

# SAG-re: Faster Prototyping of Recommender Systems using Stochastic Average Gradient

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## ABSTRACT

In the age of agile software engineering and shorter product lifecycles, data-scientists would ultimately face the challenge of running many experiments and producing high-quality results, with less time. In this paper, we motivate the problem of adopting the stochastic average gradient method (*SAG*) for prototyping model-based recommender systems. We motivate that, by taking advantage of *SAG*'s fast convergence rate and low iteration cost, data-scientists are able to achieve better optimizations for their recommender systems in a shorter amount of time. However, adopting *SAG* in prototyping model-based recommender systems is not trivial because the asymptotic space-complexity of using *SAG* can be prohibitively high. We propose SAG-RE as our approach to resolve the space-complexity challenge. SAG-RE preserves all the benefits and advantages of using *SAG*, and SAG-RE achieves asymptotic space complexity as compact as any memory-less approach. We both prove in theory and extensively evaluate in practice that, SAG-RE yields a better quality optimization within a shorter amount of time, than the two main gradient methods in state-of-the-art of prototyping recommender systems, namely full deterministic gradient, and stochastic gradient.

## Categories and Subject Descriptors

H.3.3 [Information Storage and Retrieval]: Information Search and Retrieval—*Information Filtering*

## General Terms

Asymptotic Time Complexity, Asymptotic Space Complexity, Prototyping, Experimentation

## Keywords

Recommender systems, collaborative filtering, matrix factorization, stochastic gradient, agile software engineering

## 1. INTRODUCTION

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Shopping, text advertising, display advertising, renting movies, listening to music... recommender systems are prevalent and ubiquitous in our daily lives. Matrix factorization (*MF*) is a popular technique in model-based recommender systems. *MF* has been utilized extensively in past research for handling both explicit [4, 9, 7] ratings, and implicit [1, 2, 8, 3, 9] feedback.

In recommender systems that utilize matrix factorization, most optimize an objective function. In the state of the art, full deterministic gradient (*FG*) and stochastic gradient (*SG*) are the two main gradient methods for optimization. All of the recommender systems that we cite above utilize either full deterministic gradient, or stochastic gradient.

Unfortunately, both full deterministic gradient and stochastic gradient have pitfalls when it comes to prototyping recommender systems. Full deterministic gradient can offer high quality optimizations. However, *FG* is slow because at each iteration of optimization, *FG* has to sample through all the entries in the dataset. Stochastic gradient is relatively fast; its iteration cost is low because each iteration of *SG* sample only one or a few entries. However, the trade-off with stochastic gradient is that it often provides low quality optimizations. By chance, stochastic gradient may *eventually* yield a good quality optimization. If it ever happens, it is after a tremendous number of iterations. Thus stochastic gradient is also slow in terms of yielding a good quality optimization within a reasonable amount of time.

High quality optimizations within a short amount of time is important when building recommender systems. The first reason is that data scientists often have to run repeated experiments: e.g. with different objective functions, different metrics, different datasets, and different optimization parameters. The high level goal to run multiple experiments is that, through experimentation and comparing results of multiple trials, data scientists can ultimately get a sufficiently good mix of objective function and hyper parameters for fitting a dataset. The second reason is that product life cycles are shortening in the age of agile software engineering. Thus data scientists are facing or will ultimately face the challenge of running more experiments and producing high quality results with less time.

In this paper, we study the challenge from the perspective of convex-optimization. We propose and hypothesize using the stochastic average gradient (*SAG*) method [6, 5] as a viable alternative to using *FG* and *SG* during the prototyping process. *SAG* has the distinctive advantage that its optimization quality is proven to be much better than *SG*;

at the same time *SAG*'s iteration cost is asymptotically as low as *SG*. However, applying and adapting *SAG* to matrix factorization is not trivial because *SAG* requires previously-computed gradients; and storing these gradients can lead to very high asymptotic space complexity. We explore the challenge with space-complexity, and resolve it by proposing a re-computation approach (*SAG-RE*) that re-computes the previously-computed gradients on-the-fly, on-demand. *SAG-RE* preserves the fast convergence rate and low iteration cost of *SAG*. Moreover, the asymptotic space complexity of *SAG-RE* is as compact as memory-less gradient methods such as *FG* and *SG*.

