RANDOMIZED NYSTRÖM PRECONDITIONING*

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Abstract. This paper introduces the Nyström PCG algorithm for solving a symmetric positivedefinite linear system. The algorithm applies the randomized Nyström method to form a low-rank approximation of the matrix, which leads to an efficient preconditioner that can be deployed with the conjugate gradient algorithm. Theoretical analysis shows that preconditioned system has constant condition number as soon as the rank of the approximation is comparable with the number of effective degrees of freedom in the matrix. The paper also develops adaptive methods that provably achieve similar performance without knowledge of the effective dimension. Numerical tests show that Nyström PCG can rapidly solve large linear systems that arise in data analysis problems, and it surpasses several competing methods from the literature.

Key words. Conjugate gradient, cross-validation, kernel method, linear system, Nyström approximation, preconditioner, randomized algorithm, regularized least-squares, ridge regression.

AMS subject classifications. 65F08, 68W20, 65F55, 65F22

1. Motivation. In their elegant 1997 textbook on numerical linear algebra [32], Trefethen and Bau write,

"In ending this book with the subject of preconditioners, we find ourselves at the philosophical center of the scientific computing of the future... Nothing will be more central to computational science in the next century than the art of transforming a problem that appears intractable into another whose solution can be approximated rapidly. For Krylov subspace matrix iterations, this is preconditioning... we can only guess where this idea will take us."

The next century has since arrived, and one of the most fruitful developments in matrix computations has been the emergence of new algorithms that use randomness in an essential way. This paper explores a topic at the nexus of preconditioning and randomized numerical linear algebra. We will show how to use a randomized matrix approximation algorithm to construct a preconditioner for an important class of linear systems that arises throughout data analysis and scientific computing.

1.1. The Preconditioner. Consider the regularized linear system

(1.1) $(A + \mu I)x = b$ where $A \in \mathbb{R}^{n \times n}$ is symmetric psd and $\mu \ge 0$.

Here and elsewhere, psd abbreviates the term "positive semidefinite." This type of linear system emerges whenever we solve a regularized least-squares problem. We will design a class of preconditioners for the problem (1.1).

Throughout this paper, we assume that we can access the matrix A through matrix–vector products $x \mapsto Ax$, commonly known as *matvecs*. The algorithms that

^{*}Submitted to the editors DATE.

Funding: ZF and MU were supported by NSF Award IIS-1943131, the ONR Young Investigator Program, and the Alfred P. Sloan Foundation. JAT was supported by ONR BRC Award N00014-18-1-2363 and NSF FRG Award 1952777.

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we develop will economize on the number of matvecs, and they may not be appropriate in settings where matvecs are very expensive or there are cheaper ways to interact with the matrix.

For a rank parameter $\ell \in \mathbb{N}$, the randomized Nyström approximation of A takes the form

(1.2)
$$\hat{A}_{nys} = (A\Omega)(\Omega^T A\Omega)^{\dagger} (A\Omega)^T$$
 where $\Omega \in \mathbb{R}^{n \times \ell}$ is standard normal.

This matrix provides the best psd approximation of A whose range coincides with the range of the sketch $A\Omega$. The randomness in the construction ensures that \hat{A}_{nys} is a good approximation to the original matrix A with high probability [19, Sec. 14].

We can form the Nyström approximation with sketch size ℓ , using ℓ matvecs with A, plus some extra arithmetic. See Algorithm 2.1 for the implementation details.

Given the eigenvalue decomposition $\hat{A}_{nys} = U \hat{\Lambda} U^T$ of the randomized Nyström approximation, we construct the Nyström preconditioner:

(1.3)
$$P = \frac{1}{\hat{\lambda}_{\ell} + \mu} U(\hat{\Lambda} + \mu I)U^T + (I - UU^T).$$

In a slight abuse of terminology, we refer to ℓ as the rank of the Nyström preconditioner. The key point is that we can solve the linear system Py = c very efficiently, and the action of P^{-1} dramatically reduces the condition number of the regularized matrix $A_{\mu} = A + \mu I$.

We propose to use (1.3) in conjunction with the preconditioned conjugate gradient (PCG) algorithm. Each iteration of PCG involves a single matvec with A, and a single linear solve with P. When the preconditioned matrix $P^{-1}A_{\mu}$ has a modest condition number, the algorithm converges to a solution of (1.1) very quickly. See Algorithm 5.1 for pseudocode for Nyström PCG.

The randomized Nyström preconditioner (1.3) was suggested by P.-G. Martinsson in the survey [19, Sec. 17], but it has not been implemented or analyzed.

1.2. Guarantees. This paper contains the first comprehensive study of the preconditioner (1.3), including theoretical analysis and testing on prototypical problems from data analysis and machine learning. One of the main contributions is a rigorous method for choosing the rank ℓ to guarantee good performance, along with an adaptive rank selection procedure that performs well in practice.

A key quantity in our analysis is the *effective dimension* of the regularized matrix $A + \mu I$. That is,

(1.4)
$$d_{\text{eff}}(\mu) = \text{tr}\left(A(A+\mu I)^{-1}\right) = \sum_{j=1}^{n} \frac{\lambda_j(A)}{\lambda_j(A) + \mu}.$$

The effective dimension measures the degrees of freedom of the problem after regularization. It may be viewed as a (smoothed) count of the eigenvalues larger than μ . Many real-world matrices exhibit strong spectral decay, so the effective dimension is typically much smaller than the nominal dimension n. As we will discuss, the effective dimension also plays a role in a number of machine learning papers [1, 2, 4, 7, 17] that consider randomized algorithms for solving regularized linear systems.

Our theory tells us the randomized Nyström preconditioner P is successful when its rank ℓ is proportional to the effective dimension.

THEOREM 1.1 (Randomized Nyström Preconditioner). Let $A \in \mathbb{S}_n^+(\mathbb{R})$ be a psd matrix, and write $A_{\mu} = A + \mu I$ where the regularization parameter $\mu > 0$. Define the

effective dimension $d_{\text{eff}}(\mu)$ as in (1.4). Construct the randomized preconditioner P from (1.2) and (1.3) with rank parameter $\ell = 2 \lceil 1.5 d_{\text{eff}}(\mu) \rceil + 1$. Then the condition number of the preconditioned system satisfies

(1.5)
$$\mathbb{E}\left[\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})\right] < 28.$$

Theorem 1.1 is a restatement of Theorem 5.1.

