# A hybrid randomized algorithm for image compression 

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#### Abstract

In this paper, we present a randomized algorithm for image compression by improving the Markov chain Monte Carlo algorithm and by applying the principal component analysis method. Some Numerical examples are provided to show that the proposed algorithm is effective.


Keywords: Image compression, Markov chain, Principal component analysis, Robust Monte Carlo algorithm

## 1 Introduction

Image compression is addresses the problem of reducing the amount of data required to represent a digital image. The image transformation from color to the gray-level (intensity) image $I$ belongs to the most common algorithms. Its implementation is usually based on the weighted sum of three color components $R, G$, and $B$ according to relation [1]

$$
I=\alpha R+\beta B+\gamma G
$$

The $R, G$, and $B$ matrices contain image color components, the weights $\alpha, \beta$, and $\gamma$ were determined with regards to the possibilities of human perception [2]. There is a huge amount of algorithms [1, 2, 5, 7] based on various principles leading to the image compression. Algorithms based on the image color reduction are mostly lossy but their results are still acceptable for some applications. Among various qualified statistical procedures, the principal component analysis (PCA) is the most classical and popular one. Since it uses only simple linear algebra and avoids any probabilistic assumption on the data, PCA has been widely used in applications [2, 5-7]. Our objective in this paper is to derive randomized PCA method from robust Markov chain Monte Carlo algorithm, with the aim of providing a better dimension reduction tool for real images. Consequently, the proposed algorithm it is more efficient in applications.

## 2 The PCA method and computation of PCs

Principal component analysis (PCA) is an exploratory statistical method for graphical description of the information present in large data sets. In most applications, PCA consists of studying $p$ variables measured on $n$ individuals. When $n$ and $p$ are large, the aim is to synthesize the huge quantity of information into an easy and understandable form. With minimal additional effort PCA provides a road map for how to reduce a complex data set to a lower dimension to reveal the sometimes hidden, simplified structure that often underlie it. Finding principal components (PCs) reduces to finding the eigenvalues and eigenvectors of a positive-semidefinite matrix [6, 7].
Suppose that $X$ is a vector of $p$ random variables with a known covariance matrix $\Sigma$, and that the variances of the $p$ random variables and the structure of the covariances or correlations between the pariables are of interest. The first step is to look for a linear function $v_{1}^{T} x$ of the elements of $x$ having maximum variance, where $v_{1}$ is a vector of $p$ constants $v_{11}, v_{12}, \ldots, v_{1 p}$ and so that

$$
v_{1}^{T} x=v_{11} x_{1}+v_{12} x_{2}+\cdots+v_{1 p} x_{p}
$$

Next, look for a linear function $v_{2}^{T} x$, uncorrelated with $v_{1}^{T} x$ having maximum variance, and so on, so that at the $k^{t h}$ stage a linear function $v_{k}^{T} x$ is found that has maximum variance subject to being uncorrelated with $v_{1}^{T}, v_{2}^{T}, \ldots, v_{k-1}^{T}$. The $k^{t h}$ derived variable, $v_{k} x$ is the $k^{t h}$ PC. Up to $p$ PCs could be found, but it is hoped, in general, that most of the variation in $x$ will be accounted for by $m$ PCs, where $m \ll p$. Suppose that $z$ be the vector whose $k^{t h}$ element is $z_{k}$, the $k^{t h} \mathrm{PC}, k=1,2, \ldots, p$. Let $z=A^{T} x$, where $A$ is the orthogonal matrix whose $k^{t h}$ column, $v_{k}$, is the $k^{t h}$ eigenvector of $\Sigma$. Thus, the Pcs are defined by an orthonormal linear transformation of $x$. We can write [6]

$$
\Sigma A=A \Lambda
$$

where $\Lambda$ is the orthogonal matrix whose $k^{t h}$ diagonal element is $\lambda_{k}$, the $k^{t h}$ eigenvalue of $\Sigma$ and $\lambda=\operatorname{var}\left(v_{k}^{T} x\right)=$ $\operatorname{var}\left(z_{k}\right)$.

Theorem 2.1 Suppose that we wish to predict each random variable, $x_{j}$ in $x$ by a linear function of $y$, where $y=B^{T} x$, and $B$ is a $p \times q$ matrix, If $\sigma_{j}^{2}$ is the residual variance in predicting $x_{j}$ from $y$, then $\sum_{j=1}^{p} \sigma_{j}^{2}$ is minimized if $B=A_{q}$, where $A_{q}$ consist of the first $q$ column of $A$.
Proof. See [6]
According to the above theorem, if we wish to get the best linear predictor of x in a $q$-dimensional subspace, in the sense of minimizing the sum over elements of $x$ of the residual variances, then this optimal subspace is defined by the first $q$ PCs. It can be shown that [7], finding the PCs reduces to finding the eigenvalues and eigenvectors of a positive-semidefinite matrix. Also, in [7] is presented PCA-MCMC algorithm for computing the PCs. We now propose a robust MCMC (RMCMC) algorithm based on control variates variance reduction that can be used to solve such an eigenpair problem.

