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# Randomized Least Squares

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## Abstract

**This report aims to provide an overview of the Blendenpik algorithm, which is a randomized version of a least squares solver geared towards large scale, highly overdetermined systems of the form  $Ax = b$ . Theoretical and practical results will be used to illustrate the importance of coherence and row-mixing to enable a uniform sampling of the rows which guarantees high quality solutions with high probability.**

## 1 Background

The setting which is considered throughout this report is that of a highly overdetermined system  $Ax = b$ , with  $A \in \mathbb{R}^{m \times n}$ ,  $x \in \mathbb{R}^n$ ,  $b \in \mathbb{R}^m$ , and  $m \gg n$ . The aim is to obtain the optimal value of  $x$ , namely  $x^*$ , in the least squares sense i.e.  $x^* = \operatorname{argmin}_{x \in \mathbb{R}^n} \|b - Ax\|_2$ . Traditional least squares solvers such as LAPACK make use of a  $QR$  decomposition, where we can write:

$$A = Q \begin{bmatrix} R \\ 0 \end{bmatrix} = [Q_1 \quad Q_2] \begin{bmatrix} R \\ 0 \end{bmatrix} \quad Q \in \mathbb{R}^{m \times m}, Q_1 \in \mathbb{R}^{m \times n}, Q_2 \in \mathbb{R}^{m \times m-n}, R \in \mathbb{R}^{n \times n}$$

$Q$  is an orthogonal matrix (or unitary in case  $A$  is complex), i.e. its rows and columns are orthonormal:  $QQ^T = Q^TQ = I \implies Q^T = Q^{-1}$ .  $R$  is an upper triangular matrix (in this case because  $m > n$  and because we assume  $A$  is full rank). The linear system can thus be rewritten by using  $Ax = b \implies Q^T Ax = Q^T b$ , meaning:

$$\|x - Ab\|_2 = \|Q^T b - Q^T Ax\|_2 = \left\| \begin{bmatrix} Q_1^T b - Rx \\ Q_2^T b \end{bmatrix} \right\|$$

This holds because  $Q^T A = R$ . Thus minimizing in the least squares sense using the  $QR$  decomposition boils down to solving  $x = R^{-1}Q_1^T b$ . The total cost associated with solving the least squares problem in this manner consists of the cost of a  $QR$  decomposition of  $A$ , and the matrix inverse and multiplications to compute  $x$ . Since  $m$  is assumed to be large in these settings, especially given the increasing prevalence of big data applications, this can be extremely costly.

The objective of the Blendenpik algorithm is to apply uniform random sampling and random projections to define a preconditioner which reduces the number of number of iterations and thus computational cost associated with solving large scale systems using methods such as MINRES or LSQR. This requires careful consideration of the implications of random row sampling, as well as preprocessing steps to ensure a good solution can still be obtained with high probability.

A natural attempt at reducing the cost of solving an overdetermined system would be to sample a certain number of rows  $\mathcal{R}$  and obtain the solution  $x_{\mathcal{R}}^* = \operatorname{argmin}_{x \in \mathbb{R}^n} \|A_{\mathcal{R}}x - b_{\mathcal{R}}\|$ . Under certain conditions on  $A$  and  $\Omega(n \log(m) \log(n \log(m)))$  uniformly randomly sampled rows, this leads to a residual such that  $\|Ax_{\mathcal{R}} - b\|_2 / \|Ax_{opt} - b\|_2 \leq 1 + \epsilon$ . This has two problems (1):

1. Standard stability analysis of linear least squares algorithms is done in terms of backward error.

## 2. Running time depends on $\epsilon^{-1}$

An alternative approach is to define a preconditioner based on the uniformly sampled rows, denoted by  $A_{\mathcal{R}}$ , and use this in conjunction with a standard iterative method such as LSQR or MINRES. However, depending on the matrix  $A$ , the resulting subsampled matrix may be **ill-conditioned/ rank deficient**, leading to failure in the iterative subspace method. Such an approach works *only* if the coherence of the matrix is small. This can be enforced via a randomized row-mixing preprocessing phase. Before further elaborating on these randomized sampling and row-mixing approaches, it is necessary to define some terminology. This will then be used to derive bounds on the condition number of the preconditioned overdetermined system.