Simple probability bounds follow from (1.5) via Markov's inequality. For example,

$$\mathbb{P}\left\{\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le 56\right\} > 1/2.$$

The main consequence of Theorem 1.1 is a convergence theorem for PCG with the randomized Nyström preconditioner.

COROLLARY 1.2 (Nyström PCG: Convergence). Construct the preconditioner P as in Theorem 1.1, and condition on the event { $\kappa_2(P^{-1/2}A_{\mu}P^{-1/2}) \leq 56$ }. Solve the regularized linear system (1.1) using Nyström PCG, starting with an initial iterate $x_0 = 0$. After t iterations, the relative error δ_t satisfies

$$\delta_t \coloneqq \frac{\|x_t - x_\star\|_{\text{PCG}}}{\|x_\star\|_{\text{PCG}}} < 2 \cdot (0.77)^t \quad \text{where } A_\mu x_\star = b.$$

The error norm is defined as $||u||_{PCG}^2 = u^T (P^{-1/2} A_\mu P^{-1/2}) u$. In particular, $t \geq [3.9 \log(2/\epsilon)]$ iterations suffice to achieve relative error ϵ .

Although Theorem 1.1 gives an interpretable bound for the rank ℓ of the preconditioner, we cannot instantiate it without knowledge of the effective dimension. To address this shortcoming, we have designed adaptive methods for selecting the rank in practice (subsection 5.4).

Finally, as part of our investigation, we will also develop a detailed understanding of Nyström sketch-and-solve, a popular algorithm in the machine learning literature [1, 4]. Our analysis highlights the deficiencies of Nyström sketch-and-solve relative to Nyström PCG.

1.3. Example: Ridge Regression. As a concrete example, we consider the ℓ^2 regularized least-squares problem, also known as ridge regression. This problem takes the form

(1.6)
$$\operatorname{minimize}_{x \in \mathbb{R}^d} \quad \frac{1}{2n} \|Gx - b\|^2 + \frac{\mu}{2} \|x\|^2,$$

where $G \in \mathbb{R}^{n \times d}$ and $b \in \mathbb{R}^n$ and $\mu > 0$. By calculus, the solution to (1.6) also satisfies the regularized system of linear equations

(1.7)
$$\left(\frac{1}{n}G^{T}G + \mu I\right)x = \frac{1}{n}G^{T}b.$$

A direct method to solve (1.7) requires $O(nd^2)$ flops, which is prohibitive when n and d are both large. Instead, when n and d are large, iterative algorithms, such as the conjugate gradient method (CG), become the tools of choice. Unfortunately, the ridge regression linear system (1.7) is often very ill-conditioned, and CG converges very slowly.

Nyström PCG can dramatically accelerate the solution of (1.7). As an example, consider the shuttle-rf dataset (subsection 6.2). The matrix G has dimension 43,300 × 10,000, while the preconditioner is based on a Nyström approximation with rank $\ell = 800$. Figure 1 shows the progress of the residual as a function of the iteration count. Nyström PCG converges to machine precision in 13 iterations, while CG stalls.

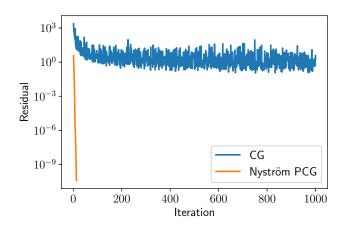


Fig. 1: Ridge regression: CG versus Nyström PCG. For the shuttle-rf data set, Nyström PCG converges to machine precision in 13 iterations while CG stalls. See subsections 1.3 and 6.2.

1.4. Comparison to prior randomized preconditioners. Prior proposals for randomized preconditioners [3, 20, 25] accelerate the solution of highly overdetermined or underdetermined least-squares problems using the sketch-and-precondition paradigm [19, Sec. 10]. For $n \ge d$, these methods require $\Omega(d^3)$ computation to factor the preconditioner. In contrast, the randomized Nyström preconditioner applies to any symmetric positive-definite linear system and can be significantly faster for regularized problems. See subsection 5.2.2 more details.

1.5. Roadmap. Section 2 contains an overview of the Nyström approximation and its key properties. Section 3 studies the role of the Nyström approximation in estimating the inverse of the regularized matrix. We analyze the Nyström sketchand-solve method in Section 4, and we give a rigorous performance bound for this algorithm. Section 5 presents a full treatment of Nyström PCG, including theoretical results and guidance on numerical implementation. Computational experiments in Section 6 demonstrate the power of Nyström PCG for three different applications involving real data sets.

1.6. Notation. We write $\mathbb{S}_n(\mathbb{R})$ for the linear space of $n \times n$ real symmetric matrices, while $\mathbb{S}_n^+(\mathbb{R})$ denotes the convex cone of real psd matrices. The symbol \leq denotes the Loewner order on $\mathbb{S}_n(\mathbb{R})$. That is, $A \leq B$ if and only if the eigenvalues of B - A are all nonnegative. The function $\operatorname{tr}[\cdot]$ returns the trace of a square matrix. The map $\lambda_j(A)$ returns the *j*th largest eigenvalue of A; we may omit the matrix if it is clear. As usual, κ_2 denotes the ℓ^2 condition number. We write ||M|| for the spectral norm of a matrix M. For a psd matrix A, we write $||u||_A^2 = u^T A u$ for the A-norm. Given $A \in \mathbb{S}_n(\mathbb{R})$ and $1 \leq \ell \leq n$, the symbol $\lfloor A \rfloor_{\ell}$ refers to any best rank- ℓ approximation to A relative to the spectral norm. For $A \in \mathbb{S}_n^+(\mathbb{R})$ and $\mu \geq 0$, the regularized matrix is abbreviated $A_{\mu} = A + \mu I$. For $A \in \mathbb{S}_n^+(\mathbb{R})$ and $\mu > 0$ effective dimension of A_{μ} is defined as $d_{\text{eff}}(\mu) = \operatorname{tr}(A(A + \mu I))^{-1}$. For $A \in \mathbb{S}_n^+(\mathbb{R})$, the *p*-stable rank of A is defined as $\operatorname{sr}_p(A) = \lambda_p^{-1} \sum_{j>p}^n \lambda_j$. For $A \in \mathbb{S}_n^+(\mathbb{R})$, we denote the time taken to compute a matvec with A by T_{mv} .

