

Fast Monte-Carlo Low Rank Approximations for Matrices

Shmuel Friedland
University of Illinois at Chicago

joint work with M. Kaveh, A. Niknejad and H. Zare

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<http://www.math.uic.edu/~friedlan>

1 Statement of the problem

Data is presented in terms of a matrix

$$A = \begin{bmatrix} a_{11} & a_{12} & \dots & a_{1n} \\ a_{21} & a_{22} & \dots & a_{2n} \\ \vdots & \vdots & \vdots & \vdots \\ a_{m1} & a_{m2} & \dots & a_{mn} \end{bmatrix}$$

Examples

1. digital picture: 512×512 matrix of pixels

2. DNA-microarrays: $60,000 \times 30$

(rows are genes and columns are experiments)

3. web pages activities:

a_{ij} -the number of times webpage j was accessed from web page i

Object: condense data and storage it effectively

2 Matrix SVD

Let $A \in \mathbb{C}^{m \times n}$. Then $A : \mathbb{C}^n \rightarrow \mathbb{C}^m$. Assume $\mathbb{C}^n, \mathbb{C}^m$ equipped with standard inner product $\langle \mathbf{x}, \mathbf{y} \rangle := \mathbf{y}^* \mathbf{x}$.

Then $A = U \Sigma V^*$, where $U \in U(m), V \in U(n)$, $\Sigma = \text{diag}(\sigma_1, \dots, \sigma_{\min(m,n)}) \in \mathbb{R}_+^{m \times n}$.

U, V transition matrices from $[\mathbf{u}_1, \dots, \mathbf{u}_m], [\mathbf{v}_1, \dots, \mathbf{v}_n]$ to the standard bases in $\mathbb{C}^m, \mathbb{C}^n$ respectively.

For $k \leq r$ let $\Sigma_k = \text{diag}(\sigma_1, \dots, \sigma_k) \in \mathbb{R}^{k \times k}$, and $U_k \in U(m, k), V_k \in U(n, k)$ having the first k columns of U, V respectively. Then $A_k := U_k \Sigma_k V_k^*$ the best rank k approximation in Frobenius and operator norm of A :

$$\min_{B \in \mathcal{R}(m,n,k)} \|A - B\| = \|A - A_k\|.$$

$A = U_r \Sigma_r V_r^*$ is Reduced SVD

$(r \geq) \nu$ numerical rank of A if $\frac{\sigma_{\nu+1}}{\sigma_\nu} \approx 0$.

A_ν is a noise reduction of A .

Noise reduction has many applications in image processing, DNA-Microarrays analysis, data compression.

3 SVD in inner product spaces

U_i is m_i -dimensional IPS over \mathbb{C} , with $\langle \cdot, \cdot \rangle_i, i = 1, 2$.

$T : U_1 \rightarrow U_2$ linear operator. $T^* : U_2 \rightarrow U_1$ the adjoint operator: $\langle T\mathbf{x}, \mathbf{y} \rangle_2 = \langle \mathbf{x}, T^*\mathbf{y} \rangle_1$.

$S_1 := T^*T : U_1 \rightarrow U_1$,

$S_2 := TT^* : U_2 \rightarrow U_2$.

S_1, S_2 self-adjoint: $S_1^* = S_1, S_2^* = S_2$ and nonnegative definite: $\langle S_i \mathbf{x}_i, \mathbf{x}_i \rangle_i \geq 0$.

$\sigma_1^2 \geq \dots \geq \sigma_r^2 > 0$ positive eigenvalues of S_1 and S_2 and $r = \text{rank } T = \text{rank } T^*$. Let

$S_1 \mathbf{v}_i = \sigma_i^2 \mathbf{v}_i, \langle \mathbf{v}_i, \mathbf{v}_j \rangle_1 = \delta_{ij}, i, j = 1, \dots, r$.

Define $\mathbf{u}_i := \sigma_i^{-1} T \mathbf{v}_i, i = 1, \dots, r$. Then

$\langle \mathbf{u}_i, \mathbf{u}_j \rangle_2 = \delta_{ij}, i, j = 1, \dots, r$.

