as information retrieval (Shahnaz et al. 2006), environmental study (Anttila et al. 1995), computer vision (Guillamet, Bressan, and Vitrià 2001) and computational social/network science (Wang et al. 2010).

Formally, what $N M F$ does is to factorize a nonnegative data matrix into the product of two (low-rank) nonnegative latent matrices. As $N M F$ requires both factorized matrices to be nonnegative, this will generally lead to sparse, partbased representation of the original data set, which is semantically much more meaningful compared to traditional factorization/basis learning methods. Due to the empirical and theoretical success of $N M F$, people have been working

[^0]on a lot of $N M F$ extensions in the last decade to fit in more application scenarios. Some representative algorithms include nonnegative sparse coding (Eggert and Korner 2004), semi and convex NMF (Ding, Li, and Jordan 2010), and orthogonal tri-NMF (Ding et al. 2006).

Many algorithms have been proposed to solve $N M F$, such as multiplicative updates (Lee and Seung 2001), active set (Kim and Park 2008) and projected gradient (Lin 2007). However, all these algorithms require to hold the whole data matrix in main memory in the entire $N M F$ process, which is quite inefficient in terms of storage cost when the data matrix large (either in data size or the feature dimensionality). To solve this problem, several researchers proposed memory efficient online implementations for NMF in recent years (Cao et al. 2007)(Wang, Li, and König 2011)(Saha and Sindhwani 2010). Rather than processing all data points in a batch mode, these approaches process the data point one at a time in a streaming fashion. Thus they only require the memory to hold one data point through the whole procedure.

In this paper, we consider the problem of $N M F$ in another scenario where the data features are evolving over time ${ }^{1}$. A straightforward solution is to rerun the whole NMF procedure at each time stamp when the data feature change. However, this poses several challenges in terms of space cost, computational time as well as privacy. Let $\mathbf{X}$ and $\widetilde{\mathbf{X}}=\mathbf{X}+\Delta \mathbf{X}$ be the old and new data feature matrices respectively. In many real applications, $\Delta \mathbf{X}$ is usually very sparse while $\widetilde{\mathbf{X}}$ is not ${ }^{2}$. It therefore is not efficient in terms of space cost to re-run $N M F$ since we need to store the whole data feature matrix $\widetilde{\mathbf{X}}$. It is also not efficient in computation since it requires some matrix-matrix multiplication between $\widetilde{\mathbf{X}}$ and the two factorized matrices. What is more, this strategy becomes infeasible for those privacy-sensitive applications where the whole data feature matrix $\widetilde{\mathbf{X}}$ might not be available at a given time stamp. For instance, Facebook's ${ }^{3}$

[^1]privacy policy prohibits the user to keep the downloaded data longer than 24 hours. So, if a data analyst wants to track the community structure on the daily-base, s/he would only have the access to the data feature within a 24 hour window.

For evolutionary data, one common assumption is that the data features evolve smoothly over time (Chi et al. 2007), i.e., the norm of the difference between the data feature matrices at two consecutive time stamps is very small. Based on this assumption, we develop a novel Evolutionary Nonnegative Matrix Factorization (eNMF) algorithm in this paper, where we assume that the factorized matrices also evolve smoothly over time. Instead of minimizing a new similar objective on the evolved feature matrix, eNMF minimizes an upper bound of the objective, and we devise an efficient projected gradient method to solve the problem. Finally we conduct experiments on several real world data sets to demonstrate the efficacy and efficiency of eNMF.

It is worthwhile to highlights several aspects of eNMF.