## 2 Key Concepts

### 2.1 Coherence

Coherence defines a notion of the degree to which the importance of the rows is concentrated onto a small number of rows. This is undesirable in the case of uniform sampling as we would need to sample a large number of rows to ensure these are picked. Therefore we prefer matrices with 'mixed' information, i.e. all rows are more or less equally important. Such matrices have a low coherence value (lower bounded by  $\frac{n}{m}$ ), whereas badly behaved matrices have high coherence values (upper bounded by 1). Given a matrix  $A \in \mathbb{R}^{m \times n}$  (where  $m > n$ ) and a matrix  $U$  with orthonormal columns that forms a basis of the column space of  $A$ , the coherence is defined as:

$$\mu(A) = \max_i \|U_{i,:}\|_2^2$$

**Lemma 1.** *The coherence  $\mu(A)$  of a matrix  $A \in \mathbb{R}^{m \times n}$  has the following bounds:*

$$\frac{n}{m} \leq \mu(A) \leq 1$$

*Proof.* First we prove the upper bound. We can define a basis with orthonormal columns by using the  $QR$  decomposition, namely  $A = QR = [Q_1 Q_2]R$ . Since  $Q$  is orthogonal, we know that the row norms are all equal to 1. Further, we only require the first  $n$  columns from  $Q$  to span the column space of  $A$ , which corresponds to the  $Q_1$  matrix. Consequently we see that the row norms are now *at most* 1, i.e.  $\mu(A) \leq 1$ . Now to prove the lower bound of the coherence, we consider the Frobenius norm of  $Q_1$ . The squared Frobenius norm of  $Q_1$  is  $\|Q_1\|_F^2 = \sum_{i \in [m], j \in [n]} Q_{1i,j}^2 = n$ , which follows from the fact that the vector 2-norm of each column is equal to 1. If  $\mu(A) < \frac{n}{m}$ , that is, the maximum squared row norm is less than  $\frac{n}{m}$ , then that implies that  $\|Q_1\|_F^2 < \sum_{i \in [m]} \frac{n}{m} = n$ , which is a contradiction as  $\|Q_1\|_F^2 = n$ . Thus we see that  $\mu(A) \geq \frac{n}{m}$  and conclude that  $\frac{n}{m} \leq \mu(A) \leq 1$ .  $\square$

Generally speaking the coherence number of randomly generated matrices is quite low. In a basic experiment that was performed using  $A_i \in \mathbb{R}^{1000}$ ,  $i \in [1000]$ , an average coherence of 0.072 was obtained, which is quite good seeing as the lower bound on such a matrix is  $\frac{50}{1000} = 0.05$ .

```
rand('state', 42);
n_matrices = 1000; n_rows = 1000; n_cols = 50;

% b) average coherences
coherences = [];
for n=1:n_matrices
    A = rand(n_rows, n_cols);
    if ~(rank(A) == n_cols)
        fprintf("rank deficient random matrix with rank: %i\n", rank(A));
    end
    [Q, R] = qr(A, 0);
    coherence = max(sum(Q.^2, 2));
    coherences = [coherences, coherence];
end
```

```
mean_coherence = mean(coherences);
fprintf("Mean coherence of random matrices: %d\n", mean_coherence);
```

In order to obtain the worst possible coherence, one can simply define a rank deficient matrix. For instance, in the case of the random matrices  $A_i \in \mathbb{R}^{1000 \times 50}$ , taking any one of these matrices and setting the first column to  $[0, \dots, 0]^T$  will yield a  $QR$  decomposition for which the first column of  $Q$  will be  $[1, 0, \dots, 0]^T$ , meaning the coherence is automatically equal to 1 i.e. the worst possible value.