Algorithm 2.	l Rano	lomized	Ν	yström	Ap	proximation	[18,	33	
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Input: Positive-semidefinite matrix $A \in \mathbb{S}_n^+(\mathbb{R})$, rank ℓ **Output:** Nyström approximation in factored form $\hat{A}_{nys} = U\hat{\Lambda}U^T$

1: $\Omega = \operatorname{randn}(n, \ell)$	\triangleright Gaussian test matrix
2: $\Omega = \operatorname{qr}(\Omega, 0)$	\triangleright Thin QR decomposition
3: $Y = A\Omega$	$\triangleright \ell$ matvecs with A
4: $\nu = \operatorname{eps}(\operatorname{norm}(Y, \operatorname{'fro'}))$	\triangleright Compute shift
5: $Y_{\nu} = Y + \nu \Omega$	\triangleright Shift for stability
6: $C = \operatorname{chol}(\Omega^T Y_{\nu})$	
7: $B = Y_{\nu}/C$	
8: $[U, \Sigma, \sim] = \operatorname{svd}(B, 0)$	\triangleright Thin SVD
9: $\hat{\Lambda} = \max\{0, \Sigma^2 - \nu I\}$	$\triangleright \operatorname{Remove \ shift, \ compute \ eigs}$

2. The Nyström approximation. Let us begin with a review of the Nyström approximation and the randomized Nyström approximation.

2.1. Definition and basic properties. The Nyström approximation is a natural way to construct a low-rank psd approximation of a psd matrix $A \in \mathbb{S}_n^+(\mathbb{R})$. Let $X \in \mathbb{R}^{n \times \ell}$ be an arbitrary test matrix. The *Nyström approximation* of A with respect to the range of X is defined by

(2.1)
$$A\langle X \rangle = (AX)(X^T A X)^{\dagger} (AX)^T \in \mathbb{S}_n^+(\mathbb{R}).$$

The Nyström approximation is the best psd approximation of A whose range coincides with the range of AX. It has a deep relationship with the Schur complement and with Cholesky factorization [19, Sec. 14].

The Nyström approximation enjoys several elementary properties that we record in the following lemma.

LEMMA 2.1. Let $A\langle X \rangle \in \mathbb{S}_n^+(\mathbb{R})$ be a Nyström approximation of the psd matrix $A \in \mathbb{S}_n^+(\mathbb{R})$. Then

- 1. The approximation $A\langle X \rangle$ is psd and has rank at most ℓ .
- 2. The approximation $A\langle X \rangle$ depends only on range(X).
- 3. In the Loewner order, $A\langle X \rangle \preceq A$.
- 4. In particular, the eigenvalues satisfy $\lambda_j(\hat{A}) \leq \lambda_j(A)$ for each $1 \leq j \leq n$.

The proof of Lemma 2.1, Item 3 is not completely obvious. It is a consequence of the fact that we may express $\hat{A}_{nys} = A^{1/2} \Pi A^{1/2}$, where Π is an orthogonal projector.

2.2. Randomized Nyström approximation. How should we choose the test matrix X so that the Nyström approximation $A\langle X \rangle$ provides a good low-rank model for A? Surprisingly, we can obtain a good approximation simply by drawing the test matrix at random. See [33] for theoretical justification of this claim.

Let us outline the construction of the randomized Nyström approximation. Draw a standard normal test matrix $\Omega \in \mathbb{R}^{n \times \ell}$ where ℓ is the sketch size, and compute the sketch $Y = A\Omega$ By Lemma 2.1, the sketch size ℓ is equal to the rank of \hat{A}_{nys} with probability 1, hence we use these terms interchangeably. The Nyström approximation (2.1) is constructed directly from the test matrix Ω and the sketch Y:

(2.2)
$$\hat{A}_{nvs} = A \langle \Omega \rangle = Y (\Omega^T Y)^{\dagger} Y^T.$$

The formula (2.2) is not numerically sound. We refer the reader to Algorithm 2.1 for a stable and efficient implementation of the randomized Nyström approximation [18, 33]. Conveniently, Algorithm 2.1 returns the truncated eigendecomposition $\hat{A}_{nys} = U\hat{\Lambda}U^T$, where $U \in \mathbb{R}^{n \times \ell}$ is an orthonormal matrix whose columns are eigenvectors and $\hat{\Lambda} \in \mathbb{R}^{\ell \times \ell}$ is a diagonal matrix listing the eigenvalues, which we often abbreviate as $\hat{\lambda}_1, \ldots, \hat{\lambda}_{\ell}$.

The randomized Nyström approximation described in this section has a key difference from the Nyström approximations that have traditionally been used in the machine learning literature [1, 4, 9, 13, 35]. In machine learning settings, the Nyström approximation is usually constructed from a sketch Y that samples random columns from the matrix (i.e., the random test matrix Ω has 1-sparse columns). In contrast, Algorithm 2.1 computes a sketch Y via random projection (i.e., the test matrix Ω is standard normal). In most applications, we have strong reasons (subsection 2.2.3) for preferring random projections to column sampling.

2.2.1. Cost of randomized Nyström approximation. Throughout the paper, we write $T_{\rm mv}$ for the time required to compute a matrix–vector product (matvec) with A. Forming the sketch $Y = A\Omega$ with sketch size ℓ requires ℓ matvecs, which costs $T_{\rm mv}\ell$. The other steps in the algorithm have arithmetic cost $O(n\ell^2)$. Hence, the total computational cost of Algorithm 2.1 is $O(T_{\rm mv}\ell + \ell^2 n)$ operations. The storage cost is $O(\ell n)$ floating-point numbers.

For Algorithm 2.1, the worst-case performance occurs when A is dense and unstructured. In this case, forming Y costs $O(n^2 \ell)$ operations. However, if we have access to the columns of A then we may reduce the cost of forming Y to $O(n^2 \log \ell)$ by using a structured test matrix Ω , such as a scrambled subsampled randomized Fourier transform (SSRFT) map or a sparse map [19, 33].

2.2.2. A priori guarantees for the randomized Nyström approximation. In this section, we present an a priori error bound for the randomized Nyström approximation. The result improves over previous analyses [13, 14, 33] by sharpening the error terms. This refinement is critical for the analysis of the preconditioner.