- eNMF is space efficient. eNMF only needs to hold the difference matrix, which is usually much sparser due to the smoothness assumption.
- eNMF is computationally efficient. One major computational cost in NMF the matrix-matrix multiplications. Our eNMF achieves computational savings by using a much sparser matrix in such matrix-matrix multiplications.
- eNMF is privacy-friendly. eNMF does not need to know the exact data feature matrix. It only requires the factorized matrices at the initial time stamp and the difference data feature matrix. This is particularly useful for those privacy-sensitive applications, e.g., the data feature is only available for a short time window.
- eNMF can be applied to different types of data. We developed two instantiations for both traditional asymmetric $N M F$ (where the feature matrix is rectangular) and symmetric $N M F$ (where the feature matrix is symmetric square, e.g., data similarity matrix).
The rest of this paper is organized as follows. Section 2 introduces the problem formulation and algorithm details. The experimental results are presented in Section 3, followed by the conclusions in Section 4.


## The Algorithm

In this section we will introduce our eNMF algorithm in detail. First we will introduce the basic notations that will be used throughout this paper and problem formulation.

## Problem Formulation

Suppose we have a nonnegative matrix $\mathbf{X} \in \mathbb{R}^{n \times d}$, and we want to factorize it into the product of two nonnegative matrices $\mathbf{F} \in \mathbb{R}^{n \times k}$ and $\mathbf{G} \in \mathbb{R}^{d \times k}$ (usually $k \ll \min (d, n)$ ) under some loss. In this paper we will concentrate on the Frobenius norm loss as it is one of the most popular loss forms, the algorithms under other loss such as KullbackLeighbler divergence (Lee and Seung 2001), $\beta$-divergence (Févotte and Idier 2010) and Bregman divergence (Dhillon and Sra 2005) can be derived similarly. The optimization
problem we need to solve is

$$
\begin{equation*}
\min _{\mathbf{F} \geqslant 0, \mathbf{G} \geqslant 0}\left\|\mathbf{X}-\mathbf{F G}^{\top}\right\|_{F}^{2} \tag{1}
\end{equation*}
$$

where $\|\mathbf{A}\|_{F}^{2}=\operatorname{tr}\left(\mathbf{A}^{\top} \mathbf{A}\right)$ is the square of the matrix Frobenius norm. This problem can be solved via multiplicative updates (Lee and Seung 2001), active set method (Kim and Park 2008) or projected gradient (Lin 2007) method.

Now suppose there is a small on variation on $\mathbf{X}$ so that $\mathbf{X}$ becomes

$$
\begin{equation*}
\widetilde{\mathbf{X}}=\mathbf{X}+\Delta \mathbf{X} \tag{2}
\end{equation*}
$$

and $\widetilde{\mathbf{X}} \in \mathbb{R}^{n \times d}$ is also nonnegative. Our goal is to factorize $\widetilde{\mathbf{X}}$ into the product of two nonnegative matrices $\widetilde{\mathbf{F}} \in \mathbb{R}^{n \times k}$ and $\widetilde{\mathbf{G}} \in \mathbb{R}^{d \times k}$, then we need to solve the following optimization problem

$$
\begin{equation*}
\min _{\widetilde{\mathbf{F}} \geqslant 0, \widetilde{\mathbf{G}} \geqslant 0}\left\|\widetilde{\mathbf{X}}-\widetilde{\mathbf{F}} \widetilde{\mathbf{G}}^{\top}\right\|_{F}^{2} \tag{3}
\end{equation*}
$$

We assume $\|\Delta \mathbf{X}\|_{F}^{2}$ is very small, and $\widetilde{\mathbf{F}}, \widetilde{\mathbf{G}}$ can be represented as

$$
\begin{align*}
\widetilde{\mathbf{F}} & =\mathbf{F}+\Delta \mathbf{F}  \tag{4}\\
\widetilde{\mathbf{G}} & =\mathbf{G}+\Delta \mathbf{G} \tag{5}
\end{align*}
$$

