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# Finding Solution to Linear System of Equations using Gibbs Sampling

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## Abstract

In this work, we explore Gibbs Sampling as a method to solve a linear system of equations evading computationally intensive matrix multiplications. We evaluate the proposed method using synthetic data generated randomly.

## 1 Introduction

There are many methods to solve a linear system of equations like the Matrix Inversion method [5] and Cramer's method [2], involves calculating the product of two matrices. Matrix Multiplication is computationally intensive since the complexity of multiplying two matrices, say  $A(m \times n)$  with  $B(n \times p)$  is of Order  $O(m \times n \times p)$ . So for two square matrices of Order  $n \times n$  is  $O(n^3)$ . This becomes very inefficient in large dimension matrices when using methods like matrix inversion to find the solution to the linear system of equations.

Methods like the steepest descent [6], conjugate gradient [6] and Jacobi Method [1] solve this issue by multiplying vectors instead of matrices which reduces the complexity to  $O(n^2)$ . In this work, we explore Gibbs Sampling as an alternative method to solve a linear system of equations evading computationally intensive matrix multiplications.

## 2 Method

Before solving the actual problem we tried using Gibbs sampling to sample from a Multi-Variate Normal(MVN) given mean and inverse of covariance matrix without finding the covariance matrix. Say we are dealing with  $n$ -dimensional multivariate which means Mean( $\mu$ ) is a  $n \times 1$  matrix and covariance ( $\Sigma$ ) is a symmetric positive definite matrix of order  $n \times n$  so  $\Sigma^{-1}$  say  $A$  is also symmetric and positive definite of order  $n \times n$ . Our matrix  $X$  is a vector  $[x_1, x_2, \dots, x_n]$ , we can break  $X$  into two parts  $X_1, X_2$  where  $X_2$  contains only one element whereas  $X_1$  contains  $X - X_2$  i.e all elements in  $X$  except  $X_2$ . Let  $X_1 \sim N(\mu_1, \Sigma_{11})$  and  $X_2 \sim N(\mu_2, \Sigma_{22})$  then we can split  $\Sigma$  into block matrices,

$$\Sigma = \left[ \begin{array}{c|c} \Sigma_{11} & \Sigma_{12} \\ \hline \Sigma_{21} & \Sigma_{22} \end{array} \right].$$

and let  $x = [x_1, x_2, \dots, x_{i-1}, x_{i+1}, \dots, x_n]$  then,

$$E[X_2|X_1 = x] = \mu_2 + \Sigma_{21}\Sigma_{11}^{-1}(x - \mu_1) \quad (1)$$

$$\Sigma_{22|1} = \Sigma_{22} - \Sigma_{21}\Sigma_{11}^{-1}\Sigma_{12} \quad (2)$$

But we don't have  $\Sigma$  we have only  $A$  so to get the block matrices of  $\Sigma$  from  $A$  we use Schur complement of matrix  $\Sigma$ . Schur Complement [7] of a sample block matrix  $P$ , below  $S = D - CA^{-1}B$

$$P = \begin{bmatrix} A & B \\ C & D \end{bmatrix}^{-1} = \begin{bmatrix} A^{-1} + A^{-1}BS^{-1}CA^{-1} & -A^{-1}BS^{-1} \\ -S^{-1}CA^{-1} & S^{-1} \end{bmatrix}.$$

Here in place of  $P$ , we have our  $\Sigma$ . We can use the matrix  $A = \Sigma^{-1}$  to get the terms  $\Sigma_{21}\Sigma_{11}^{-1}$  and  $\Sigma_{22} - \Sigma_{21}\Sigma_{11}^{-1}\Sigma_{12}$  directly without doing any matrix multiplications. After getting those terms we can update  $X_2$ ,

$$X_2 \sim N(E[X_2|X_1 = x], \Sigma_{22|1}) \quad (3)$$

and thus we will iteratively modify each dimension of  $X$  and generate our samples.

### 3 Experiments

Many experiments were done starting from comparing the steepest descent and conjugate gradient to find the solution of the linear system of equations to sampling from a multivariate normal distribution with known mean and inverse of the covariance matrix using Gibbs sampling without actually computing the inverse of the covariance matrix

#### 3.1 Experiment for comparing steepest descent and conjugate gradient

we compared the two algorithms in the number of iterations and time they take before converging to a solution with an error less than  $10^{-9}$  while solving the linear system of equations  $Ax = b$  where  $A, b, x_0$  (initial starting point) were generated randomly. For a fixed dimension, the value of  $A, b, x_0$  were kept the same for both the steepest descent and Conjugate Gradient algorithm. Here  $A$  is of Order  $n \times n$ ,  $b$  is of Order  $n \times 1$  and  $x$  is order  $n \times 1$

n	Steepest Descent		Conjugate Gradient	
	iterations	time (in sec)	iterations	time (in sec)
10	507682	10.5	10	$338\mu$
50	5419350	117	50	$1.61m$
100(sparse)	2490	$95.3m$	64	$16.5m$