PROPOSITION 2.2 (Randomized Nyström approximation: Error). Consider a psd matrix $A \in \mathbb{S}_n^+(\mathbb{R})$ with eigenvalues $\lambda_1 \geq \cdots \geq \lambda_n$. Choose a sketch size $\ell \geq 4$, and draw a standard normal test matrix $\Omega \in \mathbb{R}^{n \times \ell}$. Then the rank- ℓ Nyström approximation \hat{A}_{nys} computed by Algorithm 2.1 satisfies

$$(2.3) \quad \mathbb{E}\|A - \hat{A}_{\text{nys}}\| \le \min_{2 \le p \le \ell - 2} \left[\left(1 + \frac{2(\ell - p)}{p - 1}\right) \lambda_{\ell - p + 1} + \frac{2e^2\ell}{p^2 - 1} \left(\sum_{j > \ell - p} \lambda_j\right) \right].$$

The proof of Proposition 2.2 may be found in subsection SM1.1.

Proposition 2.2 shows that, in expectation, the randomized Nyström approximation \hat{A}_{nys} provides a good rank- ℓ approximation to A. The first term in the bound is comparable with the spectral-norm error $\lambda_{\ell-p+1}$ in the optimal rank- $(\ell - p)$ approximation, $\lfloor A \rfloor_{\ell-p}$. The second term in the bound is comparable with the trace-norm error $\sum_{j>\ell-p} \lambda_j$ in the optimal rank- $(\ell - p)$ approximation.

Proposition 2.2 is better understood via the following simplification.

COROLLARY 2.3 (Randomized Nyström approximation). Instate the assumptions of Proposition 2.2. For $p \ge 2$ and $\ell = 2p - 1$, we have the bound

$$\mathbb{E}\|A - \hat{A}_{nys}\| \le \left(3 + \frac{4e^2}{p} \operatorname{sr}_p(A)\right) \lambda_p.$$

NYSTRÖM PCG

The p-stable rank, $\operatorname{sr}_p(A) = \lambda_p^{-1} \sum_{j=p}^n \lambda_j$, reflects decay in the tail eigenvalues.

Corollary 2.3 shows that the Nyström approximation error is on the order of λ_p when the rank parameter $\ell = 2p - 1$. The constant depends on the *p*-stable rank $\operatorname{sr}_p(A)$, which is small when the tail eigenvalues decay quickly starting at λ_p . This bound is critical for establishing our main results (Theorems 4.2 and 5.1).

2.2.3. Random projection versus column sampling. Most papers in the machine learning literature [1, 4] construct Nyström approximations by sampling columns at random from an adaptive distribution. In contrast, for most applications, we advocate using an oblivious random projection of the matrix to construct a Nyström approximation.

Random projection has several advantages over column sampling. First, column sampling may not be practical when we only have black-box matvec access to the matrix, while random projections are natural in this setting. Second, it can be very expensive to obtain adaptive distributions for column sampling. Indeed, computing approximate ridge leverage scores costs just as much as solving the ridge regression problem directly using random projections [10, Theorem 2]. Third, even with a good sampling distribution, column sampling produces higher variance results than random projection, so it is far less reliable.

On the other hand, we have found that there are a few applications where it is more effective to compute a randomized Nyström preconditioner using column sampling in lieu of random projections. In particular, this seems to be the case for kernel ridge regression (subsection 6.5). Indeed, the entries of the kernel matrix are given by an explicit formula, so we can extract full columns with ease. Sampling ℓ columns may cost only $O(\ell n)$ operations, whereas a single matvec generally costs $O(n^2)$. Furthermore, kernel matrices usually exhibit fast spectral decay, which limits the performance loss that results from using column sampling in lieu of random projection.

3. Approximating the regularized inverse. Let us return to the regularized linear system (1.1). The solution to the problem has the form $x_{\star} = (A + \mu I)^{-1}b$. Given a good approximation \hat{A} to the matrix A, it is natural to ask whether $\hat{x} = (\hat{A} + \mu I)^{-1}b$ is a good approximation to the desired solution x_{\star} .

There are many reasons why we might prefer to use \hat{A} in place of A. In particular, we may be able to solve linear systems in the matrix $\hat{A} + \mu I$ more efficiently. On the other hand, the utility of this approach depends on how well the inverse $(\hat{A} + \mu I)^{-1}$ approximates the desired inverse $(A + \mu I)^{-1}$. The next result addresses this question for a wide class of approximations that includes the Nyström approximation.

PROPOSITION 3.1 (Regularized inverses). Consider psd matrices $A, A \in \mathbb{S}_n^+(\mathbb{R})$, and assume that the difference $E = A - \hat{A}$ is psd. Fix $\mu > 0$. Then

(3.1)
$$\|(\hat{A} + \mu I)^{-1} - (A + \mu I)^{-1}\| \le \frac{1}{\mu} \frac{\|E\|}{\|E\| + \mu}.$$

Furthermore, the bound (3.1) is attained when $\hat{A} = \lfloor A \rfloor_{\ell}$ for $1 \leq \ell \leq n$.

The proof of Proposition 3.1 may be found in subsection SM1.1.1. It is based on [5, Lemma X.1.4].

Proposition 3.1 has an appealing interpretation. When $||A - \hat{A}||$ is small in comparison to the regularization parameter μ , then the approximate inverse $(\hat{A} + \mu I)^{-1}$ can serve in place of the inverse $(A + \mu I)^{-1}$. Note that $||(A + \mu I)^{-1}|| \leq 1/\mu$, so we can view (3.1) as a relative error bound.

Algorithm 4.1 Nyström sketch-and-solve

Input: Psd matrix $A \in \mathbb{S}_n^+(\mathbb{R})$, right-hand side *b*, regularization μ , rank ℓ **Output:** Approximate solution \hat{x} to (1.1)

[U, Λ] = RandomizedNyströmApproximation(A, ℓ)
 Use (4.1) to compute x̂ = (Â_{nvs} + μI)⁻¹b

4. Nyström sketch-and-solve. The simplest mechanism for using the Nyström approximation is an algorithm called Nyström sketch-and-solve. This section introduces the method, its implementation, and its history. We also provide a general theoretical analysis that sheds light on its performance. In spite of its popularity, the Nyström sketch-and-solve method is rarely worth serious consideration.