Bringing Eq.(4) and Eq.(5) into problem (3), we can get that

$$
\begin{aligned}
& \left\|\widetilde{\mathbf{X}}-\widetilde{\mathbf{F}} \widetilde{\mathbf{G}}^{\top}\right\|_{F}^{2}=\left\|\mathbf{X}+\Delta \mathbf{X}-(\mathbf{F}+\Delta \mathbf{F})(\mathbf{G}+\Delta \mathbf{G})^{\top}\right\|_{F}^{2} \\
& =\left\|\mathbf{X}+\Delta \mathbf{X}-\mathbf{F} \mathbf{G}^{\top}-\Delta \mathbf{F} \mathbf{G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right\|_{F}^{2}
\end{aligned}
$$

and the constraint here is that

$$
\begin{align*}
\widetilde{\mathbf{F}} & =\mathbf{F}+\Delta \mathbf{F} \geqslant 0  \tag{7}\\
\widetilde{\mathbf{G}} & =\mathbf{G}+\Delta \mathbf{G} \geqslant 0 \tag{8}
\end{align*}
$$

For matrix Frobenius norm, we have the following triangle inequality

$$
\begin{aligned}
& \left\|\mathbf{X}+\Delta \mathbf{X}-\mathbf{F G}^{\top}-\Delta \mathbf{F G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right\|_{F} \\
& \leqslant\left\|\mathbf{X}-\mathbf{F G}^{\top}\right\|_{F}+\left\|\Delta \mathbf{X}-\Delta \mathbf{F G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right\|_{F}
\end{aligned}
$$

In our evolutionary setting, we already got the optimal $\mathbf{F}$ and $\mathbf{G}$ by solving problem (1), thus $\left\|\mathbf{X}-\mathbf{F G}^{\top}\right\|_{F}$ is already minimized. In order to minimize the objective of problem (3), we propose to solve the following optimization problem

$$
\begin{array}{cl}
\min _{\Delta \mathbf{F}, \Delta \mathbf{G}} & \left\|\Delta \mathbf{X}-\Delta \mathbf{F} \mathbf{G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right\|_{F}^{2} \\
\text { s.t. } & \mathbf{F}+\Delta \mathbf{F} \geqslant 0, \mathbf{G}+\Delta \mathbf{G} \geqslant 0 \tag{9}
\end{array}
$$

This is an optimization problem with box constraints, and we propose to apply Projected Gradient (PG) (Lin 2007) method to solve it.

## Projected Gradient

In this section we will introduce how to make use of PG to solve problem (9). For notational convenience, we introduce a box projection operator $P_{\mathbf{B}}[\mathbf{A}]$ as

$$
\left(P_{\mathbf{B}}[\mathbf{A}]\right)_{i j}= \begin{cases}A_{i j} & \text { if } A_{i j} \geqslant B_{i j}  \tag{10}\\ B_{i j} & \text { otherwise }\end{cases}
$$

```
Algorithm 1 Projected Gradient
Require: \(0<\beta<1,0<\sigma<1\). Initialization \(\mathbf{A}^{(0)}\).
Ensure: \(\mathbf{A}^{(0)} \geqslant \mathbf{B}\)
    for \(k=1,2, \cdots\) do
        \(\mathbf{A}^{(k)}=P_{\mathbf{B}}\left[\mathbf{A}^{(k-1)}-\alpha_{k} \nabla f\left(\mathbf{A}^{(k-1)}\right)\right]\)
        where \(\alpha_{k}=\beta^{t_{k}}\), and \(t_{k}\) is the first nonnegative integer
        for which
\(f\left(\mathbf{A}^{(k)}\right)-f\left(\mathbf{A}^{(k-1)}\right) \leqslant \sigma\left\langle\nabla f\left(\mathbf{A}^{(k-1)}\right),\left(\mathbf{A}^{(k)}-\mathbf{A}^{k-1)}\right)\right\rangle\)
    end for
```

Then the PG method for solving the problem

$$
\begin{equation*}
\min _{\mathbf{A} \geqslant \mathbf{B}} f(\mathbf{A}) \tag{11}
\end{equation*}
$$

can be presented in Algorithm 1, where $\langle\cdot, \cdot\rangle$ is the sum of elementwise multiplication, and the rule for determining the step size in Algorithm 1 is usually referred to as the Armijo rule (Bertsekas 1999).