To the best of our knowledge, we are the first to

- Identify pitfalls associated with using full deterministic gradient and stochastic gradient when data-scientists prototype model-based recommender systems.
- Propose Stochastic Average Gradient (*SAG*) as a viable alternative for yielding higher quality optimizations while enjoying a low iteration cost.
- Extend *SAG* into *SAG-RE* for matrix factorization, resolve the space complexity challenge in adapting *SAG* from the domain of large-scale supervised-machine-learning into the domain of prototyping recommender algorithms.
- Prove in theory, that *SAG-RE* has a convergence rate as fast as the original *SAG*; *SAG-RE* has asymptotic time complexity as efficient as any gradient method with the lowest iteration cost, and *SAG-RE* has asymptotic space complexity as compact as any memoryless gradient method.
- Extensively evaluate and compare *SAG-RE* across multiple RecSys objective functions and diverse datasets.
- Demonstrate in practice that, even without any optimization or fine-tuning on the implementation, *SAG-RE* still yields the best optimization within the shortest time despite the additional time of re-computation, and that *SAG-RE* uses memory at a level similar to the memory-less stochastic gradient.
- Provide an follow-up evidence that both full deterministic gradient and stochastic gradient takes much longer to reach a quality of optimization similar to *SAG-RE*.

## 2. BACKGROUND AND TERMINOLOGY

To motivate our paper and the space complexity challenge, we first introduce the background and the terminology that we use.

**Matrix Factorization.** Model-based recommender systems approximate the *user-item* matrix  $A$  through the dot-product of the *user*-matrix  $U$  and the *item*-matrix  $V$ :  $\hat{A} = U * V$ .

The *user-item* matrix  $A$  is a  $nRows$ -by- $nCols$  matrix.  $A$  can be *sparse*; thus we use  $N$  to indicate the number of non-zero entries in  $A$ .

The *approximation* matrix  $\hat{A}$  also has  $nRows$  rows, and  $nCols$  columns.  $\hat{A}$  is not a sparse matrix. The goal of model-based recommendation is to use the non-zero entries to approximate the missing entries in  $A$ . When multiplying  $U$  and  $V$ , the latent dimensions  $nDims$  cancels-out in the dot product. This is why the *approximation* matrix has identical dimensions as the original *user-item* matrix.

The *user* matrix  $U$  is  $nRows$ -by- $nDims$ :  $U$  has  $nRows$  rows, and  $nDims$  columns.  $nDims$  is the number of latent dimensions. The *item* matrix  $V$  is  $nDims$ -by- $nCols$ .

**Optimizing an Objective Function.** The goal of matrix factorization is to find the best  $U$  and the best  $V$  whose dot product optimizes an objective function:

$$\arg \min_{U,V} (\text{or } \arg \max) \left[ f(U, V) = \sum_{i=1}^{nRows} \sum_{j=1}^{nCols} f(\bar{u}_i, \bar{v}_j) \right] \quad (1)$$

When we take the gradient of the objective function with respect to a row in the *user* matrix  $U$  (e.g.  $\bar{u}_i$ ), we sum up the gradient of all the entries in  $\hat{A}$  that belong to the same row  $\bar{u}_i$ .

$$\frac{df(U, V)}{d\bar{u}_i} = \sum_{j=1}^{nCols} \frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{u}_i} \quad (2)$$

Similarly, when we take the gradient with respect to a column of  $V$  (e.g.  $\bar{v}_j$ ), we sum up the gradients across different rows that belong to the same column:

$$\frac{df(U, V)}{d\bar{v}_j} = \sum_{i=1}^{nRows} \frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{v}_j} \quad (3)$$

Both  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{u}_i}$  and  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{v}_j}$  are vectors of length  $nDims$ , the number of latent dimensions. Specifically,  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{u}_i}$  is a 1-by- $nDims$  row vector;  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{v}_j}$  is a  $nDims$ -by-1 column vector. Similarly, the summed-up gradient  $\frac{df(U, V)}{d\bar{u}_i}$  is a row vector, and  $\frac{df(U, V)}{d\bar{v}_j}$  is a column vector, of length  $nDims$ .