## 3 The proposed algorithm

Monte Carlo methods gives statistical estimates for the functional of the solution by performing random sampling of a certain random variable whose mathematical expectation is the desired functional [3]. The basic power method [8] is the traditional starting point for a Monte Carlo determination of the eigenpair associated with the eigenvalues of largest absolute value $\lambda_{1}$. While various versions of the Monte Carlo power method often compute this dominant eigenvalue very well, computing subdominant eigenvalues $\lambda_{2}, \lambda_{3}, \ldots$ has often proven much more difficult and is much less frequently attempted. We will present a robust Markov chain Monte Carlo (RMCMC) algorithm that allows the simultaneous determination of a few extremal eigenpairs of a very large matrix without the need to orthogonalize pairs of vectors to each other or store all the components of any vector.
Consider the following problem of calculating eigenvalues performed by,

$$
\begin{equation*}
A x=\lambda x \tag{1}
\end{equation*}
$$

Suppose $A \in R^{n \times n}$ is a symmetric matrix and also

$$
\lambda_{\min }=\lambda_{n}<\lambda_{n-1} \leq \cdots \leq \lambda_{2}<\lambda_{1}=\lambda_{\max }
$$

Consider the following Markov chain $T_{i}$ with length $i$ :

$$
T_{i}: k_{0} \rightarrow k_{1} \rightarrow \cdots \rightarrow k_{i}
$$

where $k_{j} \in\{1,2, \cdots, n\}$ for $j=1, \cdots, i$ are natural numbers. The statistical nature of constructing the above Markov chain follow as:

$$
p\left(k_{0}=\alpha\right)=p_{\alpha}, \quad p\left(k_{j}=\beta \mid k_{j-1}=\alpha\right)=p_{\alpha \beta}
$$

where $p_{\alpha}$ and $p_{\alpha \beta}$ show the probability of starting chain at $\alpha$ and transition probability from state $\alpha$ to $\beta$, respectively [3].
Now, define the random variable $W_{j}$ using the following recursion for

$$
\begin{equation*}
W_{j}=W_{j-1} \frac{a_{k_{j-1} k_{j}}}{p_{k_{j-1} k_{j}}}, \quad j=1,2, \ldots, i, \quad W_{0}=1 \tag{2}
\end{equation*}
$$

From all possible permissible densities, we choose the following

$$
\begin{aligned}
p_{\alpha} & =\frac{\left|h_{\alpha}\right|}{\sum_{\alpha=1}^{n}\left|h_{\alpha}\right|} \\
p_{\alpha \beta} & =\frac{\left|a_{\alpha \beta}\right|}{\sum_{\beta=1}^{n}\left|a_{\alpha \beta}\right|} \quad \alpha=1,2, \ldots, n
\end{aligned}
$$

where $h \in R^{n}$. It can be shown that [3] for a vector $f \in R^{n}$ and large value of $i$ :

$$
\begin{equation*}
\lambda_{1} \approx \frac{E\left[W_{i} f_{k_{i}}\right]}{E\left[W_{i-1} f_{k_{i-1}}\right]} \tag{3}
\end{equation*}
$$

Using deflation techniques $[7,8]$, once having the eigenpair $\left(\lambda_{i}, v_{i}\right)$, we can obtain eigenpairs $\left(\lambda_{i+1}, v_{i+1}\right), i=$ $1,2, \ldots, n$.
In [4] is shown that the accuracy of the eigenpairs has been soared by using RMCMC. In this paper, we use this algorithm for finding PCs.

Algorithm. Execute the following steps:
(1) Set $[M, N]=\operatorname{size}($ data $)$.
(2) Set $\mathrm{mn}=\operatorname{mean}($ data, 2$)$.
(3) Set data=data-repmat $(m n, 1, N)$.
(4) Compute COV=covariance(data).
(5) Compute $[\mathrm{PC}, \mathrm{V}]=\boldsymbol{R M C M C}$ algorithm (covariance).
(6) $\mathrm{V}=\operatorname{diag}(\mathrm{V})$.
(7) $\mathrm{V}=\mathrm{V}$ (rindices).
(8) $\mathrm{PC}=\mathrm{PC}(:$, rindices $)$.
(9) Signals $=\mathrm{PC}^{t} *$ data.
(10) Input color image $C$.
(11) Decompose $C$ to $R, G, B$.
(12) Call the MCMC-PCA algorithm for $R, G, B$.
(13) Output the compressed image $\hat{C}=R \bigoplus G \bigoplus B$.

## 4 Numerical experiments

We performed two images using PCA-MCMC and PCA-RMCMC algorithms and some of results for finding PC's are presented. In Table 1 and Figs. 1 and 2 we can see that the RMCMC algorithm gives better results than the MCMC algorithm. Also, the original images and their compressed images are shown in Figs 3-6. The numerical tests are made on $\operatorname{Intel}(\mathrm{R})$, Core(TM)i5 CPU, 2.5 GHz , personal machine using MATLAB software.

Table 1: Relative error for computing three eigenvalue by using MCMC and RMCMC algorithms

| $\lambda_{1}(M C M C)$ | $\lambda_{1}($ RMCMC $)$ | $\lambda_{2}($ MCMC $)$ | $\lambda_{2}($ RMCMC $)$ | $\lambda_{3}($ MCMC $)$ | $\lambda_{3}($ RMCMC $)$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $0.1600 \times 10^{-2}$ | $3.1289 \times 10^{-4}$ | $1.0526 \times 10^{-2}$ | $5.1689 \times 10^{-4}$ | $2.8726 \times 10^{-2}$ | $6.8916 \times 10^{-4}$ |

## 5 Conclusion

The paper presents a new algorithm for image compression based on PCA and RMCMC methods. The hybrid algorithm has enabled us to have better accuracy than the standard MCMC algorithm.

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Figure 1: Comparison of relative error between MCMC and RMCMC algorithms


Figure 2: Average PSNR vs. number of PCs by using MCMC and RMCMC algorithms


Figure 3: Original image, Size $=845941$ bytes


Figure 4: Compressed image with $q=20$ PCs, Size $=67910$ bytes


Figure 5: Original image, Size $=45,414$ bytes


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Figure 6: Compressed image with $q=20$ PCs, Size $=14,101$ bytes
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