Now let us return to problem (9). If we denote the objective of the above problem by

$$
\begin{equation*}
\mathcal{J}=\left\|\Delta \mathbf{X}-\Delta \mathbf{F} \mathbf{G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right\|_{F}^{2} \tag{13}
\end{equation*}
$$

Then the gradient of $\mathcal{J}$ with respect to $\Delta \mathbf{F}$ and $\Delta \mathbf{G}$ are

$$
\begin{align*}
& \frac{\partial \mathcal{J}}{\partial \Delta \mathbf{F}}=-2\left(\Delta \mathbf{X}-\Delta \mathbf{F G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right)(\mathbf{G}+\Delta \mathbf{G})(14) \\
& \frac{\partial \mathcal{J}}{\partial \Delta \mathbf{G}}=-2\left(\Delta \mathbf{X}-\Delta \mathbf{F G}^{\top}-\mathbf{F} \Delta \mathbf{G}^{\top}-\Delta \mathbf{F} \Delta \mathbf{G}^{\top}\right)^{\top}(\mathbf{F}+\Delta \mathbf{F})(15) \tag{15}
\end{align*}
$$

We can observe that there are two variables, $\Delta \mathbf{F}$ and $\Delta \mathbf{G}$, in problem (9). It is not easy to solve for $\Delta \mathbf{F}$ and $\Delta \mathbf{G}$ simultaneously. However, if we fix one variable, then the problem is convex with respect to the other. Therefore it is natural to adopt the block coordinate descent scheme (Bertsekas 1999), which is an alternating optimization strategy, to solve it. At each round of the iteration, we fix one variable and solve the other (via PG), until some stopping criterion is satisfied. As the objective is lower bounded by zero and after each round its value will decrease, the algorithm is guaranteed to converge. The basic algorithm sketch is summarized in Algorithm 2.

```
Algorithm 2 eNMF
Require: Initialization \(\Delta \mathbf{F}^{(0)}, \Delta \mathbf{G}^{(0)}\).
Ensure: \(\Delta \mathbf{F}^{(0)} \geqslant 0, \Delta \mathbf{G}^{(0)} \geqslant 0\)
    for \(t=1,2, \cdots\) do
        Fix \(\Delta \mathbf{F}=\Delta \mathbf{F}^{(t-1)}\), update \(\Delta \mathbf{G}^{(t)}\) using PG
        Fix \(\Delta \mathbf{G}=\Delta \mathbf{G}^{(t)}\), update \(\Delta \mathbf{F}^{(t)}\) using PG
    end for
```


## Complexity Analysis

For Algorithm 2, we need to hold $\mathbf{F}, \mathbf{G}$ and $\Delta \mathbf{F}, \Delta \mathbf{G}$ in the main memory, thus the total storage complexity is $O(2 k(n+$ $d)$ ). Actually in our experiments, we usually find that the
obtained $\Delta \mathbf{F}$ or $\Delta \mathbf{G}$ is sparse, therefore the storage cost can be further reduced by only storing the nonzero elements.
For computational complexity, as both eNMF and NMF need to evaluate the function objective value (in Armijo rule) when applying PG, the main difference would lie in the evaluation of the function gradient. Suppose that we have $m$ and $\hat{m}$ non-zero elements in the matrices $\mathbf{X}+\Delta \mathbf{X}$ and $\Delta \mathbf{X}$ respectively, then the time cost for eq. (14) and Eq. (15) is $O(\hat{m} k)+O\left(n k^{2}\right)+O\left(d k^{2}\right)$. In contrast, the time complexity for computing the gradient for the original NMF is $O(m k)+O\left(n k^{2}\right)+O\left(d k^{2}\right)$. In many real applications, the matrix $\Delta \mathbf{X}$ is usually much more sparser than the matrix $\mathbf{X}+\Delta \mathbf{X}$ (i.e., $\hat{m} \ll m$ ). Moreover, since $k \ll n, l<m$, $O(m k)$ is dominant term of the time complexity for the original NMF. Therefore, we would expect that the proposed algorithm is much more efficient in computation compared with the original NMF.