Complete $\{\mathbf{v}_1, \dots, \mathbf{v}_r\}$ and $\{\mathbf{u}_1, \dots, \mathbf{u}_r\}$ to orthonormal bases $[\mathbf{v}_1, \dots, \mathbf{v}_{m_1}]$ and $[\mathbf{u}_1, \dots, \mathbf{u}_{m_2}]$ in U_1 and U_2 .

4 RANDOM k -SVD

Stable numerical algorithms of SVD introduced by Golub-Kahan 1965, Golub-Reinsch 1970:

Implicit QR Algo to reduce to upper bidiagonal form using Householder matrices, then Golub-Reinsch SVD algo to zero superdiagonal elements.

Complexity: $O(mn \min(m, n))$.

In applications for massive data:

$A \in \mathbb{R}^{m \times n}$, $m, n \gg 1$ needed a good approximation

$$A_k = \sum_{i=1}^k \mathbf{x}_i \mathbf{y}_i^T, \mathbf{x}_i \in \mathbb{R}^m, \mathbf{y}_i \in \mathbb{R}^n, i = 1, \dots, k \ll \min(m, n).$$

Random A_k approximation algo:

Find a good algo by reading l rows or columns of A at random and update the approximations.

Frieze-Kannan-Vempala FOCS 1998 suggest algo without updating.

5 FKNZ RANDOM ALGO [4]

Fast k -rank approximation and SVD algorithm

Input: positive integers m, n, k, l, N , $m \times n$ matrix A , $\epsilon > 0$.

Output: an $m \times n$ k -rank approximation B_f of A , with the ratios $\frac{\|B_0\|}{\|B_t\|}$ and $\frac{\|B_{t-1}\|}{\|B_t\|}$, approximations to k -singular values and k left and right singular vectors of A .

1. Choose k -rank approximation B_0 using k columns, (or rows), of A .

2. **for** $t = 1$ **to** N

- Select l columns, (or rows), from A at random and update B_{t-1} to B_t .

- Compute the approximations to k -singular values, and k left and right singular vectors of A .

- If $\frac{\|B_{t-1}\|}{\|B_t\|} > 1 - \epsilon$ let $f = t$ and finish.

Complexity: $O(mnk)$.

Each iteration $\|A - B_{t-1}\|_F \geq \|A - B_t\|_F$.

6 DETAILS

Choose at random k columns of A . Apply modified Gram-Schmidt algo to obtain $\mathbf{x}_1, \dots, \mathbf{x}_q \in \mathbb{R}^m, q \leq k$.

Set $B_0 := \sum_{i=1}^q \mathbf{x}_i (\mathbf{A}^T \mathbf{x}_i)^T$.

$$\|A - B_0\|_F^2 = \text{tr } A^T A - \text{tr } B_0^T B_0 = \text{tr } A^T A - \sum_{i=1}^q (\mathbf{A}^T \mathbf{x}_i)^T (\mathbf{A}^T \mathbf{x}_i).$$

Choose at random another l columns of A : $\mathbf{w}_1, \dots, \mathbf{w}_l$.

Apply modified Gram-Schmidt algo to

$\mathbf{x}_1, \dots, \mathbf{x}_q, \mathbf{w}_1, \dots, \mathbf{w}_l$ to obtain o.n.s.

$\mathbf{x}_1, \dots, \mathbf{x}_q, \mathbf{x}_{q+1}, \dots, \mathbf{x}_p$. Form

$$C_0 := B_0 + \sum_{i=q+1}^p \mathbf{x}_i (\mathbf{A}^T \mathbf{x}_i)^T.$$

Find the first left k -o.n. left singular vectors $\mathbf{v}_1, \dots, \mathbf{v}_k$ of

C_0 . Then $B_1 := \sum_{i=1}^k \mathbf{v}_i (\mathbf{A}^T \mathbf{v}_i)^T$ and

$$\text{tr } B_0^T B_0 \leq \text{tr } B_1^T B_1.$$

Obtain B_t from B_{t-1} as above.