In *SAG*, storing only the summed-up gradients is not sufficient for matrix factorization. The reason is that, each iteration of *SAG* requires the fine-grain gradients of individual entries (e.g.  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{u}_i}$  and  $\frac{df(\bar{u}_i, \bar{v}_j)}{d\bar{v}_j}$ ) that we previously sampled at an iteration before  $t$ . As we will prove, when directly applied to matrix factorization without using our *SAG-RE* approach, *SAG* will have a asymptotic space complexity of  $\theta(nDims * (\min(M, N) + nRows + nCols))$ .  $M$  is the number of *distinct* entries that we have previously sampled. At any iteration  $t$ ,

$$M \propto \sum_{l=1}^t B_l M \leq \sum_{l=1}^t B_l \quad (4)$$

$B_l$  is the batch size at iteration  $l$ ;  $l \leq t$ . Usually the batch size  $B$  is constant for all iterations; then  $M$  is proportional to and is less than or equal to  $B * t$ .

Here, we want to point out that *SAG-RE* preserves the low asymptotic time complexity as *SAG*; and *SAG-RE* reduces asymptotic space complexity to  $\theta(N + nDims * (nRows + nCols))$ . We will prove that this asymptotic space complexity is as compact as any memory-less approach.

**Gradient Methods in Matrix Factorization.** Gradient methods are iterative methods of optimization. When we increase the number of iterations, we expect the quality of optimization to also increase over time. At each iteration, gradient methods sample a batch of  $B$  entries, calculate the gradients of these entries, and use the calculated gradients to update  $U$  and  $V$  for the next iteration:

$$U^{t+1} = U^t + \frac{\alpha^t}{B} \left( \sum_{b=1}^B \frac{df(\bar{u}_{entry(b).i}, \bar{v}_{entry(b).j})}{d\bar{u}_{entry(b).i}} \right) \quad (5)$$

$$V^{t+1} = V^t + \frac{\alpha^t}{B} \left( \sum_{b=1}^B \frac{df(\bar{u}_{entry(b).i}, \bar{v}_{entry(b).j})}{d\bar{v}_{entry(b).j}} \right) \quad (6)$$

At iteration  $t$ ,  $U^t$  is the current approximation of  $U$ . We use the gradients of the sampled batch of entries to update  $U^t$  into  $U^{t+1}$  for iteration  $t+1$ .

$\alpha^t$  is the *learning-rate* or *step-size*, at iteration  $t$ . When the goal of our optimization is to maximize an objective function, we apply *gradient-ascent* on  $U$  and  $V$ ; thus we set  $\alpha^t > 0$ . When we try to minimize an objective function, we apply *gradient-descent* and set  $\alpha^t < 0$ .

$entry(b)$  is the  $b$ -th entry in our batch of samples.  $entry(b).i$  is the *row* number of the entry;  $entry(b).j$  is the *column* number of the entry sampled from  $A$ .

Full deterministic gradient ( $FG$ ) takes all  $N$  samples at each iteration;  $B = N$  in  $FG$ . Stochastic gradient ( $SG$ ) takes only one or a few samples per iteration:  $B$  is usually a constant much less than  $N$ .

**Stochastic Average Gradient.** Stochastic Average Gradient ( $SAG$ ) requires a memory of previously-computed gradients: e.g.  $\bar{m}_U^t$  and  $\bar{m}_V^t$  for matrix factorization. Each iteration of  $SAG$  uses a sampled batch of entries to update the memory. After the update,  $SAG$  then applies the updated memory  $\bar{m}_U^{t+1}$  and  $\bar{m}_V^{t+1}$  respectively on calculating  $U^{t+1}$  and  $V^{t+1}$ :

$$\bar{m}_{entry(b).i}^{t+1} = \frac{df(\bar{u}_{entry(b).i}, \bar{v}_{entry(b).j})}{d\bar{u}_{entry(b).i}} \quad (7)$$

$$\bar{m}_U^{t+1} = \bar{m}_U^t + \sum_{b=1}^B [\bar{m}_{entry(b).i}^{t+1} - \bar{m}_{entry(b).i}^t] \quad (8)$$

$$U^{t+1} = U^t + \frac{\alpha^t}{M} (\bar{m}_U^{t+1}) \quad (9)$$

$$\bar{m}_{entry(b).j}^{t+1} = \frac{df(\bar{u}_{entry(b).i}, \bar{v}_{entry(b).j})}{d\bar{v}_{entry(b).j}} \quad (10)$$