## 2.2 Unitary Seed Transforms - Discrete Cosine Transform

Todo

## 2.3 Randomized Row-Mixing

Randomized row-mixing preprocessing phase to reduce coherence of a matrix. This enables us to apply uniform sampling to the mixed matrix, as this only works well for matrices with low coherence values.

Todo

## 3 Path to Blendenpik

Noting these definitions, we can formally prove bounds on a matrix  $A$  preconditioned via randomized techniques. General thought process of the procedure:

1. Given an overdetermined system  $Ax = b$
2. Don't know (or don't want to compute) its coherence number, but we still want to apply random uniform sampling approach to avoid computing  $QR$  decomposition of original  $A$  matrix. For this we need to define a good preconditioner which results in a good condition number  $\kappa(AR^{-1})$ .
3. To guarantee a good coherence number, we apply a unitary and randomized seed transform  $\mathcal{F} = FD$ , where  $F$  is a unitary matrix and  $D$  is a diagonal matrix with IID Rademacher random variables.
4. Given our incoherent (low coherence), randomly row-mixed matrix  $\mathcal{F}A$ , we can now apply uniform random row sampling and obtain a matrix  $\tilde{A}$  with a good condition number with high probability. Now we can compute the  $QR$  decomposition  $\tilde{A} = QR$  and use  $R$  as a preconditioner for  $A$  with a standard method like MINRES or LSQR.

### 3.1 Proofs

Since the point of the Blendenpik algorithm is to reduce the required number of iterations until convergence via a suitable preconditioner, it is useful to define an upper bound on the condition number of the preconditioned matrix  $\kappa(AR^{-1})$ , which is guaranteed with high probability.

**Theorem 2.** *Let  $A$  be an  $m \times n$  full rank matrix, and let  $S$  be a random sampling operator that samples  $r \geq n$  rows from  $A$  uniformly. Let  $\tau = C\sqrt{m\mu(A)}\log(r)/r$ , where  $C$  is a constant. Further assume  $\delta^{-1}\tau < 1$ . With probability at least  $1 - \delta$ , the sampled matrix  $SA$  is full rank, and if  $SA = QR$  is a reduced  $QR$  factorization of  $SA$ , we have*

$$\kappa(AR^{-1}) \leq \sqrt{\frac{1 + \delta^{-1}\tau}{1 - \delta^{-1}\tau}}$$

To prove this, we first consider Theorem 7 from (2).

**Theorem 3.** (2) *Suppose  $A \in \mathbb{R}^{m \times n}$ ,  $B \in \mathbb{R}^{n \times p}$ , and  $c \leq n$ . Construct  $C$  and  $R$  with [Algorithm 6](#), using the *EXPECTED*( $c$ ) algorithm. If the sampling probabilities  $\{p_i\}_{i=1}^n$  used by the algorithm are of the form (44) or (45), then*

$$\mathbb{E} [\|AB - CR\|_F] \leq \frac{1}{\sqrt{\beta c}} \|A\|_F \|B\|_F$$

If, in addition,  $B = A^T$ , then

$$\mathbb{E} [\|AA^T - CC^T\|_2] \leq O(1) \sqrt{\frac{\log c}{\beta c}} \|A\|_F \|A\|_2$$

We apply this theorem using  $A = U^T$ , where  $U \in \mathbb{R}^{m \times n}$  is a basis (with orthonormal columns) for the column space of  $A$ . We note that the sampling probabilities  $p_i$  satisfy the following inequality:

$$p_i \geq \beta \frac{\|A^{(i)}\|_2^2}{\|A\|_F^2} \iff \beta \leq p_i \frac{\|A\|_F^2}{\|A^{(i)}\|_2^2}$$