7 Lifting body original

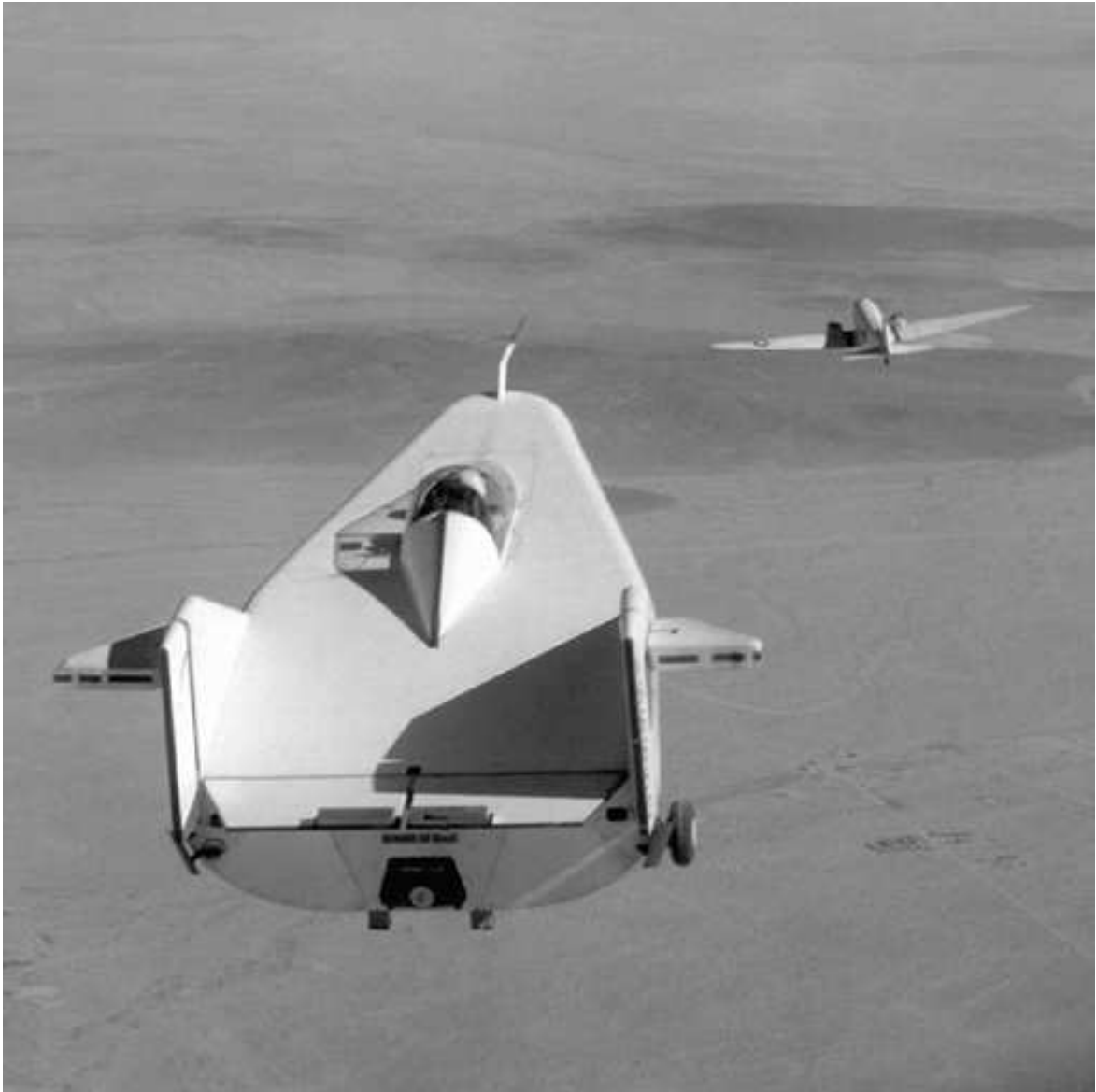


Figure 1: Lifting body image 512×512 .