4.1. Overview. Given a rank- ℓ Nyström approximation \hat{A}_{nys} of the psd matrix A, it is tempting to replace the regularized linear system $(A + \mu I)x = b$ with the proxy $(\hat{A}_{nys} + \mu I)x = b$. Indeed, we can solve the proxy linear system in $O(\ell n)$ time using the Sherman–Morrison–Woodbury formula [15, Eqn. (2.1.4)]:

LEMMA 4.1 (Approximate regularized inversion). Consider any rank- ℓ matrix \hat{A} with eigenvalue decomposition $\hat{A} = U\hat{\Lambda}U^T$. Then

(4.1)
$$(\hat{A} + \mu I)^{-1} = U(\hat{\Lambda} + \mu I)^{-1}U^T + \frac{1}{\mu}(I - UU^T).$$

We refer to the approach in this paragraph as the Nyström sketch-and-solve algorithm because it is modeled on the sketch-and-solve paradigm that originated in [27].

See Algorithm 4.1 for a summary of the Nyström sketch-and-solve method. The algorithm produces an approximate solution \hat{x} to the regularized linear system (1.1) in time $O(T_{\rm mv}\ell + \ell^2 n)$. The arithmetic cost is much faster than a direct method, which costs $O(n^3)$. It can also be faster than running CG for a long time at a cost of $O(T_{\rm mv})$ per iteration. The method looks attractive if we only consider the runtime, and yet...

Nyström sketch-and-solve only has one parameter, the rank ℓ of the Nyström approximation, which controls the quality of the approximate solution \hat{x} . When $\ell \ll n$, the method has an appealing computational profile. As ℓ increases, the approximation quality increases but the computational burden becomes heavy. We will show that, alas, an accurate solution to the linear system actually requires $\ell \approx n$, at which point the computational benefits of Nyström sketch-and-solve evaporate completely.

In summary, Nyström sketch-and-solve is almost never the right algorithm to use. We will see that Nyström PCG generally produces much more accurate solutions with a similar computational cost.

4.2. Guarantees and deficiencies. Using Proposition 3.1 together with the a priori guarantee in Proposition 2.2, we quickly obtain a performance guarantee for Algorithm 4.1.

THEOREM 4.2. Fix $p \geq 2$, and set $\ell = 2p - 1$. For a psd matrix $A \in \mathbb{S}_n^+(\mathbb{R})$, construct a randomized Nyström approximation \hat{A}_{nys} using Algorithm 2.1. Then the approximation error for the inverse satisfies

(4.2)
$$\mathbb{E} \| (A + \mu I)^{-1} - (\hat{A}_{nys} + \mu I)^{-1} \| \le \left(3 + \frac{4e^2}{p} \operatorname{sr}_p(A) \right) \frac{\lambda_p}{\mu \cdot (\lambda_p + \mu)}.$$

Define $x_{\star} = (A + \mu I)^{-1}b$, and select $\ell = 2 \lceil 1.5 d_{\text{eff}}(\epsilon \mu) \rceil + 1$. Then the approximate solution \hat{x} computed by Algorithm 4.1 satisfies

(4.3)
$$\mathbb{E}\left[\frac{\|\hat{x} - x_\star\|_2}{\|x_\star\|_2}\right] \le 26\epsilon$$

The proof of Theorem 4.2 may be found in Appendix A.1.

Theorem 4.2 tells us how accurately we can hope to solve linear systems using Nyström sketch-and-solve (Algorithm 4.1). To obtain relative error ϵ , we should choose $\ell = O(d_{\text{eff}}(\epsilon\mu))$. When $\epsilon\mu$ is small, we anticipate that $d_{\text{eff}}(\epsilon\mu) \approx n$. In this case, Nyström sketch-and-solve has no computational value. Our analysis is sharp in its essential respects, so the pessimistic assessment is irremediable.

4.3. History. Nyström sketch-and-solve has a long history in the machine learning literature. It was introduced in [35] to speed up kernel-based learning, and it plays a role in many subsequent papers on kernel methods. In this context, the Nyström approximation is typically obtained using column sampling [1, 4, 35], which has its limitations (subsection 2.2.3). More recently, Nyström sketch-and-solve has been applied to speed up approximate cross-validation [31].

The analysis of Nyström sketch-and-solve presented above differs from previous analysis. Prior works [1, 4] focus on the kernel setting, and they use properties of column sampling schemes to derive learning guarantees. In contrast, we bound the relative error for a Nyström approximation based on a random projection. Our overall approach extends to column sampling if we replace Proposition 2.2 by an appropriate analog, such as Gittens's results [13].

5. Nyström Preconditioned Conjugate Gradients. We now present our main algorithm, Nyström PCG. This algorithm produces high accuracy solutions to a regularized linear system by using the Nyström approximation \hat{A}_{nys} as a preconditioner. We provide a rigorous estimate for the condition number of the preconditioned system, and we prove that Nyström PCG leads to fast convergence for regularized linear systems. In contrast, we have shown that Nyström sketch-and-solve cannot be expected to yield accurate solutions.

5.1. The preconditioner. In this section, we introduce the optimal low-rank preconditioner, and we argue that the randomized Nyström preconditioner provides an approximation that is easy to compute.

5.1.1. Motivation. As a warmup, suppose we knew the eigenvalue decomposition of the best rank- ℓ approximation of the matrix: $\lfloor A \rfloor_{\ell} = V_{\ell} \Lambda_{\ell} V_{\ell}^{T}$. How should we use this information to construct a good preconditioner for the regularized linear system (1.1)?