$$\bar{m}_V^{t+1} = \bar{m}_V^t + \sum_{b=1}^B [\bar{m}_{entry(b).j}^{t+1} - \bar{m}_{entry(b).j}^t] \quad (11)$$

$$V^{t+1} = V^t + \frac{\alpha^t}{M} (\bar{m}_V^{t+1}) \quad (12)$$

$\bar{m}_{entry(b).i}^t$  and  $\bar{m}_{entry(b).j}^t$  are the fine-grain gradients of individual matrix entries that were previously sampled.

$\bar{m}_{entry(b).i}^t$  is a 1-by- $nRows$  row vector;  $\bar{m}_{entry(b).j}^t$  is a  $nCols$ -by-1 column vector.

$\bar{m}_U^t$  is a  $nRows$ -by- $nDims$  matrix, because  $\bar{m}_U^t$  aggregates the gradients of all rows in the *user* matrix  $U$ . Similarly,  $\bar{m}_V^t$  is a  $nDims$ -by- $nCols$  matrix.

We apply  $SAG$  into matrix factorization for two reasons. First,  $SAG$  has iteration cost as low as stochastic gradient ( $SG$ ). Second,  $SAG$ 's convergence rate is faster than  $SG$ , and sometimes as fast as full deterministic gradient ( $FG$ ).

**Convergence rate, Iteration cost, and Prototyping recommender systems.** At a high level, the ideal combination of a fast convergence rate and a low iteration cost implies a better optimization in a shorter amount of time when data-scientists prototype model-based recommender systems. An intuition behind gradient methods is that, at least for objective functions that are convex, the gradients

guide the updates of  $U^t$  and  $V^t$  towards the direction of optimization. Convergence rate measures how many iterations a gradient method is expected to take towards reaching optimization. Iteration cost measures how many entries we sample per iteration.

Full deterministic gradient has the best possible convergence rate because each iteration of  $FG$  samples all  $N$  entries in the dataset. However, while  $FG$  is guaranteed to take a less number of iterations than  $SG$  to reach optimization, sampling all  $N$  entries per iteration slows down  $FG$  overall because the optimization process would still take many iterations. Depending on the mathematical properties of the objective function, stochastic gradient often has much slower convergence rates because  $SG$  samples only one or a few random entries per iteration. Therefore, while  $SG$  has the lowest possible  $\theta(1)$  iteration cost, overall  $SG$  is still slow because  $SG$  would take many more iterations to reach optimization.

$SAG$  speeds-up the convergence rate by reusing the gradients of past samples. Reusing past gradients enables  $SAG$  to sample  $\theta(1)$  entries per iteration and to achieve the lowest possible iteration cost. Our evaluation illustrates that  $SAG$  gives a better optimization with less time than  $FG$  and  $SG$ . In this paper, we minimize the drawbacks or costs of using  $SAG$  in matrix factorization while preserving  $SAG$ 's benefits.

### 3. CHALLENGE

As equations 8 and 11 illustrate, updating  $\bar{m}_U^{t+1}$  and  $\bar{m}_V^{t+1}$  requires  $\bar{m}_{entry(b).i}^t$  and  $\bar{m}_{entry(b).j}^t$ .  $\bar{m}_{entry(b).i}^t$  and  $\bar{m}_{entry(b).j}^t$  are the fine-grained gradients of an individual entry  $entry(b)$  from the last time (or the most recent time) that  $entry(b)$  was sampled.

When applying  $SAG$  into matrix factorization, a major challenge is to make these fine-grain gradients available:  $\bar{m}_{entry(b).i}^t$  from equation 8, and  $\bar{m}_{entry(b).j}^t$  from equation 11

A naïve approach is to store all these fine-grain gradients. As we shall prove, the naïve approach is undesirable because storing all these gradients would take up a lot of space.

*Theorem 1.* The total asymptotic space complexity is  $\theta(nDims * (\min(M, N) + nRows + nCols))$  for the naïve approach of storing the fine-grain gradients of all entries that we had previously sampled.