8 Lifting body compressed

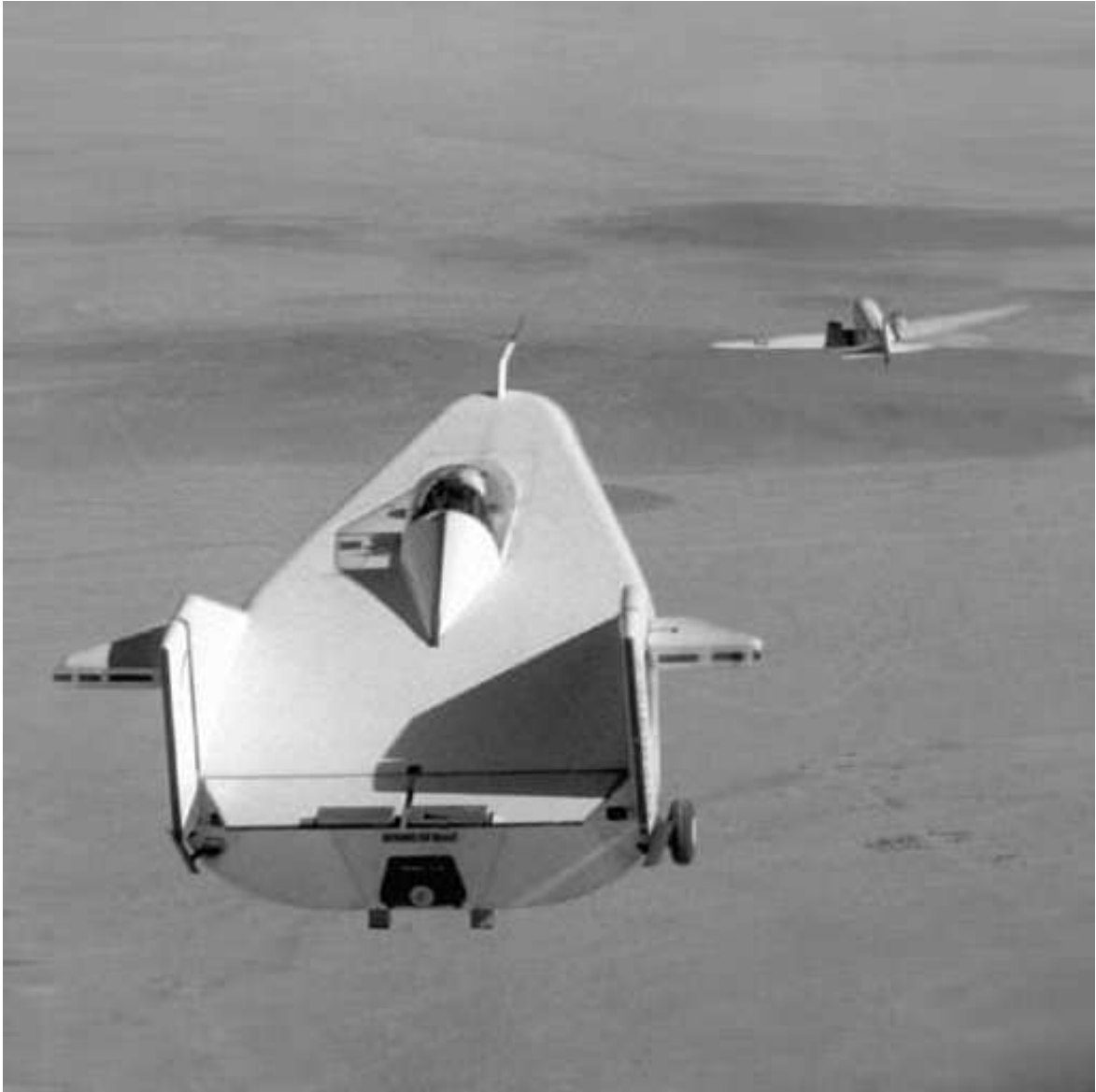


Figure 2: 80-rank approximation of Lifting body image 512×512 .

