Equilibrating low-rank approximations with Gaussian priors

&

High-performance finite DPP sampling via mirror-image Cholesky

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Google Research

ELSI Conference, August 2018

Overview

1 Equilibrating low-rank approximations with Gaussian priors

2 High-performance finite DPP sampling via mirror-image Cholesky

Recommender systems and language models often involve low-rank approximations of a large, sparse matrix A, e.g., a local minimum of:

$$\mathcal{L}(X,Y) = \frac{1}{2} \|W \circ (A - XY^*)\|_F^2 + \frac{\lambda}{2} \left(\|X\|_F^2 + \|Y\|_F^2 \right),$$

where W is a weighting matrix (often a function of A).¹

This is Maximum Likelihood inference with $(XY^*)_{i,j} \sim \mathcal{N}(A_{i,j}, W_{i,j}^{-2})$ and priors $X_{i,j}, Y_{i,j} \sim \mathcal{N}(0, 1/\lambda)$.

One can find an approximate local minimum via a few iterations of Weighted Alternating Least Squares.³

¹See, for example, [Hu et al.-2008] Collaborative filtering for implicit feedback datasets

²Cf. [Srebro/Jaakkola-2003] Weighted low-rank approximations ³http://www.tensorflow.org/api_docs/python/tf/contrib/

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Definition 1. Given $S \in \operatorname{Sym}(n,\mathbb{R})$, we will use the shorthand P(S) for the linear operator $P(S) : \operatorname{Sym}(n,\mathbb{R}) \to \operatorname{Sym}(n,\mathbb{R})$ via P(S)A = SAS.

Definition 2. The **geometric mean** of $A, B \in S_{++}^n$ is $A \sharp B = B \sharp A = P(A^{1/2})(P(A^{-1/2})B)^{1/2}$.

Proposition 1. For any $A, B \in S_{++}^n$, there is a unique $S \in S_{++}^n$ such that P(S)A = B.⁴

Proof. For existence, put $S = A^{-1} \sharp B$.

For uniqueness, if P(S)A = P(T)A, then $X^*AX = A$, with $X = T^{-1}S$. Then the spectral decomposition $(S^{1/2}T^{-1}S^{1/2})(S^{1/2}Z) = (S^{1/2}Z)\Lambda$ implies $XZ = Z\Lambda$, $\Lambda \succ 0$. And $Z^*AZ = Z^*(X^*AX)Z = \Lambda Z^*AZ\Lambda$, so $\Lambda = I$ and T = S. \square

⁴[Anderson/Trapp-1980] Operator means and electrical networks, Cf [Bhatia-2007] Positive Definite Matrices

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Lemma 4 (P.). Given $(X,Y) \in \mathbb{R}^{m \times r} \times \mathbb{R}^{n \times r}$, $S \in S_{++}^n$ minimizes $f: S_{++}^n \to \mathbb{R}_+$, where

$$f(S) = ||XS||_F^2 + ||YS^{-1}||_F^2,$$

iff $P(S)(X^*X) = P(S^{-1})(Y^*Y)$. And, if X and Y have full column rank, then $S = ((X^*X)^{-1} \sharp (Y^*Y))^{1/2}$ is the unique minimizer.

Proof. Decompose f as $g \circ h$, where $h: S_{++}^n \to S_{++}^n$ via $h(S) = S^2$ an $g: S_{++}^n \to \mathbb{R}_+$ via $g(T) = \langle X^*X, T \rangle + \langle Y^*Y, T^{-1} \rangle$.

Then h is a diffeomorphism and $dg_T: (T_TS_{++}^n \cong \operatorname{Sym}(n,\mathbb{R})) \to (T_{g(T)}\mathbb{R} \cong \mathbb{R})$ via $dg_T(dT) = \langle X^*X - T^{-1}Y^*YT^{-1}, dT \rangle$.