Consider the family of symmetric psd matrices that act as the identity on the orthogonal complement of range(V_{ℓ}). Within this class, we claim that the following matrix is the *optimal preconditioner*:

(5.1)
$$P_{\star} = \frac{1}{\lambda_{\ell+1} + \mu} V_{\ell} (\Lambda_{\ell} + \mu I) V_{\ell}^{T} + (I - V_{\ell} V_{\ell}^{T}).$$

The optimal preconditioner P_{\star} requires $O(n\ell)$ storage, and we can solve linear systems in P_{\star} in $O(n\ell)$ time. Whereas the regularized matrix A_{μ} has condition number $\kappa_2(A_{\mu}) = (\lambda_1 + \mu)/(\lambda_n + \mu)$, the preconditioner yields

(5.2)
$$\kappa_2(P_{\star}^{-1/2}A_{\mu}P_{\star}^{-1/2}) = \frac{\lambda_{\ell+1}+\mu}{\lambda_n+\mu}.$$

Algorithm 5.1 Nyström PCG

Input: Psd matrix A, righthand side b, initial guess x_0 , regularization parameter μ , sketch size ℓ , solution tolerance η

Output: Approximate solution \hat{x} to regularized system (1.1)

1: $[U, \hat{\Lambda}] = \text{RandomizedNyströmApproximation}(A, \ell)$ 2: $r_0 = b - (A + \mu I)x_0$ 3: $z_0 = P^{-1} r_0$ \triangleright using (5.3) 4: $p_0 = z_0$ 5: while $||r||_2 > \eta$ do $v = (A + \mu I)p_0$ 6: $\alpha = (r_0^T z_0) / (p_0^T v_0)$ 7: ▷ compute step size $x = x_0 + \alpha p_0$ ▷ update solution 8: $r = r_0 - \alpha v$ ▷ update residual 9: $z=P^{-1}r$ \triangleright find search direction via (5.3) 10: $\beta = (r^T z) / (r_0^T z_0)$ 11: 12: $x_0 \leftarrow x, r_0 \leftarrow r, p_0 \leftarrow z + \beta p_0, z_0 \leftarrow z$

This is the minimum possible condition number attainable by a preconditioner from the class that we have delineated. It represents a significant improvement when $\lambda_{\ell+1} \ll \lambda_1$. The proofs of these claims are straightforward; for details, see subsection SM1.1.2.

5.1.2. Randomized Nyström preconditioner. It is expensive to compute the best rank- ℓ approximation $[A]_{\ell}$ accurately. In contrast, we can compute the rank- ℓ randomized Nyström approximation \hat{A}_{nys} efficiently (Algorithm 2.1). Furthermore, we have seen that \hat{A}_{nys} approximates A nearly as well as the optimal rank- ℓ approximation (Corollary 2.3). These facts lead us to study the randomized Nyström preconditioner, proposed in [19, Sec. 17] without a complete justification.

Consider the eigenvalue decomposition $\hat{A}_{nys} = U \hat{\Lambda} U^T$, and write $\hat{\lambda}_{\ell}$ for its ℓ th eigenvalue. The randomized Nyström preconditioner and its inverse take the form

(5.3)
$$P = \frac{1}{\hat{\lambda}_{\ell} + \mu} U(\hat{\Lambda} + \mu I) U^{T} + (I - UU^{T});$$
$$P^{-1} = (\hat{\lambda}_{\ell} + \mu) U(\hat{\Lambda} + \mu I)^{-1} U^{T} + (I - UU^{T}).$$

Like the optimal preconditioner P_{\star} , the randomized Nyström preconditioner (5.3) is cheap to apply and to store. We may hope that it damps the condition number of the preconditioned system $P^{-1/2}A_{\mu}P^{-1/2}$ nearly as well as the optimal preconditioner P_{\star} . We will support this intuition with a rigorous bound (Proposition 5.3).

5.2. Nyström PCG. We can obviously use the randomized Nyström preconditioner within the framework of PCG. We call this approach Nyström PCG, and we present a basic implementation in Algorithm 5.1.

More precisely, Algorithm 5.1 uses left-preconditioned CG. This algorithm implicitly works with the unsymmetric matrix $P^{-1}A_{\mu}$, rather than the symmetric matrix $P^{-1/2}A_{\mu}P^{-1/2}$. The two methods yield identical sequences of iterates [26], but the former is more efficient. For ease of analysis, our theoretical results are presented in terms of the symmetrically preconditioned matrix.

NYSTRÖM PCG

5.2.1. Complexity of Nyström PCG. Nyström PCG has two steps. First, we construct the randomized Nyström approximation, and then we solve the regularized linear system using PCG. We have already discussed the cost of constructing the Nyström approximation (subsection 2.2.1). The PCG stage costs $O(T_{\text{max}})$ operations per iteration, and it uses a total of O(n) additional storage.

For the regularized linear system (1.1), Theorem 5.1 and Corollary 5.2 demonstrate that it is enough to choose the sketch size $\ell = 2 \lceil 1.5d_{\text{eff}}(\mu) \rceil + 1$. In this case, the overall runtime of Nyström PCG is

 $O(d_{\text{eff}}(\mu)^2 n + T_{\text{mv}}(d_{\text{eff}}(\mu) + \log(1/\epsilon)))$ operations.

When the effective dimension $d_{\text{eff}}(\mu)$ is modest, Nyström PCG is very efficient.

In contrast, subsection 4.2 shows that the running time for Nyström sketch-andsolve has the same form—with $d_{\text{eff}}(\epsilon\mu)$ in place of $d_{\text{eff}}(\mu)$. This is a massive difference. Nyström PCG can produce solutions whose residual norm is close to machine precision; this type of successful computation is impossible with Nyström sketch-and-solve.

5.2.2. Comparison to other randomized preconditioning methods. In this subsection we give a more comprehensive discussion of how Nyström PCG compares to prior work on randomized preconditioning [3, 17, 20, 25] based on sketch-and-precondition and related ideas. All these prior methods were developed for least squares problems. We summarize the complexity of each method for regularized least-squares problems in Table 1.

Table 1: Regularized least-squares: complexity of prior randomized preconditoning methods vs. Nyström PCG. The table compares the complexity of Nyström PCG and state-of-the-art randomized preconditioning methods in the overdetermined case $n \ge d$, assuming we can access A only via matrix-vector products. The sketch-and-precondition preconditioner is constructed from a sketch SA, where $S \in \mathbb{R}^{m \times n}$ is a $(1 \pm \gamma)$ Gaussian subspace embedding with sketch size $\Omega(d/\gamma)$ and $\gamma \in (0, 1)$. The time to compute the sketch is $O(T_{mv}d/\gamma)$ and the iteration complexity follows from the argument in [37, Sec 2.6]. For AdaIHS we use a sketch constructed from a Gaussian subspace embedding with sketch size $O(d_{\text{eff}}(\mu)/\rho)$ where $\rho \in (0, 0.18)$. The complexity of AdaIHS follows from [17, Theorem 5]. The complexity of Nyström PCG is derived from Theorem 5.1 and Corollary 5.2.