9 SIMULATIONS 1

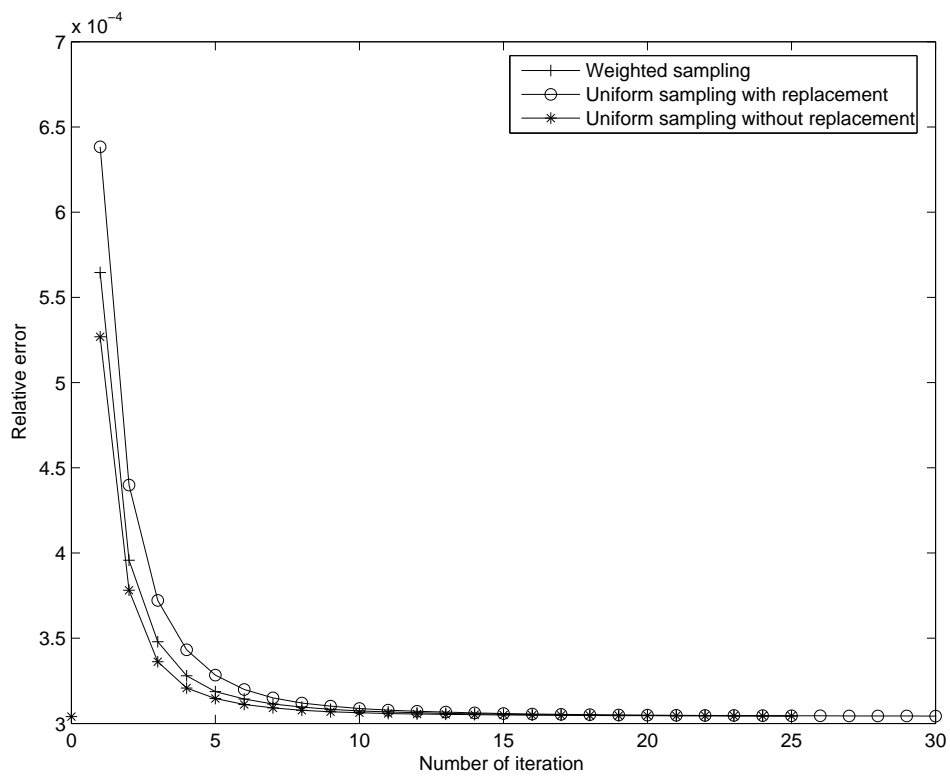


Figure 3: Convergence property of the Monte-Carlo method for Liftingbody image(512×512), $k = 80$.