So $S \in S_{++}^n$ is a critical point of f iff $df_S = dg_{S^2} \circ dh_S = 0$ iff $X^*X - S^{-2}Y^*YS^{-2} = 0$. \square

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Theorem 5 (P.). If $\ell: \mathbb{R}^{m \times n} \to \mathbb{R}$ is continuous, the local minima of $\mathcal{L}: \mathbb{R}^{m \times r} \times \mathbb{R}^{n \times r} \to \mathbb{R}$, where

$$\mathcal{L}(X, Y) = \ell(XY^*) + \frac{\lambda}{2} \left(||X||_F^2 + ||Y||_F^2 \right),$$

satisfy $X^*X = Y^*Y$. And, given any candidate (X,Y), the **equilibration**, $(XS^{1/2},YS^{-1/2})$, where $S = (X^*X)^{-1} \sharp (Y^*Y)$, minimizes the regularization while preserving the input to ℓ .

Proof. Given (X, Y), $\ell(XY^*)$ is invariant under any transformation $(X, Y) \mapsto (XZ, YZ^{-*})$ where $Z \in GL(n, \mathbb{R})$. Thus, any local minimum must satisfy

$$||X||_F^2 + ||Y||_F^2 = \min_{Z \in GL(n,\mathbb{R})} \{||XZ||_F^2 + ||YZ^{-*}||_F^2\}$$
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where we exploited the polar decomposition Z = SQ, Q unitary. The result then follows from our lemma. \square

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$$\mathcal{L}(X, Y) = \ell(XY^*) + \frac{\lambda}{2} \left(||X||_F^2 + ||Y||_F^2 \right),$$

insert an equilibration step between each block coordinate descent step. E.g., if X and Y have full column rank, replace

$$(X, Y) \mapsto (XS^{1/2}, YS^{-1/2}), \quad S = (X^*X)^{-1} \sharp (Y^*Y),$$

which can be computed in $O((m+n+r)r^2)$ time.

Equilibration is essentially free and keeps the regularization minimized (with the constraint of preserving the loss function input).

If one thinks of (X^*X, Y^*Y) as analogous to a primal/dual pair in an SDP IPM, this is similar to centering the Newton step about the NT point.

Equilibration has a much more pronounced effect for small regularization values.

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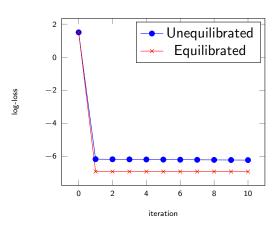
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Equilibration has a much more pronounced effect for small regularization values

A trivial example

Consider minimizing $(\alpha - \chi \eta)^2 + \lambda(\chi^2 + \eta^2)$ given $\alpha = 1$, $\lambda = 0.001$, $\chi_0 = \eta_0 = 2$.



The Nesterov-Todd equilibration obviously makes assumptions about the invertibility of the Gramians.

Geometrically, $S = A \sharp B$, when $A, B \in S_{++}^n$, is well-known to be the Euclidean midpoint between $\log(A)$ and $\log(B)$ and the midpoint of the geodesic between A and B when S_{++}^n is equipped with the left-invariant metric $g_X(S,T) = \langle X^{-1}S, X^{-1}T \rangle$.

One could extend the geometric mean to the boundary via:

$$A \sharp B = \lim_{\epsilon \downarrow 0} (A + \epsilon I) \sharp (B + \epsilon I).$$

But this extension is discontinuous [Bhatia-2007]: Let

$$A = \begin{pmatrix} 4 & 0 \\ 0 & 1 \end{pmatrix}, B = \begin{pmatrix} 20 & 6 \\ 6 & 2 \end{pmatrix}, X_n = \begin{pmatrix} 1 & 0 \\ 0 & 1/n \end{pmatrix} \rightarrow X = \begin{pmatrix} 1 & 0 \\ 0 & 0 \end{pmatrix}.$$

$$\lim_{n \uparrow \infty} \Phi_n(A) \sharp \Phi_n(B) = \lim_{n \uparrow \infty} \Phi_n(A \sharp B) = \Phi(A \sharp B) = \begin{pmatrix} 0 & 0 \\ 0 & 0 \end{pmatrix},$$
$$\left(\lim_{n \uparrow \infty} \Phi_n(A)\right) \sharp \left(\lim_{n \uparrow \infty} \Phi_n(B)\right) = \Phi(A) \sharp \Phi(B) = \begin{pmatrix} \sqrt{80} & 0 \\ 0 & 0 \end{pmatrix}.$$