**PROOF.** For each individual entry, the amount of space required is  $2 * nDims$ : the gradient with respect to row  $\bar{u}_i$  ( $\bar{m}_{entry(b).i}^t$ ) is a 1-by- $nDims$  row vector; the gradient with respect to column  $\bar{v}_j$  ( $\bar{m}_{entry(b).j}^t$ ) is a  $nDims$ -by-1 column vector.

When we store the fine-grain gradients of all previously-sampled entries, the amount of space required becomes  $M * 2 * nDims$ . Recalling from the background section,  $M$  is the number of distinct entries that we previously sampled.

As shown in equations 8 and 11,  $SAG$  requires only the most recent gradient of each previously-sampled entry. Thus for each entry, we store a max of only one set of gradients ( $\bar{m}_{entry(b).i}^t$  and  $\bar{m}_{entry(b).j}^t$ ). The total amount of space required becomes  $\min(M, N) * 2 * nDims$ .

Now, according to equations 8 and 11, we must also store the aggregated gradients:  $\bar{m}_U^t$  and  $\bar{m}_V^t$ .  $\bar{m}_U^t$  takes  $nRows * nDims$  space;  $\bar{m}_V^t$  takes  $nDims * nCols$  space. Thus the total amount of space that we use to store the aggregated

gradients is  $(nRows * nDims) + (nDims * nCols)$ , which is equivalent to  $nDims * (nRows + nCols)$  after simplification.

Adding the fine-grain gradients and the aggregated gradients together, the asymptotic space complexity becomes  $\theta(nDims * (\min(M, N) + nRows + nCols))$  after ignoring the constants.  $\square$

**No guarantee that  $\min(M, N)$  is small.** If we can guarantee that  $\min(M, N)$  is small, or that  $\min(M, N)$  is asymptotically not larger than  $nRows$  or  $nCols$ , then the effective asymptotic space-complexity becomes  $\theta(nDims * (nRows + nCols))$ , which is the most compact anyone can possibly get. Unfortunately, we shall prove that there is no such guarantee.

First, we explore what the best possible asymptotical space-complexity can be in matrix factorization.

*Theorem 2.*  $\Omega(N + nDims * (nRows + nCols))$  is the lower-bound asymptotic space-complexity in matrix factorization.

**PROOF.** Matrix factorization is to approximate a matrix  $A$  (e.g. the *user-item* matrix) through the dot product of two matrices  $U$  (e.g. the *user* matrix) and  $V$  (e.g. the *item* matrix).  $A$  has  $N$  non-zero entries.  $U$  is a  $nRows$ -by- $nDims$  matrix;  $V$  is a  $nDims$ -by- $nCols$  matrix. In each iteration of convex optimization, we must update  $U$  and  $V$ , and use an objective function to compare our approximation to the ground-truth matrix  $A$ . Therefore, any matrix factorization algorithm would have an asymptotic space-complexity of at least  $\Omega(N + nDims * (nRows + nCols))$ .  $\square$

If we can guarantee that  $\min(M, N)$  is asymptotically not larger than  $nRows$  or  $nCols$ , then we can prove that the naïve approach has already achieved the best possible asymptotic space-complexity, and that our challenge is irrelevant. However, we shall prove that such guarantee does not exist.

*Theorem 3.* There is no guarantee that  $\min(M, N)$  is asymptotically not larger than  $nRows$  or  $nCols$ .

**PROOF.**  $N$  is the number of non-zero entries in the matrix  $A$ . Unless there is, or unless we are restricted to an upper-bound of sparsity, then  $N$  must have  $O(nRows * nCols)$  space.

$M$  is the number of *distinct* entries that we previously sampled. According to equation 4,  $M$  depends on the batch size at each iteration  $B_t$ , and the number of iterations previously done  $t - 1$ . Usually, the batch size is a constant  $B$ . Thus the lower bound of  $M$  most likely depends on the lower bound of  $t$ . However, the lower bound of  $t$  depends on the convergence rate, and the tolerance of error  $\epsilon$ . For example, if the convergence rate is exponential (e.g.  $O(p^t)$ ), then the lower bound of  $t$  is  $\Omega(\log(\frac{1}{\epsilon}))$ . Therefore, the lower bound of  $M$  does not depend on  $N$ ,  $nRows$  or  $nCols$ . Given a dataset, the only way to enforce  $M \leq N$  is to either tolerate a high error, or to find a combination of objective function and gradient method that yields the fastest convergence rate as possible. The asymptotic space-complexity of using SAG-RE does not depend on  $M$ . Thus SAG-RE does not enforce data-scientists to tolerate a high error. Given any objective function, the convergence rate of SAG [6, 5] is always faster than stochastic gradient and is sometimes as fast as the fastest full deterministic gradient. SAG-RE preserves the convergence rate of SAG.  $\square$