Consequently setting  $\beta = p_i \frac{\|A\|_F^2}{\mu(A)}$ , the above inequality is still satisfied as the coherence  $\mu(A)$  is the maximum squared row norm. By also setting  $p_i = n/m$  we obtain:

$$\begin{aligned} \mathbb{E} [\|AA^T - CC^T\|_2] &\leq O(1) \sqrt{\frac{\log c}{\beta c}} \|A\|_F \|A\|_2 \\ &= O(1) \sqrt{\frac{m\mu(A) \log c}{cn\|A\|_F^2}} \|A\|_F \|A\|_2 \\ &= O(1) \sqrt{\frac{m\mu(A) \log c}{c}} \frac{\|A\|_2}{\sqrt{n}} \\ &\leq O(1) \sqrt{\frac{m\mu(A) \log c}{c}} \\ &= \tau \end{aligned} \tag{1}$$

The final inequality uses the fact that  $\|A\|_2 = \max\{\|Ax\|_2 : \|x\|_2 = 1\} \leq \sqrt{n}$ , since the rows of  $A$  have unit norm. We can now apply Markov's inequality, which states that  $\mathbb{P}(X \geq a) \leq \mathbb{E}[X]/a$ , where  $X$  is a non-negative random variable and  $a > 0$  a non-negative constant. In particular, we consider the random variable  $\|I_{n \times n} - \frac{m}{r} U_S^T U_S\|_2$ , set  $a = \delta^{-1} \tau$ , and obtain the following:

$$\mathbb{P}(\|I_{n \times n} - \frac{m}{r} U_S^T U_S\|_2 \geq \delta^{-1} \tau) \leq \frac{\mathbb{E}[\|I_{n \times n} - \frac{m}{r} U_S^T U_S\|_2]}{\delta^{-1} \tau} \leq \delta \tag{2}$$

If  $\mathbb{P}(X \geq a) \leq \mathbb{E}[X]/a$ , this implies that  $\mathbb{P}(X < a) = 1 - \mathbb{P}(X \geq a) \geq 1 - \mathbb{E}[X]/a$ . Consequently we have that:

$$\mathbb{P}(\|I_{n \times n} - \frac{m}{r} U_S^T U_S\|_2 < \underbrace{\delta^{-1} \tau}_{< 1}) \geq 1 - \delta$$

Thus both the above inequality is satisfied and  $A_S$  is full rank with probability  $1 - \delta$ . Now we can make use of Theorem 1 from (3), which states the following.

**Theorem 4.** (3) Suppose that  $l, m$ , and  $n$  are positive integers such that  $m \geq l \geq n$ . Suppose further that  $A$  is a full-rank  $m \times n$  matrix, and that the SVD of  $A$  is

$$A_{m \times n} = U_{m \times n} \Sigma_{n \times n} V_{n \times n}^*$$

Suppose in addition that  $T$  is an  $l \times m$  matrix such that the  $l \times n$  matrix  $TU$  has full rank. Then, there exist an  $n \times n$  matrix  $P$ , and an  $l \times n$  matrix  $Q$  whose columns are orthonormal, such that

$$T_{l \times m} A_{m \times n} = Q_{l \times n} P_{n \times n}.$$

Furthermore, if  $P$  is any  $n \times n$  matrix, and  $Q$  is any  $l \times n$  matrix whose columns are orthonormal, such that  $P$  and  $Q$  satisfy the latter equation, then the condition numbers of  $AP^{-1}$  and  $TU$  are equal.