## Evolutionary Symmetric NMF

Another interesting scenario is Symmetric NMF (Wang et al. 2010), where we have a symmetric square nonnegative feature matrix $\mathbf{S} \in \mathbb{R}^{n \times n}$ (e.g., the connectivity matrix of an undirected graph). The goal is to factorize it into the product of a nonnegative matrix $\mathbf{G} \in \mathbb{R}^{n \times k}$ (usually $k \ll n$ ) and its transpose by solving the following optimization problem

$$
\begin{equation*}
\min _{\mathbf{G} \geqslant 0}\left\|\mathbf{S}-\mathbf{G G}^{\top}\right\|_{F}^{2} \tag{16}
\end{equation*}
$$

Wang et al.(2010) derived an multiplicative update approach to solve problem (16). Actually, as problem (16) is also a minimization problem with box constraint, we can also apply PG to solve it. Specifically, if we denote the objective of the above problem as

$$
\begin{equation*}
\mathcal{J}_{S}=\left\|\mathbf{S}-\mathbf{G G}^{\top}\right\|_{F}^{2} \tag{17}
\end{equation*}
$$

then we can also solve it by PG using Algorithm 1 with $\mathbf{A}=$ $\mathbf{G}, f(\mathbf{A})=\mathcal{J}_{S}, \mathbf{B}=\mathbf{O}\left(\mathbf{O} \in \mathbb{R}^{n \times k}\right.$ is an all-zero matrix $)$, and the gradient

$$
\begin{equation*}
\nabla f(\mathbf{A})=\frac{\partial \mathcal{J}_{S}}{\partial \mathbf{G}}=-4\left(\mathbf{S}-\mathbf{G G}^{\top}\right) \mathbf{G} \tag{18}
\end{equation*}
$$

In the evolutionary setting, $\mathbf{S}$ is changed to $\widetilde{\mathbf{S}}=\mathbf{S}+\Delta \mathbf{S}$ with a small $\|\Delta \mathbf{S}\|_{F}^{2}$. Then we want to factorize $\widetilde{\mathbf{S}}$ by solving the following optimization problem

$$
\begin{equation*}
\min _{\widetilde{\mathbf{G}} \geqslant 0}\left\|\widetilde{\mathbf{S}}-\widetilde{\mathbf{G}} \widetilde{\mathbf{G}}^{\top}\right\|_{F}^{2} \tag{19}
\end{equation*}
$$

We assume $\widetilde{\mathbf{G}}$ takes the following form

$$
\begin{equation*}
\widetilde{\mathbf{G}}=\mathbf{G}+\Delta \mathbf{G} \tag{20}
\end{equation*}
$$

with a small $\|\Delta \mathbf{G}\|_{F}^{2}$. Bringing Eq.(20) into the objective of problem (19), we obtain

$$
\begin{aligned}
& \left\|\widetilde{\mathbf{S}}-\widetilde{\mathbf{G}} \widetilde{\mathbf{G}}^{\top}\right\|_{F}=\left\|\mathbf{S}+\Delta \mathbf{S}-(\mathbf{G}+\Delta \mathbf{G})(\mathbf{G}+\Delta \mathbf{G})^{\top}\right\|_{F} \\
& \leqslant\left\|\mathbf{S}-\mathbf{G G}^{\top}\right\|_{F}+\left\|\Delta \mathbf{S}-\mathbf{G} \Delta \mathbf{G}^{\top}-\Delta \mathbf{G} \mathbf{G}^{\top}-\Delta \mathbf{G} \Delta \mathbf{G}^{\top}\right\|_{F}(21)
\end{aligned}
$$

Similar as in the asymmetric case, we also minimize the upper bound instead of the original objective in problem (19). As $\left\|\mathbf{S}-\mathbf{G G}^{\top}\right\|_{F}^{2}$ already minimized, we solve the following optimization problem instead for evolutionary SNMF

$$
\begin{array}{cl}
\min _{\Delta \mathbf{G}} & \left\|\Delta \mathbf{S}-\mathbf{G} \Delta \mathbf{G}^{\top}-\Delta \mathbf{G G}^{\top}-\Delta \mathbf{G} \Delta \mathbf{G}^{\top}\right\|_{F}^{2}  \tag{22}\\
\text { s.t. } & \mathbf{G}+\Delta \mathbf{G} \geqslant 0
\end{array}
$$