10 SIMULATIONS 2

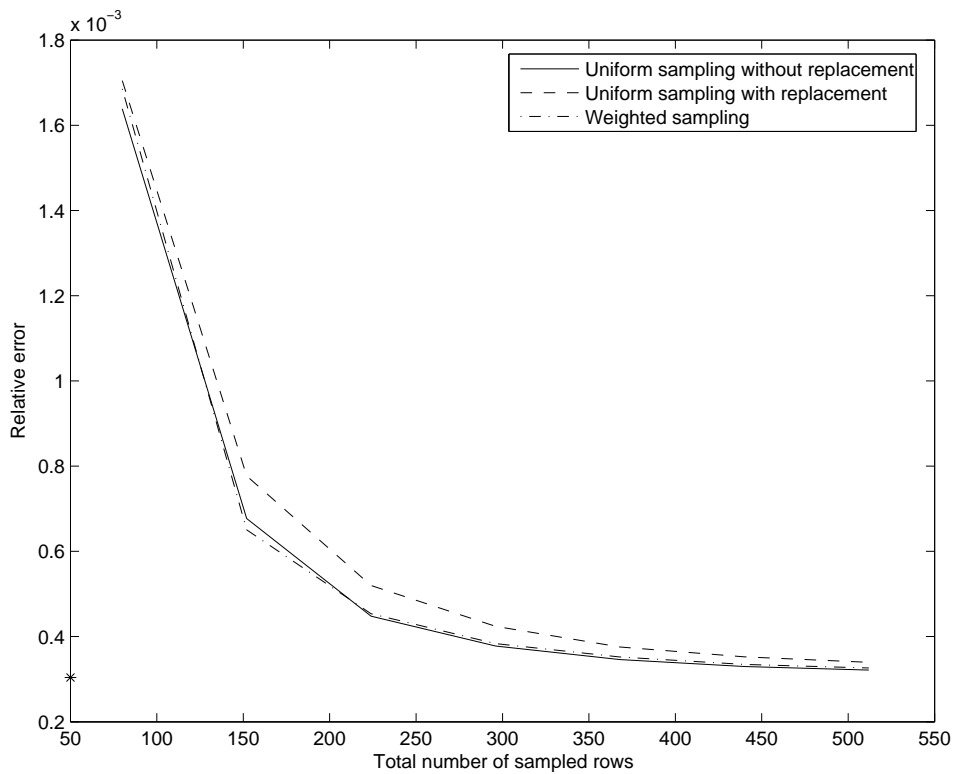


Figure 4: Liftingbody: relative errors versus total number of sampled rows, $k = 100$

11 Camera man original



Figure 5: Camera man image 256×256 .

12 Camera man compressed



Figure 6: 80-rank approximation of Camera man 256×256 .

13 SIMULATIONS 3

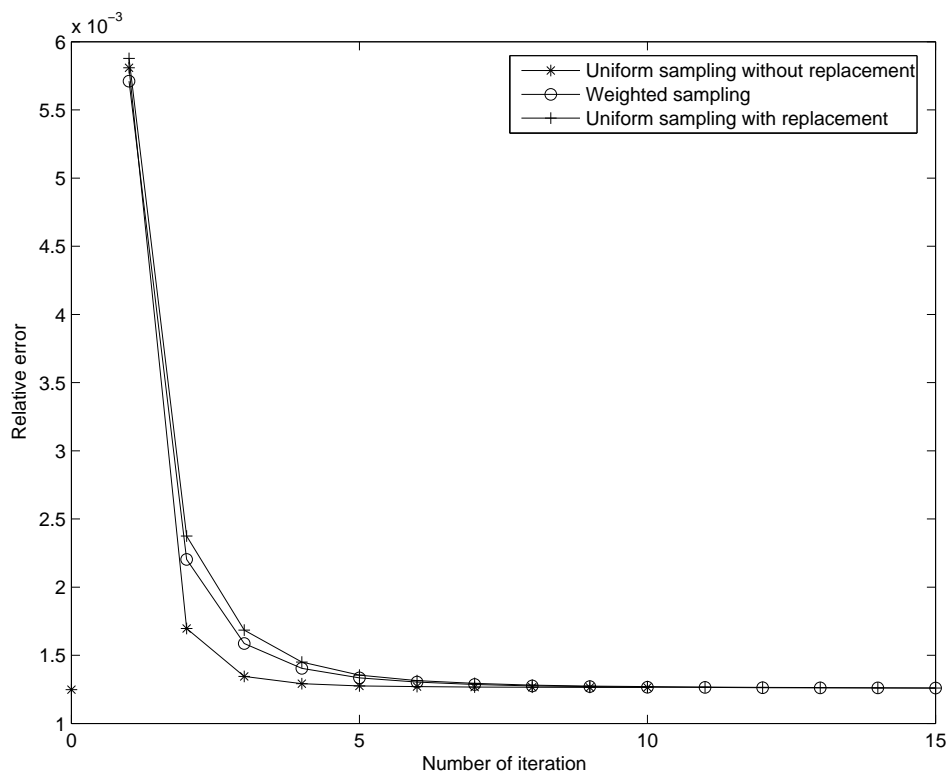


Figure 7: Convergence property of the Monte-Carlo method for Cameraman image(256×256), $k = 80$.