Method	Complexity	References
Sketch-and-precondition	$O\left(T_{ m mv}d/\gamma+d^3/\gamma+T_{ m mv}rac{\log(2/\epsilon)}{\log(1/\gamma)} ight)$	[3, 20, 25]
AdaIHS	$O\left((T_{\rm mv}d_{\rm eff}/\rho + dd_{\rm eff}^2/\rho^2)\log(d_{\rm eff}/\rho) + T_{\rm mv}\frac{\log(1/\epsilon)}{\log(1/\rho)}\right)$	[17]
Nyström PCG	$O\left(T_{\rm mv}d_{\rm eff} + dd_{\rm eff}^2 + T_{\rm mv}\log(2/\epsilon)\right)$	This work

The time to construct the sketch-and-precondition preconditioner is always larger than the Nyström preconditioner, since $d_{\text{eff}} < d$ and $\gamma < 1$. Indeed, constructing the preconditioner for sketch-and-precondition costs $\Omega(d^3)$, which is the same as a direct method when $d = \Omega(n)$ and is prohibitive for high-dimensional problems. Hence Nyström PCG is amenable to problems with large d and runs much faster than sketchand-precondition whenever $d_{\text{eff}}(\mu) \ll d$. The Nyström preconditioner also enjoys wider applicability then sketch-precondition: it applies to square-ish systems, whereas the others only work for strongly overdetermined or underdetermined problems. In addition to sketch-and-precondition, Nyström PCG also enjoys better complexity than

AdaIHS. AdaIHS has linear dependence in d, but possesses additional logarithmic factors and scales in terms of $d_{\text{eff}}(\mu)/\rho$ where $\rho < 0.18$, leading to a worse runtime.

In the context of kernel ridge regression (KRR), the random features method of [2] may be viewed as a randomized preconditioning technique. [2] prove convergence guarantees for the polynomial kernel with a potentially prohibitive sketch size $\ell = O(d_{\text{eff}}(\mu)^2)$. In contrast, Nyström PCG can be used for KRR with any kernel and requires smaller sketch size $\ell = O(d_{\text{eff}}(\mu))$ to obtain fast convergence.

In summary, Nyström PCG applies to a wider class of problems than prior randomized preconditioners and enjoys stronger theoretical guarantees for regularized problems. Nyström PCG also outperforms other randomized preconditioners numerically (section 6).

5.2.3. Block Nyström PCG. We can also use the Nyström preconditioner with the block CG algorithm [21] to solve regularized linear systems with multiple right-hand sides. For this approach, we also use an orthogonalization pre-processing proposed in [11] that ensures numerical stability without further orthogonalization steps during the iteration.

5.3. Analysis of Nyström PCG. We now turn to the analysis of the randomized Nyström preconditioner P. Theorem 5.1 provides a bound for the rank ℓ of the Nyström preconditioner that reduces the condition number of A_{μ} to a constant. In this case, we deduce that Nyström PCG converges rapidly (Corollary 5.2).

THEOREM 5.1 (Nyström preconditioning). Suppose we construct the Nyström preconditioner P in (5.3) using Algorithm 2.1 with sketch size $\ell = 2 \lceil 1.5 d_{\text{eff}}(\mu) \rceil + 1$. Using P to precondition the regularized matrix A_{μ} results in the condition number bound

$$\mathbb{E}\big[\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})\big] < 28.$$

The proof of Theorem 5.1 may be found in subsection 5.3.3.

Theorem 5.1 has several appealing features. Many other authors have noticed that the effective dimension controls sample size requirements for particular applications such as discriminant analysis [7], ridge regression [17], and kernel ridge regression [1, 4]. In contrast, our result holds for any regularized linear system.

Our argument makes the role of the effective dimension conceptually simpler, and it leads to explicit, practical parameter recommendations. Indeed, the effective dimension $d_{\text{eff}}(\mu)$ is essentially the same as the sketch size ℓ that makes the approximation error $||A - \hat{A}_{nys}||$ proportional to μ . In previous arguments, such as those in [1, 4, 7], the effective dimension arises because the authors reduce the analysis to approximate matrix multiplication [8], which produces inscrutable constant factors.

Theorem 5.1 ensures that Nyström PCG converges quickly.

COROLLARY 5.2 (Nyström PCG: Convergence). Define P as in Theorem 5.1, and condition on the event $\{\kappa_2 (P^{-1/2}A_\mu P^{-1/2}) \leq 56\}$. Let $M = P^{-1/2}A_\mu P^{-1/2}$. If we initialize Algorithm 5.1 with initial iterate $x_0 = 0$, then the relative error δ_t in the iterate x_t satisfies

$$\delta_t = \frac{\|x_t - x_\star\|_M}{\|x_\star\|_M} < 2 \cdot (0.77)^t \quad where \ A_\mu x_\star = b.$$

In particular, after $t = \lceil 3.8 \log(2/\epsilon) \rceil$ iterations, we have relative error $\delta_t < \epsilon$.

The proof of Corollary 5.2 is an immediate consequence of the standard convergence result for CG [32, Theorem 38.5, p. 299]. See Appendix A.2.

NYSTRÖM PCG

5.3.1. Analyzing the condition number. The first step in the proof of Theorem 5.1 is a deterministic bound on how the preconditioner (5.3) reduces the condition number of the regularized matrix A_{μ} . Let us emphasize that this bound is valid for any rank- ℓ Nyström approximation, regardless of the choice of test matrix.

PROPOSITION 5.3 (Nyström preconditioner: deterministic bound). Let $\hat{A} =$ $U\hat{\Lambda}U^T$ be any rank- ℓ Nyström approximation, with ℓ th largest eigenvalue $\hat{\lambda}_{\ell}$, and let $E = A - \hat{A}$ be the approximation error. Construct the Nyström preconditioner P as in (5.3). Then the condition number of the preconditioned matrix $P^{-1/2}A_{\mu}P^{-1/2}$ satisfies

(5.4)
$$\max\left\{\frac{\hat{\lambda}_{\ell}+\mu}{\lambda_{n}+\mu},1\right\} \leq \kappa_{2}(P^{-1/2}A_{\mu}P^{-1/2})$$
$$\leq \left(\hat{\lambda}_{\ell}+\mu+\|E\|\right)\min\left\{\frac{1}{\mu}, \ \frac{\hat{\lambda}_{\ell}+\lambda_{n}+2\mu}{(\hat{\lambda}_{\ell}+\mu)(\lambda_{n}+\mu)}\right\}$$

For the proof of Proposition 5.3 see Appendix A.1.1.