Consequently we have that  $\kappa(AR^{-1}) = \kappa(U_S)$ . Thus using the bound derived from the Markov inequality above, we can derive a bound on the condition number of the preconditioned matrix  $AR^{-1}$ . **We are dealing with symmetric matrices, so we can use the fact that  $\lambda_{max} = \max_{x \in \mathbb{R}^n: \|x\|=1} x^T A x$ , which holds analogously for the minimum eigenvalue. We want to use this to derive a bound on  $\kappa(U_S)$ .** We proceed by using the Rayleigh quotient, namely  $R(A, x) = \frac{x^T A x}{x^T x}$ , or equivalently  $R(A, x)$  is applied to  $\frac{m}{r} U_S^T U_S$ . We obtain the following:

$$\begin{aligned}
R\left(\frac{m}{r} U_S^T U_S, x\right) &= \frac{x^T \left(\frac{m}{r} U_S^T U_S\right) x}{x^T x} \\
&= \frac{x^T (I_{n \times n} + \frac{m}{r} U_S^T U_S - I_{n \times n}) x}{x^T x} \\
&= \frac{x^T x + x^T \left(\frac{m}{r} U_S^T U_S - I_{n \times n}\right) x}{x^T x} \\
&= 1 - \underbrace{R(I_{n \times n} - \frac{m}{r} U_S^T U_S, x)}_M
\end{aligned} \tag{3}$$

We note that  $M$  is symmetric, meaning that the Rayleigh quotient  $R(M, x) \in [\lambda_{min}, \lambda_{max}]$ , and since  $\|M\|_2 < \delta^{-1} \tau$  (with probability  $1 - \delta$ ), it holds that  $|R(M, x)| < \delta^{-1} \tau$ , which implies that the eigenvalues of  $\frac{m}{r} U_S^T U_S$  are contained in the interval  $[1 - \delta^{-1} \tau, 1 + \delta^{-1} \tau]$ . This allows us to bound the condition number of  $U_S$  and thus the preconditioned matrix  $AR^{-1}$ . In particular we have that:

$$\begin{aligned}
\kappa(U_S) &= \frac{\sigma_{max}(U_S)}{\sigma_{min}(U_S)} \\
&= \sqrt{\frac{\lambda_{max}(U_S^T U_S)}{\lambda_{min}(U_S^T U_S)}} \\
&= \sqrt{\frac{\lambda_{max}(\frac{m}{r} U_S^T U_S)}{\lambda_{min}(\frac{m}{r} U_S^T U_S)}}
\end{aligned} \tag{4}$$

Note that the last inequality holds as scaling a matrix  $A$  by some constant  $c$  leads to the same eigenvalues but scaled by  $c$  i.e.  $\lambda_i(cA) = c\lambda_i(A)$ , which does not influence the ratio of the maximum and minimum eigenvalues. In addition we have used the fact that  $\sigma_i^2(U_S) = \lambda_i(U_S^T U_S)$ , which we have just shown to be contained in the interval  $[1 - \delta^{-1} \tau, 1 + \delta^{-1} \tau]$ . Finally, we can conclude the bound on the condition number of the preconditioned matrix.

$$\kappa(AR^{-1}) = \kappa(U_S) \leq \sqrt{\frac{1 + \delta^{-1} \tau}{1 - \delta^{-1} \tau}}$$

### 3.2 Algorithm

We apply the Blendenpik algorithm using two different iterative solvers, namely MINRES and LSQR. In the case of MINRES we are required to use the normal equations  $A^T A x = A^T b$  as a symmetric, square matrix is required, unlike for LSQR. **Not clear on which type of preconditioning required for each approach. Should also probably indicate stopping criterion used internally by matlab. Also not sure if we are correctly applying preconditioner for minres case (see ch4 part 4.9 on when the left and split preconditioned matrices have the same condition number) - preconditioner for minres has to be SPD**

## 4 Experiments

We define matrices  $A_1$  with:

```
rng(11);
U = orth(rand(20000,400));
```

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**Algorithm 1** Blendenpik Algorithm (DCT)

---

$x = \text{blendenpik}(A \in \mathbb{R}^{m \times n}, b \in \mathbb{R}^n, \gamma)$