This problem is still a minimization problem with box constraints, which can be solved by PG. We denote the objective of the above problem by
$\mathcal{J}_{S}^{e}(\Delta \mathbf{G})=\left\|\Delta \mathbf{S}-\mathbf{G} \Delta \mathbf{G}^{\top}-\Delta \mathbf{G} \mathbf{G}^{\top}-\Delta \mathbf{G} \Delta \mathbf{G}^{\top}\right\|_{F}^{2}$
Then problem (22) can be solved using PG in Algorithm 1 with $\mathbf{A}=\Delta \mathbf{G}, f(\mathbf{A})=\mathcal{J}_{S}^{e}, \mathbf{B}=-\mathbf{G}$, and the gradient
$\frac{\partial \mathcal{J}_{S}^{e}}{\partial \Delta \mathbf{G}}=-4\left(\Delta \mathbf{S}-\mathbf{G} \Delta \mathbf{G}^{\top}-\Delta \mathbf{G} \mathbf{G}^{\top}-\Delta \mathbf{G} \Delta \mathbf{G}^{\top}\right)(\mathbf{G}+\Delta \mathbf{G})$
We summarize the procedure of eSNMF in Algorithm 3.

$$
\begin{aligned}
& \text { Algorithm } 3 \text { eSNMF } \\
& \text { Require: } 0<\beta<1,0<\sigma<1 \text {. Initialization } \Delta \mathbf{G}^{(0)} . \\
& \text { Ensure: } \Delta \mathbf{G}^{(0)} \geqslant-\mathbf{G} \\
& \text { for } k=1,2, \cdots \mathbf{d o} \\
& \Delta \mathbf{G}^{(k)}=P_{-\mathbf{G}}\left[\Delta \mathbf{G}^{(k-1)}-\alpha_{k} \nabla \mathcal{J}_{S}^{e}\left(\Delta \mathbf{G}^{(k-1)}\right)\right] \\
& \text { where } \alpha_{k}=\beta^{t_{k}}, \text { and } t_{k} \text { is the first nonnegative integer } \\
& \text { for which } \\
& \mathcal{J}_{S}^{e}\left(\Delta \mathbf{G}^{(k)}\right)-\mathcal{J}_{S}^{e}\left(\Delta \mathbf{G}^{(k-1)}\right) \leqslant \sigma\left\langle\nabla \mathcal{J}_{S}^{e}\left(\Delta \mathbf{G}^{(k-1)}\right),\left(\Delta \mathbf{G}^{(k)}-\Delta \mathbf{G}^{(k-1)}\right)\right\rangle \\
& \text { end for }
\end{aligned}
$$

Suppose that we have $m$ and $\hat{m}$ non-zero elements in the matrices $\mathbf{S}+\Delta \mathbf{S}$ and $\Delta \mathbf{S}$ respectively, then the time cost for eq. (24) is $O(\hat{m} k)+O\left(n k^{2}\right)$. In contrast, the time complexity for computing the gradient for the original NMF is $O(m k)+O\left(n k^{2}\right)$. In many real applications, the matrix $\Delta \mathbf{S}$ is usually much more sparser than the matrix $\mathbf{S}+\Delta \mathbf{S}$ (i.e., $\hat{m} \ll m)$. Moreover, since $k \ll n, l<m, O(m k)$ dominates the time complexity for the original SNMF. Thus we would expect that the proposed algorithm is computationally more efficient compared with the original SNMF.