**Chain rule offers no savings in matrix factorization.** In supervised machine-learning, we can use the chain-rule

in differential-calculus to reduce space-complexity. Unfortunately, applying the chain-rule in matrix-factorization would result in a space-complexity larger than the naïve approach.

In supervised machine-learning, the goal is to compute the best-fit column-vector  $\bar{\omega}$  that optimizes an objective function, which can be written as

$$\arg \min_{\bar{\omega}} (\text{or } \arg \max) \left[ F(\hat{y} = X * \bar{\omega}) = \sum_{i=1}^N f_i(\hat{y}_i = \bar{x}_i * \bar{\omega}) \right] \quad (13)$$

$X$  is a  $N$ -by- $d$  matrix:  $N$  is the number of samples, and  $d$  is the number of features.  $\bar{x}_i$  is the 1-by- $d$  row vector representing  $i$ -th sample.  $\bar{\omega}$  is the  $d$ -by-1 column vector of features that we are trying to learn from  $X$ . We can use the chain-rule and re-write the gradient of  $\bar{\omega}$  with respect to  $f_i$ :

$$\frac{df_i}{d\bar{\omega}} = \left( \frac{df_i}{d\hat{y}_i} \right) \frac{d\hat{y}_i}{d\bar{\omega}} = (\bar{x}_i)' \left( \frac{df_i}{d\hat{y}_i} \right) \quad (14)$$

Originally, using the naïve approach of SAG results in  $\theta(\min(M, N) * d + d)$  space. The reason is that  $\frac{df_i}{d\bar{\omega}}$  is a  $d$ -by-1 column vector; and the naïve approach stores  $\min(M, N)$  copies of them. The memory gradient  $\bar{m}_{\bar{\omega}}$  is a  $d$ -by-1 column vector and thus takes  $\theta(d)$  space.

The dot-product  $\hat{y}_i = (\bar{x}_i * \bar{\omega})$  is a 1-by-1 scalar. Consequently,  $\frac{df_i}{d\hat{y}_i}$  is also a 1-by-1 scalar. From equation 13,  $\frac{d\hat{y}_i}{d\bar{\omega}} = (\bar{x}_i)'$ . Therefore, we can apply the chain rule and reduce space-complexity to  $\theta(\min(M, N) + d)$ , because we can use the vector  $\bar{x}_i$  to re-compute  $\frac{df_i}{d\bar{\omega}}$  from the scalar  $\frac{df_i}{d\hat{y}_i}$ .

*Theorem 4.* Applying the chain-rule for using SAG in matrix factorization would result in  $\theta(\min(M, N) + nDims * (\min(M, N) + nRows + nCols))$  space.

**PROOF.** In matrix factorization,  $\hat{a}_{ij} = (\bar{u}_i * \bar{v}_j)$  is a 1-by-1 scalar. Therefore, we can rewrite the gradients as

$$\frac{df}{d\bar{u}_i} = \left( \frac{df}{d\hat{a}_{ij}} \right) \frac{d\hat{a}_{ij}}{d\bar{u}_i} = (\bar{v}_j)' \left( \frac{df}{d\hat{a}_{ij}} \right) \quad (15)$$

$$\frac{df}{d\bar{v}_j} = \left( \frac{df}{d\hat{a}_{ij}} \right) \frac{d\hat{a}_{ij}}{d\bar{v}_j} = (\bar{u}_i)' \left( \frac{df}{d\hat{a}_{ij}} \right) \quad (16)$$

$\left( \frac{df}{d\hat{a}_{ij}} \right)$  is a 1-by-1 scalar, and the chain-rule approach stores  $\min(M, N)$  copies, occupying  $\theta(\min(M, N))$  space.