$\tilde{m} \leftarrow \lceil m/1000 \rceil \times 1000$

$M \leftarrow \begin{bmatrix} A \\ 0 \end{bmatrix} \in \mathbb{R}^{\tilde{m} \times n}$

while true

1.  $M \leftarrow DCT(DM)$ , where  $D \in \mathbb{R}^{\tilde{m} \times \tilde{m}}$  is diagonal with  $D_{ii} = \pm 1$  with equal probability

2. Define diagonal matrix  $S \in \mathbb{R}^{\tilde{m} \times \tilde{m}}$  with  $S_{ii} = \begin{cases} 1 & \text{with probability } \gamma n / \tilde{m} \\ 0 & \text{with probability } 1 - \gamma n / \tilde{m} \end{cases}$

3. Compute reduced  $QR$  decomposition  $SM = QR$  with  $R \in \mathbb{R}^{n \times n}$

4. if  $\kappa_{est}(R)^{-1} > 5\epsilon_{machine}$

(a)  $x \leftarrow \begin{cases} \text{MINRES applied to } A^T A, A^T b \text{ with } R^{-T} \text{ and } R^{-1} \text{ as left and right preconditioners} \\ \text{LSQR applied to } A, b \text{ using } R \text{ as a preconditioner} \end{cases}$

(b) **return**

5. **endif**

**end while**

---

```
S = diag(linspace(1,1e5,400));  
V = orth(rand(400));  
A1 = U*S*V';
```

Matrix  $A_2$  is given by:

```
rng(11);  
A2 = [ diag(linspace(1,e5,400)); zeros(19600,400) ];  
A2 = A2 + 1e-8*ones(20000,400);
```

In the following figures we show the required number of iterations to converge for different values of  $\gamma$ . The  $b$  vector is randomly initialized. **Note that we only implement one single row-mixing step, although it may be beneficial to implement more (refer to pg 13 of blendenpik paper)**

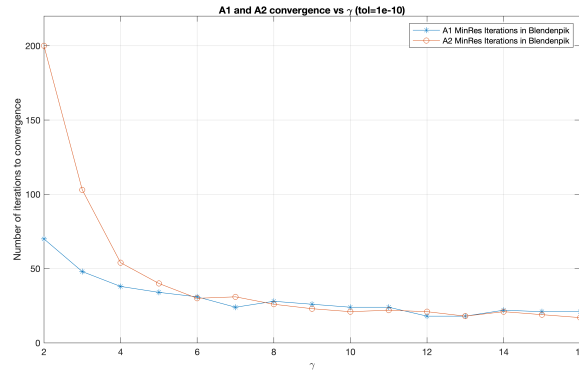


Figure 1: Convergence of  $A_1$  and  $A_2$  as function of  $\gamma$

The following figure investigates the convergence of the inner LSQR/MINRES steps for the two matrices defined above. **Generate these plots. Can we just use the residual resulting from the lsqr/minres methods? Which gamma value do we use?... lsqr returns LSVEC which is pretty close to what is indicated in the y axis**

of figure 5.6 but not exactly the same, is this ok? Not sure how we can easily extract the residual vector at each iteration to compute the indicated norm. MinRes doesn't have LSVEC...

## References

- [1] P. Drineas, M. W. Mahoney, S. Muthukrishnan, and T. Sarlós, “Faster least squares approximation,” *CoRR*, vol. abs/0710.1435, 2007. [Online]. Available: <http://arxiv.org/abs/0710.1435>
- [2] P. Drineas, M. Mahoney, and S. Muthukrishnan, “Relative-error *cur* matrix decompositions,” *SIAM Journal on Matrix Analysis and Applications*, vol. 30, pp. 844–881, 05 2008.
- [3] V. Rokhlin and M. Tygert, “A fast randomized algorithm for overdetermined linear least-squares regression,” *Proceedings of the National Academy of Sciences of the United States of America*, vol. 105, pp. 13 212–7, 09 2008.

## A Appendix

Optionally include extra information (complete proofs, additional experiments and plots) in the appendix. This section will often be part of the supplemental material.