14 SIMULATIONS 4

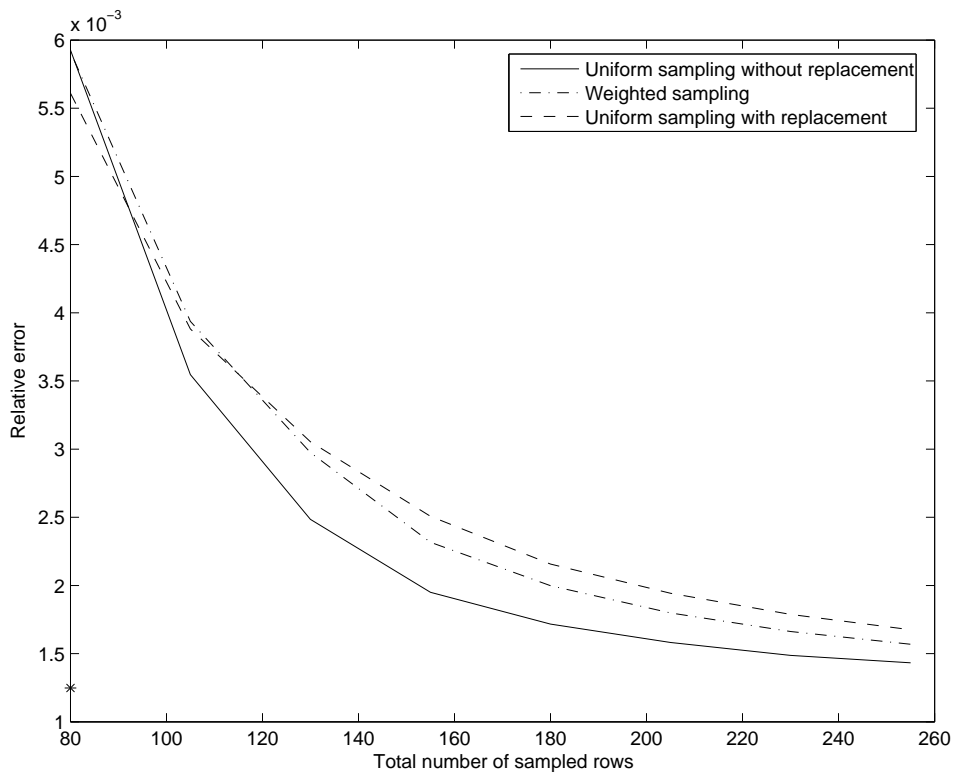


Figure 8: Cameraman: Relative error versus total number of sampled rows, $k = 80$.