## Experiments

We conduct experimental results to evaluate the proposed algorithms from the following three aspects:

1 Convergency. How does the overall reconstruction error change wrt the iteration steps?
2 Effectiveness. How effective are the proposed algorithms, compared with the original NMF and SNMF, respectively.
3 Speed. How fast are the proposed algorithms?
The data set we used for evaluation is from DBLP ${ }^{4}$. We construct time-evolving matrices using the publication records from one of the following four conferences: AAAI,

[^2]KDD, SIGIR, NIPS. For each conference, we first construct the author-paper and the co-authorship snapshot matrices from each of its publication years. For the author-paper snapshot matrices, they are asymmetric where each rows are the authors and columns are the papers. If a given author wrote a paper, the corresponding element in the matrix is 1 and 0 otherwise. We aggregate the first 6 snapshot matrices as the initial $\mathbf{X}$ matrix, and treat each of the remaining snapshot matrices as the $\Delta \mathbf{X}$ matrix in Algorithm 2. We denote these four asymmetric time-evolving matrices as AAAI-AP, KDD-AP, SIGIR-AP, NIPS-AP respectively, which are summarized in Table 1. Each of these four asymmetric timeevolving matrices typically contains a few thousands of rows and columns $(n \times d)$, and a few thousands of non-zeros elements $(m)$ in $\mathbf{X}$, a few ( $T$ ) $\Delta \mathbf{X}$ matrices, and a few hundreds of non-zero elements ( $\hat{m}$ ) in the $\Delta \mathbf{X}$ matrix on average.
For the co-authorship snapshot matrices, they are symmetric where each row/column corresponds to an author and edge weights are the number of the co-authored papers. We also aggregate the first 6 snapshot matrices as the initial $\mathbf{S}$ matrix, and treat each of the remaining snapshot matrices as the $\Delta \mathbf{S}$ matrix in Algorithm 3. We denote these four symmetric time-evolving matrices as $A A A I-A A, K D D-A A$, SIGIR-AA, NIPS-AA respectively, which are summarized in Table 1. For each of these four symmetric time-evolving matrices, it typically contains a few thousands of rows/columns $(n \times n)$, and a few thousand, or a few tens of thousands of non-zeros elements ( $m$ ) in the initial $\mathbf{S}$ matrix, a few $(T) \Delta \mathbf{S}$ matrices, and a few thousands of non-zero elements $(\hat{m})$ in the $\Delta \mathbf{S}$ matrix on average.

Table 1: Summary of the data sets

| Name | $\mathbf{n} \times \mathbf{n}(\mathbf{n} \times \mathbf{d})$ | $\mathbf{m}$ | $\mathbf{T}$ | $\mathbf{\mathbf { m }}$ |
| :---: | :---: | :---: | :---: | :---: |
| AAAI-AP | $3,659 \times 2,651$ | 5,762 | 9 | 265 |
| KDD-AP | $1,974 \times 1,118$ | 3,202 | 7 | 256 |
| SIGIR-AP | $2,489 \times 1,867$ | 4,584 | 22 | 124 |
| NIPS-AP | $3,417 \times 2,927$ | 7,111 | 13 | 355 |
| AAAI-AA | $3,659 \times 3,659$ | 10,849 | 9 | 5,059 |
| KDD-AA | $1,974 \times 1,974$ | 3,717 | 7 | 5,639 |
| SIGIR-AA | $2,489 \times 2,489$ | 3,957 | 22 | 6,336 |
| NIPS-AA | $3,417 \times 3,417$ | 6,063 | 13 | 6,860 |

Convergence. In both $e N M F$ and $e S N M F$, instead of minimizing the true reconstruction error directly, we try to minimize its upper bound. Here, we test how the true reconstruction error change wrt the iteration steps. Figures 1-2 show the results on NIPS-AP and NIPS-AA for one time stamp, respectively. We compared our algorithms with the original $N M F$ and $S N M F$ respectively. From the figures, it can be seen that for both $e N M F$ and $e S N M F$, the overall reconstruction error decreases quickly and reaches an steady state wrt the iteration steps, suggesting that our algorithms indeed converge fast. It is worth pointing out that the final reconstruction error of $e N M F$ is very close to that of the original $N M F$. We have similar observation for $e S N M F$ and SNMF.