Methods like steepest descent and conjugate gradient work for large dimensions but the conjugate gradient is better because it converges in less number of iterations and time, especially in the case of sparse matrices whereas the positive side of steepest descent is it can converge a bit closer to the actual value than conjugate gradient due to more number of iterations and time taken per iteration is less in steepest descent. Plain steepest descent takes many iterations but with certain optimizations, we can make the steepest descent converge in less number of iterations. [4]

#### 3.2 Experiment to sample from bivariate and draw the KDE contour plots

We sampled from a bivariate with a Mean( $\mu$ ) and covariance( $\Sigma$ ) using the Cholesky decomposition [3]. For any affine transformation of  $X \sim N(0, I)$  to  $Y = ZX + M$  then  $Y \sim N(M, Z.(Z.T))$  but here we want samples with Mean  $\mu$  so we keep  $M = \mu$  and covariance  $\Sigma$  so  $Z.(Z.T) = \Sigma$  which is nothing but Cholesky decomposition [3] of  $\Sigma$ . Hence here  $Z = \text{Cholesky}(\Sigma)$ . In this experiment for simplicity, we took Mean as zero,  $\mu = 0$ . We get circular contours for bivariate when the two random variables are uncorrelated. When they are correlated, we get elliptical contours.

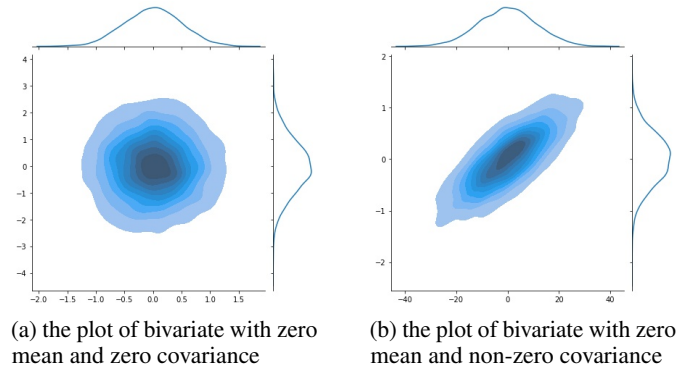


Figure 1: bivariate joint density KDE plots

Contour KDE plots of two gaussian random variables which are uncorrelated are circular. But when we add correlation between those two random variables, the KDE contour plots become elliptical. We can also say that each circular contour  $C$  with a radius  $r$  has its corresponding elliptical contour  $E$  related by  $E = LC$ , where  $L$  is the Cholesky decomposition [3] of  $\Sigma$

### 3.3 Experiment verifying the Gibbs sampler

The samples drawn from the Gibbs sampler were plotted to check whether they match with the KDE contour plots for a normal bivariate. We took 100,1000,10000 samples and plotted the respective KDE contour plots for all the samples. For the bivariate, we took Mean  $\mu = 0$  and the covariance matrix

$$\Sigma = \begin{bmatrix} 10.10549468 & 15.08509136 \\ 15.08509136 & 24.89129378 \end{bmatrix}.$$

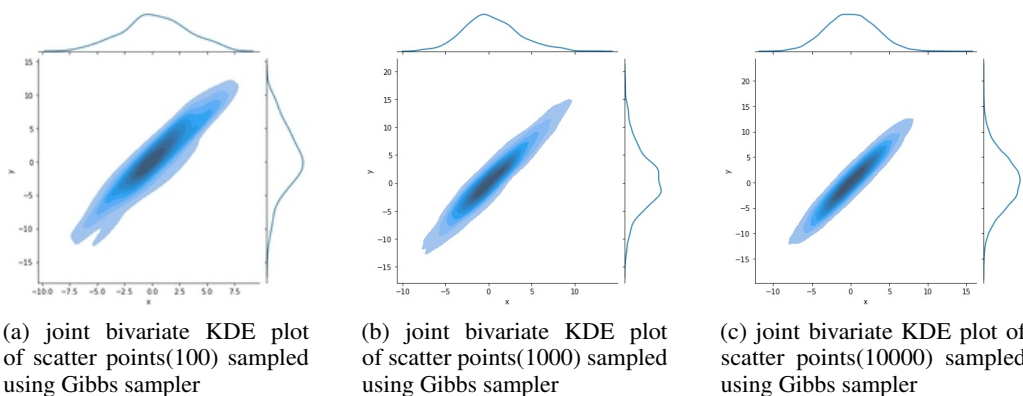


Figure 2: Gibbs sampler plots

The KDE contour plots for all the three plots match with the standard bivariate KDE plots which shows that our sampler is indeed generating samples from bivariate normal with a given distribution but the error in the covariance matrix generated from the samples drawn to the actual covariance matrix varies with the number of samples.

## References

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