15 SIMULATIONS 5

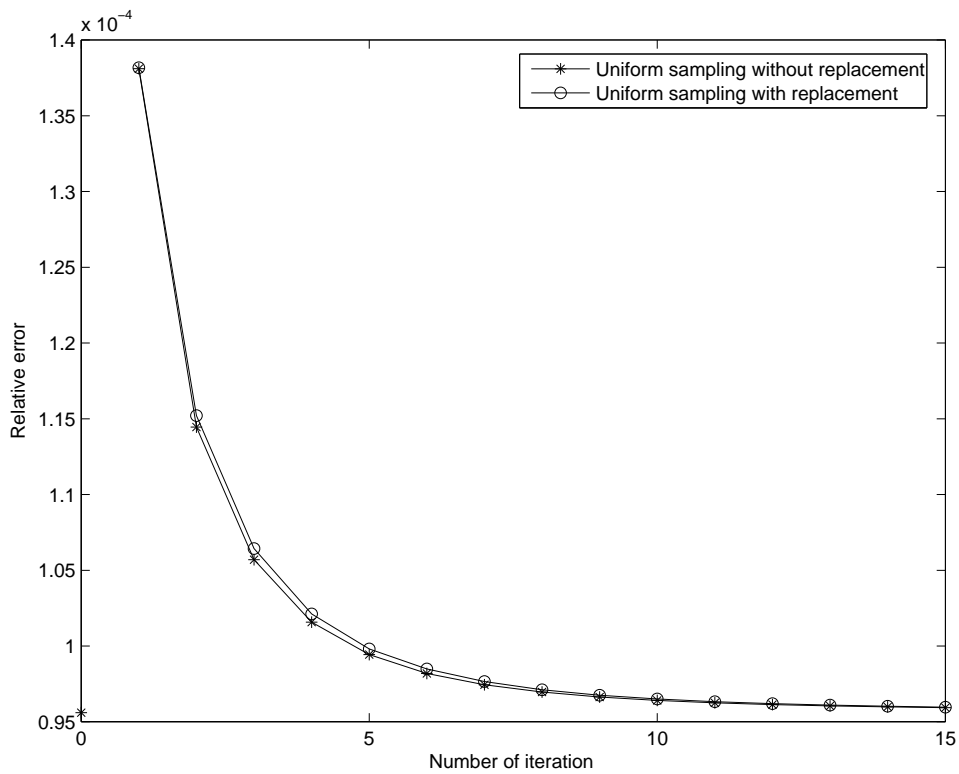


Figure 9: Convergence property of the Monte-Carlo method for random data matrix(3000×500) $k = l = 100$.

16 COMPARISONS

Table 1: Comparison of relative error and speed up of our algorithm with optimum k -rank approximation algorithm

Data sets	Speed up	Re. ratio
Cameraman(256 \times 256), $k = 80$	1.145	1.083
Liftingbody (512 \times 512), $k = 100$	8	1.08
Map image(627 \times 865) $k = 200$	3.33	1.067
Random matrix(8000 \times 200) $k = 100$	42	1.1

17 Choosing columns of A

Frieze, Kannan and Vempala [8] suggest to choose column $\mathbf{c}_i(A)$ with probability $\frac{\|\mathbf{c}_i(A)\|^2}{\|A\|_F^2}$.

If $s \geq k$ are chosen then the k -approximation satisfies A_k
 $\|A - A_k\|_F^2 \leq \sum_{i=k+1}^m \sigma_i(A)^2 + \frac{10k}{s} \|A\|_F^2$.

If $s \geq \frac{k}{10\epsilon}$ then

$$\|A - A_k\|_F^2 \leq \sum_{i=k+1}^m \sigma_i(A)^2 + \epsilon \|A\|_F^2.$$

Deshpande, Rademacher, Vempala and Wang [2] improved the sampling by modifying the sampling $\mathbf{c}_i(A)$ according to new probabilities $\frac{\|\mathbf{c}_i(A - A_k)\|^2}{\|A - A_k\|_F^2}$.

Perhaps our algorithm can be combined with above sampling of columns to get better results.

References

- [1] O. Alter, P.O. Brown and D. Botstein, Singular value decomposition for genome-wide expression data processing and modelling, *Proc. Nat. Acad. Sci. USA* 97 (2000), 10101-10106.
- [2] A. Deshpande, L. Rademacher, S. Vemapala and G. Wang, Matrix Approximation and Projective Clustering via Volume Sampling, *SODA*, 2006.
- [3] S. Friedland, A New Approach to Generalized Singular Value Decomposition, *SIMAX* 27 (2005), 434-444.
- [4] S. Friedland, M. Kaveh, A. Niknejad and H. Zare, Fast Monte-Carlo Low Rank Approximations for Matrices, *Proc. IEEE SoSE*, 2006, 6 pp., to appear.
- [5] S. Friedland, M. Kaveh, A. Niknejad and H. Zare, An Algorithm for Missing Value Estimation for DNA Microarray Data, *Proceedings of ICASSP 2006*, Toulouse, France, 4 pp., to appear.
- [6] S. Friedland, A. Niknejad and L. Chihara, A simultaneous reconstruction of missing data in DNA

microarrays, to appear in *Linear Algebra and Its Applications*.

- [7] S. Friedland, J. Nocedal and M. Overton, The formulation and analysis of numerical methods for inverse eigenvalue problems, *SIAM J. Numer. Anal.* 24 (1987), 634-667.
- [8] A. Frieze, R. Kannan and S. Vempala, Fast Monte-Carlo algorithms for finding low rank approximations, *Proceedings of the 39th Annual Symposium on Foundation of Computer Science*, 1998.
- [9] G.H. Golub and C.F. Van Loan, *Matrix Computation*, John Hopkins Univ. Press, 3rd Ed., 1996.