Effectiveness Comparison. Here, we evaluate the effectiveness of the proposed $e N M F$ and $e S N M F$ in terms of


Figure 1: The reconstruction error vs. iteration steps on NIPS-AA data. The proposed eNMF converges quickly, leading to a similar reconstruction error as NMF after convergence.


Figure 2: The reconstruction error vs. iteration steps on NIPS-AA data. The proposed eSNMF converges quickly, leading to a similar reconstruction error as SNMF after convergence.
the final reconstruction error (i.e., the reconstruction error after the algorithms converge). For each time stamp of a given data set, we run $e N M F$ (or $e S N M F$ for the symmetric matrix) and $N M F$ (or $S N M F$ for the symmetric matrix) until convergence, and then compare the reconstruction errors. The results are summarized in figures 3-4. It can be seen that in terms of the final reconstruction error, our algorithms are very close to the original methods ( $N M F$ and $S N M F$ ). In most cases, the final reconstruction error of the proposed $e N M F$ and $e S N M F$ is within the range of $1 \pm 10 \%$ of that by the original $N M F$ and $S N M F$ respectively. For those exceptions (e.g., $K D D-A P, K D D-A A, S I G I R-A A$ ), our algorithms actually lead to smaller reconstruction error. On the other hand, our proposed methods do not require the original $\mathbf{X}$ and $\mathbf{S}$ matrices, which in turn leads to some nice properties for the applications (e.g., space and computation efficient, privacy-friendly, etc).

Speed Comparison. We also compared the speed between our algorithms and the original $N M F$ and $S N M F$. Figure 5 show the average wall-clock time on each data set.

The results are consistent with the complexity analysis in Section. In most cases, our $e N M F$ and $e S N M F$ are faster than the original $N M F$ and $S N M F$ respectively. The only exception is the NIPS-AA data set, where the proposed eSNMF is slightly slower than the original $S N M F$. This is because for this data set, we have more non-zero elements in the $\Delta \mathbf{S}$ matrix $(\hat{m}=6,860)$ than that of the original the $\mathbf{S}$ matrix ( $m=6,063$ ) on average. Compared with $e N M F$ and $e S$ $N M F$, we can see that the speed saving is more significant in $e N M F$. This is because the $\Delta \mathbf{X}$ matrix is much more sparser than the $\Delta \mathbf{S}$ matrix, - on average, there are a few hundred of non-zero elements in $\Delta \mathbf{X}$, and a few thousand of non-zero elements in $\Delta \mathbf{S}$.


Figure 5: Speed comparison of eNMF/eSNMF and NMF. Our eNMF is much faster than NMF

## Conclusion

We present a novel evolutionary Nonnegative Matrix Factorization (eNMF) strategy to efficient perform NMF in the scenario where the data features are evolving over time. Our method is both storage and computational efficient as well as privacy friendly. The experimental results on real world time evolving networks are presented to demonstrate the effectiveness of our proposed methods.

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Figure 3: Comparison for asymmetric matrices. The x-axis is time stamp (each corresponds to a publication year.), and y-axis is the final reconstruction error. The reconstruction error of the proposed eNMF is very close to that of the original NMF.


Figure 4: Comparison for symmetric matrices. The $x$-axis is time stamp (each corresponds to a publication year.), and $y$-axis is the final reconstruction error. The reconstruction error of the proposed eSNMF is very close to that of the original SNMF.
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[^1]:    ${ }^{1}$ The difference between this setting and online learning is that in online learning, the data points are processed one by one, i.e., the elements in the data matrix are changed one column at a time. However, in our scenario, we allow any elements in the data matrix to change from time to time.
    ${ }^{2}$ Even if $\widetilde{\mathbf{X}}$ is also sparse, it is usually much denser compared with the $\Delta \mathbf{X}$ matrix. See table 1 for some examples.
    ${ }^{3}$ http://www.facebook.com/

[^2]:    4http://www.informatik.uni-trier.de/~ley/db/