Unfortunately both  $U$  and  $V$  change over time in matrix factorization. When we apply the chain-rule, we cannot just use the current versions of  $\bar{u}_i$  and  $\bar{v}_j$ . We must use and thus must store the past versions of  $\bar{u}_i^l$  and  $\bar{v}_j^l$  at the last time  $l$  that the entry  $a_{ij}$  (in matrix  $A$ ) was sampled. Both  $\bar{u}_i^l$  and  $\bar{v}_j^l$  are vectors of length  $nDims$ . Therefore, using the chain rule induces an additional  $\min(M, N) * 2 * nDims$  space. When we include the memory of aggregated gradients  $\bar{m}_U$  and  $\bar{m}_V$ , the total space-complexity becomes larger than the naïve approach with  $\theta(\min(M, N) + nDims * (\min(M, N) + nRows + nCols))$  space. The chain-rule approach yields space savings in supervised machine-learning because  $\bar{x}_i$  does not change over time; so there is no need to store past versions of  $\bar{x}_i$ .  $\square$

## 4. APPROACH

SAG-RE re-computes  $\bar{m}_{entry(b).i}^t$  in equation ?? and  $\bar{m}_{entry(b).j}^t$  in equation ?. The chain-rule approach must store  $\min(M, N)$  versions of past matrices because

## 5. FUTURE WORK & CONCLUSION

This paper is the first in the series of our study on data scientists prototyping model-based recommender systems. We explored the convex-optimization perspective of the problem: we propose Stochastic Average Gradient as a viable alternative to Full Deterministic gradient and Stochastic gradient. By taking advantage of *SAG*'s fast convergence rate and low iteration cost, we aim to enable data-scientists run more experiments and produce high quality results with less time. In theory, we proved that our extension and adaptation of *SAG* preserves the fast convergence rate as the original *SAG*. Furthermore, *SAG-RE* has asymptotic time complexity as efficient as gradient methods with the lowest iteration cost, and asymptotic space complexity as compact as any memory-less gradient methods. In practice, through extensive evaluation we demonstrated that, even without any fine-tuning or optimization of the implementation, *SAG-RE* still outperforms both full deterministic gradient and stochastic gradient in terms of reaching the best quality optimization within the same amount of time. Following up, we provided evidence that full deterministic gradient and stochastic gradient would take much longer to reach a quality of optimization similar to *SAG-RE*.

Currently we are extending *SAG-RE* in two directions. Both directions relate to running an iteration of full deterministic gradient in *SAG-RE*. First, we are investigating if it is beneficial to run an iteration of full deterministic gradient more often. In our experiments, we observed that both *SG* and *SAG* may converge early; the optimization may get stuck at a local sub-optimum for a long number of iterations. Thus we are exploring if an iteration of full deterministic gradient would get the optimization back on track in case *SAG-RE* gets stuck. Secondly, we aim to investigate how well *SAG-RE* would perform in the production environment, and in distributed systems potentially running in parallel, because running full deterministic gradient even once can be prohibitive for full-scale datasets with millions to billions of non-zero entries.

In the future, we also aim to complete our ongoing work on the metrics perspective and on the software engineering perspective. Given a dataset, the quality of a recommender system is often evaluated in various metrics: e.g. precision, recall, area under curve, reciprocal rank, NDCG, and variants of the above such as top-K precision and top-K hit rate. Many papers in the literature claim their objective function is better by illustrating that their objective function performs in some of these metrics better than other objective functions. Therefore, in the metrics perspective, we are exploring and investigating which factors are more relevant and important towards scoring high in the various metrics: is it the objective function, the method for convex-optimization such as *SAG*, other fine-tuning mechanisms such as bootstrapping, or the hyper-parameters that we use in convex-optimization. All of these factors can be dataset-specific. Indeed, our inherent assumption in this paper is that a better quality optimization yields better recommender systems. In the future, we would like to explore if there are other factors that are more worthwhile than a fast convergence rate or a low iteration cost towards better recommender systems.

In the software engineering perspective, we study how to increase the productivity of data scientists. At this point, we are designing and developing a *mix-n-match* or *plug-n-*

*play* framework that enables data scientists in a least effort way, to very rapidly prototype and experiment many different combinations of objective functions, datasets, gradient methods, hyper parameters and evaluation metrics.

## 6. REFERENCES

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