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$$A = \begin{pmatrix} 4 & 0 \\ 0 & 1 \end{pmatrix}, B = \begin{pmatrix} 20 & 6 \\ 6 & 2 \end{pmatrix}, X_n = \begin{pmatrix} 1 & 0 \\ 0 & 1/n \end{pmatrix} \rightarrow X = \begin{pmatrix} 1 & 0 \\ 0 & 0 \end{pmatrix}.$$

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The Nesterov-Todd equilibration obviously makes assumptions about the invertibility of the Gramians.

Geometrically, $S = A \sharp B$, when $A, B \in S_{++}^n$, is well-known to be the Euclidean midpoint between $\log(A)$ and $\log(B)$ and the midpoint of the geodesic between A and B when S_{++}^n is equipped with the left-invariant metric $g_X(S,T) = \langle X^{-1}S, X^{-1}T \rangle$.

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We thus saw that the extension:

$$A \sharp B = \lim_{\epsilon \downarrow 0} (A + \epsilon I) \sharp (B + \epsilon I)$$

can lead to singular geometric means (in addition to being discontinuous).

But if we only care about **backwards stability**, then there is no issue. One can compute $S = \widehat{X^*X}^{-1} \sharp \widehat{Y^*Y}$, where $\widehat{Z} = Z + \alpha \|Z\|_F$ for some $\alpha \ll 1$, equilibrate with S, and perhaps repeat.

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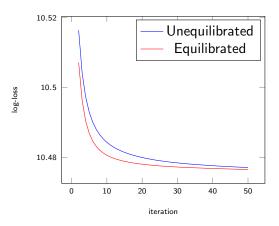
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Another toy example

Consider minimizing $||A - XY^*||_F^2 + \lambda(||X||_F^2 + ||Y||_F^2)$, given A = randn(200, 400), $\lambda = 0.1$, $X_0 = \text{randn}(200, 10)$, $Y_0 = [\text{randn}(400, 9), \text{zeros}(400, 1)]$.



Jordan-algebraic interpretations

Recall our definition P(S): Sym $(n, \mathbb{R}) \to \text{Sym}(n, \mathbb{R})$ via P(S)A = SAS.

This is a special case of the quadratic representation of a **Jordan algebra** V, where $P(x) = 2L(x)^2 - L(x^2)$ and $L(x) : V \to V$ is left application of $x \in V$.

For $V = \operatorname{Sym}(n, \mathbb{R})$ with Jordan product $A \circ B \equiv \frac{1}{2}(AB + BA)$, $L(A)B \equiv A \circ B$.

$$P(A)B = 2(A \circ (A \circ B)) - A^{2} \circ B = ABA.$$

The 1-to-1 correspondence between symmetric cones and squares of Euclidean Jordan algebras [Faraut/Koranyi-1998] is commonly exploited in Interior Point Methods (especially for Lorentz cones).⁷

One can easily build on Prop'n 1 to show: given $A, B \in \text{int}(V^2)$, there is a unique $S \in \text{int}(V^2)$ such that $P(S)A = B.^8$ The definitions of geometric means and Nesterov-Todd scaling points carry over through usage of P.

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Determinantal Point Processes

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Definition 7. A (finite) **Determinantal Point Process (DPP)** is a random variable **Y** over the power set of $\mathcal{Y} = \{1,...,k\} \subset \mathbb{N}$ generated by a $k \times k$ marginal kernel matrix K via the rule

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[Gillenwater-2014] reduced the factored elementary DPP sampling down to cubic complexity via what is equivalent to diagonally-pivoted Cholesky.¹⁰

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Conditioning and Schur complements

Proposition 2. Given disjoint subsets $A, B \subset \mathcal{Y}$,

$$P[B \subseteq \mathbf{Y}|A \subseteq \mathbf{Y}] = \det(K_B - K_{B,A}K_A^{-1}K_{A,B}),$$

