# A Randomized Algorithm for Approximating the Log Determinant of a Symmetric Positive Definite Matrix

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LogDet Problem

Given an SPD matrix  $A \in \mathbb{R}^{n \times n}$ , compute (exactly or approximately) log det (A).

**Additive Error Approximation** 

**Relative Error Approximation** 

Let  $\mathbf{A} \in \mathbb{R}^{n \times n}$  be an SPD matrix. For any  $\alpha$  with  $\lambda_1(\mathbf{A}) < \alpha$ , define  $\mathbf{B} = \mathbf{A}/\alpha$  and  $\mathbf{C} = \mathbf{I}_n - B$ . Then,

Let  $\mathbf{A} \in \mathbb{R}^{n \times n}$  be an SPD matrix whose eigenvalues lie in the interval  $(\theta_1, 1)$ , for some  $0 < \theta_1 < 1$ . Let  $\mathbf{C} = \mathbf{I}_n - A$ . Then,

$$\log \det(A) = n \log(\alpha) - \sum_{k=1}^{\infty} \frac{\operatorname{Tr}\left(\log\left(\mathbf{C}^{\kappa}\right)\right)}{k}.$$

- (  $\sim l_{a})$ 

$$\log \det(A) = -\sum_{k=1}^{\infty} \frac{\operatorname{Tr}\left(\log\left(\mathbf{C}^{\kappa}\right)\right)}{k}.$$

## Algorithm 1

Input: A ∈ ℝ<sup>n×n</sup>, accuracy parameter ε > 0, integer m > 0.
Compute an estimate to the largest eigenvalue of A,λ<sub>1</sub>(A), using the Power Method.

2 C = I<sub>n</sub> - A/(7λ<sub>1</sub>(A)).
3 Create p = [20 log(2/δ)/ε<sup>2</sup>] i.i.d random Gaussian vectors, g<sub>1</sub>, g<sub>2</sub>,..., g<sub>p</sub>.
4 Estimate Σ<sub>k=1</sub><sup>∞</sup> Tr(log(C<sup>k</sup>))/k with a truncated Taylor Series type randomized trace estimator that computes Σ<sub>k=1</sub><sup>m</sup> (1/p Σ<sub>i=1</sub><sup>p</sup> g<sub>i</sub><sup>T</sup> C<sup>k</sup> g<sub>i</sub>).

Let  $\widehat{\log \det}(\mathbf{A})$  be the  $\log \det$  approximation of the above procedure. Then, we prove that with probability at least  $1 - 2\delta$ ,

 $|\widehat{\log \det}(\mathbf{A}) - \log \det(A)| \le 2\epsilon\Gamma$ where  $\Gamma = \sum_{i=1}^{n} \log \left(7 \cdot \frac{\lambda_1(\mathbf{A})}{\lambda_i(\mathbf{A})}\right)$  and  $m \ge \lceil 7\kappa(\mathbf{A}) \log(\frac{1}{\epsilon}) \rceil$ .

## Algorithm 2

Input: A ∈ ℝ<sup>n×n</sup>, accuracy parameter ε > 0, integer m > 0.
C = I<sub>n</sub> - A
Create p = [20 log(2/δ)/ε<sup>2</sup>] i.i.d random Gaussian vectors, g<sub>1</sub>, g<sub>2</sub>,..., g<sub>p</sub>.
Estimate Σ<sup>∞</sup><sub>k=1</sub> Tr(log(C<sup>k</sup>))/k with a truncated Taylor Series type randomized trace estimator that computes Σ<sup>m</sup><sub>k=1</sub> (<sup>1</sup>/<sub>p</sub>Σ<sup>p</sup><sub>i=1</sub> g<sup>T</sup><sub>i</sub>C<sup>k</sup>g<sub>i</sub>)

Let  $\widehat{\log \det}(\mathbf{A})$  be the  $\log \det$  approximation of the above procedure on inputs  $\mathbf{A}$  and  $\epsilon$ . Then, we **prove** that with probability at least  $1 - \delta$ ,  $\widehat{|\log \det}(\mathbf{A}) - \log \det(A)| \le 2\epsilon \cdot |\log \det(\mathbf{A})|.$ 

### **Intensive Computational Kernel:**

- Exact computation complexity  $\mathcal{O}(n^3)$ , prohibitive for

Big Data!!!

Approximation using Algorithm 1:

 $\mathcal{O}(nnz(A) \cdot (\mathbf{A}) \cdot (m\epsilon^{-2} + \log(n))).$ 

**Real world applications:** Multivariate Statistics (e.g. Computation of Maximum Likelihood), Spatial-Temporal (e.g. GIS, GPS e.t.c.), Data Mining (e.g. Classification of data) e.t.c.

### **More RandNLA Techniques**

**Krylov Subspace Methods:** Given a matrix  $\mathbf{A} \in \mathbb{R}^{m \times n}$  and a starting guess matrix  $\mathbf{X} \in \mathbb{R}^{n \times s}$ , we want to use the block Krylov space

Table 1: Real-World data from University of Florida Sparse Matrix Collection and C++ Parallel Implementation. (Names: thermal2, ecology2, Idoor, thermomech\_TC, boneS01)

	$\log \det(\mathbf{A})$			time (sec)		
n	exact	approx		ovact	approx	m
		mean	std	CACU	mean	
1228045	1.3869e6	1.3928e6	964.79	31.28	31.24	149
999999	3.3943e6	3.403e6	1212.8	18.5	10.47	125
952203	1.4429e7	1.4445e7	1683.5	117.91	17.60	33
102158	-546787	-546829.4	553.12	57.84	2.58	77
127224	1.1093e6	1.106e6	247.14	130.4	8.48	125

We **prove** that with probability at least 3/4 the sparse principal component is close to the actual principal component and its 2-norm is close to 1. We **demonstrate** applications on real world data, e.g. Genomics, Text Clustering.

 $\mathcal{K}_q(\mathbf{A}\mathbf{A}^{\top}, \mathbf{A}\mathbf{X})$  to approximate the left singular vector space of  $\mathbf{A}$ . We prove:

- Spectral & Frobenius bounds for the distance between the approximate and the actual space.
- Quality measurements of the bounds relative to the best low-rank approximation.

**Sparse Principal Component Analysis:** We relax the Sparse PCA problem to a **problem with convex constraints (but non-convex objective)**:

maximize  $\mathbf{w}^{\top} \mathbf{X}^{\top} \mathbf{X} \mathbf{w}$ subject to  $\|\mathbf{w}\|_{1} \leq \sqrt{k}, \|\mathbf{w}\|_{2} \leq 1, \mathbf{w} \in \mathbb{R}^{n}$ .

We **design** a two-phase algorithm that first approximates the relaxed problem and then uses a randomized rounding strategy to sparsify the approximation.

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