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Proof. The first claim follows from

$$\det(K_{A\cup B}) = \det(K_A)\det(K_B - K_{B,A}K_A^{-1}K_{A,B})$$

and

$$P[B \subseteq \mathbf{Y}|A \subseteq \mathbf{Y}] = \frac{\det(K_{A \cup B})}{\det(K_A)}.$$

The second claim follows from applying the first result to the complementary DPP to find

$$P[B \subseteq \mathbf{Y}^c | A \subseteq \mathbf{Y}^c] = \det((I - K)_B - K_{B,A}(I - K)_A^{-1} K_{A,B}).$$

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Sampling w/ mirror-image Cholesky

```
samples = {}
for j=1:n
  J2 = [j+1:n]
  keep_index = Bernoulli(K(j,j))
  if keep_index
    scale = -1; samples.insert(j)
    K(j,j) = sqrt(K(j,j))
  else
    scale = +1
    K(j,j) = sqrt(1-K(j,j))
  K(J2,j) /= K(j,j)
  K(J2,J2) += scale*tril(K(J2,j)*K(J2,j)')
```

This is a small tweak of unblocked Cholesky factorization; the majority of the work is in Hermitian rank-1 updates. And the standard Cholesky optimizations apply (e.g., blocking and sparse-direct factorization)!

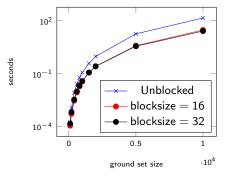
Sampling w/ mirror-image Cholesky

```
samples = \{\}
for j=1:n
  J2 = [j+1:n]
  keep_index = Bernoulli(K(j,j))
  if keep_index
    scale = -1; samples.insert(j)
    K(i,i) = sqrt(K(i,i))
  else
    scale = +1
    K(j,j) = sqrt(1-K(j,j))
 K(J2,j) /= K(j,j)
  K(J2,J2) += scale*tril(K(J2,j)*K(J2,j)')
```

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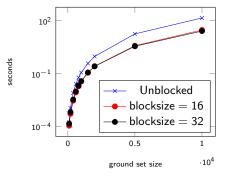
Blocked mirror-image sampling

```
samples = {}
J1_beg = 1
while J1_beg <= n
  J1_end = min(n, J1_beg+blocksize-1)
  J1 = [J1\_beg:J1\_end]; J2 = [J1\_end+1:n]
  J1\_samples, K(J1,J1) = sample(K(J1,J1))
  A21 = zeros(len(J2), len(J1_samples))
  B21 = zeros(len(J2), len(J1)-len(J1_samples))
  num_keep_packed = num_drop_packed = 0
 for k in J1
    K(J2,k) /= K(k,k)
    if (k-J1_beg+1) in J1_samples
      A21(:,num\_keep\_packed++) = K(J2,k); scale = -1
    else
      B21(:,num\_drop\_packed++) = K(J2,k); scale = +1
    J1R = [k+1:J1\_end]
    K(J2,J1R) += scale*K(J2,k)*K(J1R,k)
  K(J2,J2) += tril(B21*B21' - A21*A21')
  J1\_beg = J1\_end + 1
```



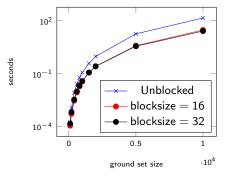
HPC dense Cholesky implementations can be trivially modified.

Maximum Likelihood inference and elementary DPP sampling are similar but involve diagonal pivoting; the former uses the largest diagonal and the latter samples from the PDF implied by the diagonal. One can modify a blocked dense diagonally-pivoted Cholesky.



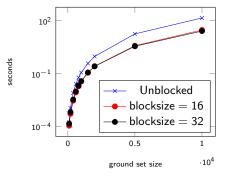
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Acknowledgements/Questions/Comments

Acknowledgements:

- Rasmus Larsen and John Anderson:
 For introducing me to the WALS problem.
- Steffen Rendle:
 For noticing that the Gramians were equal.
- Matt Knepley and Sameer Agarwal:
 For pointing out the gauge transformation analogy.
- Alex Kulesza and Jenny Gillenwater:
 For answering DPP sampling questions.

Questions/comments?