To interpret the result, recall the expression (5.2) for the condition number induced by the optimal preconditioner. Proposition 5.3 shows that the Nyström preconditioner may reduce the condition number almost as well as the optimal preconditioner.

In particular, when $||E|| = O(\mu)$, the condition number of the preconditioned system is bounded by a constant, independent of the spectrum of A. In this case, Nyström PCG is guaranteed to converge quickly.

5.3.2. The effective dimension and sketch size selection. How should we choose the sketch size ℓ to guarantee that $||E|| = O(\mu)$? Corollary 2.3 shows how the error in the rank- ℓ randomized Nyström approximation depends on the spectrum of A through the eigenvalues of A and the tail stable rank. In this section, we present a lemma which demonstrates that the effective dimension $d_{\text{eff}}(\mu)$ controls both quantities. As a consequence of this bound, we will be able to choose the sketch size ℓ proportional to the effective dimension $d_{\text{eff}}(\mu)$.

Recall from (1.4) that the effective dimension of the matrix A is defined as

(5.5)
$$d_{\text{eff}}(\mu) = \text{tr}(A(A+\mu I)^{-1}) = \sum_{j=1}^{n} \frac{\lambda_j(A)}{\lambda_j(A) + \mu}.$$

As previously mentioned, $d_{\text{eff}}(\mu)$ may be viewed as a smoothed count of the eigenvalues larger than μ . Thus, one may expect that $\lambda_k(A) \leq \mu$ for $k \geq d_{\text{eff}}(\mu)$. This intuition is correct, and it forms the content of Lemma 5.4.

LEMMA 5.4 (Effective dimension). Let $A \in \mathbb{S}_n^+(\mathbb{R})$ with eigenvalues $\lambda_1 \geq \lambda_2 \geq$ $\dots \geq \lambda_n$. Let $\mu > 0$ be regularization parameter, and define the effective dimension as in (5.5). The following statements hold.

- 1. Fix $\gamma > 0$. If $j \ge (1 + \gamma^{-1})d_{\text{eff}}(\mu)$, then $\lambda_j \le \gamma \mu$. 2. If $k \ge d_{\text{eff}}(\mu)$, then $k^{-1} \sum_{j>k} \lambda_j \le (d_{\text{eff}}(\mu)/k) \cdot \mu$.

The proof of Lemma 5.4 may be found in Appendix A.1.2.

Lemma 5.4, Item 1 captures the intuitive fact that there are no more than $2d_{\text{eff}}(\mu)$ eigenvalues larger than μ . Similarly, Item 2 states that the effective dimension controls the sum of all the eigenvalues whose index exceeds the effective dimension. It is instructive to think about the meaning of these results when $d_{\text{eff}}(\mu)$ is small.

5.3.3. Proof of Theorem 5.1. We are now prepared to prove Theorem 5.1. The key ingredients in the proof are Proposition 2.2, Proposition 5.3, and Lemma 5.4.

Proof of Theorem 5.1. Fix the sketch size $\ell = 2 \lceil 1.5 d_{\text{eff}}(\mu) \rceil + 1$. Construct the rank- ℓ randomized Nyström approximation \hat{A}_{nys} with eigenvalues $\hat{\lambda}_j$. Write $E = A - \hat{A}_{\text{nys}}$ for the approximation error. Form the preconditioner P via (5.3). We must bound the expected condition number of the preconditioned matrix $P^{-1/2}A_{\mu}P^{-1/2}$

First, we apply Proposition 5.3 to obtain a deterministic bound that is valid for any rank- ℓ Nyström preconditioner:

$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le \frac{\hat{\lambda}_\ell + \mu + \|E\|}{\mu} \le 2 + \frac{\|E\|}{\mu}.$$

The second inequality holds because $\hat{\lambda}_{\ell} \leq \lambda_{\ell} \leq \mu$. This is a consequence of Lemma 2.1, Item 4 and Lemma 5.4, Item 1 with $\gamma = 1$. We rely on the fact that $\ell \geq 2 d_{\text{eff}}(\mu)$.

Decompose $\ell = 2p - 1$ where $p = \lceil 1.5 d_{\text{eff}}(\mu) \rceil + 1$. Take the expectation, and invoke Corollary 2.3 to obtain

$$\mathbb{E}\left[\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})\right] \le 2 + \left(3 + \frac{4\mathrm{e}^2}{p}\mathrm{sr}_p(A)\right)(\lambda_p/\mu).$$

By definition, $\operatorname{sr}_p(A) \cdot \lambda_p = \sum_{j \ge p} \lambda_j$. To complete the bound, apply Lemma 5.4 twice. We use Item 1 with $\gamma = 2$ and Item 2 with $k = p - 1 = \lfloor 1.5 d_{\text{eff}}(\mu) \rfloor$ to reach

$$\mathbb{E}\left[\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})\right] \le 2 + \frac{3 \cdot 2\mu + 4e^2 \cdot 2\mu/3}{\mu} < 2 + 26 = 28,$$

which is the desired result.

5.4. Practical parameter selection. In practice, we may not know the regularization parameter μ in advance, and we rarely know the effective dimension $d_{\text{eff}}(\mu)$. As a consequence, we cannot enact the theoretical recommendation for the rank of the Nyström preconditioner: $\ell = 2 \lceil 1.5 d_{\text{eff}}(\mu) \rceil + 1$. Instead, we need an adaptive method for choosing the rank ℓ . Below, we outline three strategies.

5.4.1. Strategy 1: Adaptive rank selection by a posteriori error estimation. The first strategy uses the posterior condition number estimate adaptively in a procedure the repeatedly doubles the sketch size ℓ as required. Recall that Proposition 5.3 controls the condition number of the preconditioned system:

(5.6)
$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le \frac{\hat{\lambda}_\ell + \mu + \|E\|}{\mu} \text{ where } E = A - \hat{A}_{nys}$$

We get $\hat{\lambda}_{\ell}$ for free from Algorithm 2.1 and we can compute the error ||E|| inexpensively with the randomized power method [16]; see Algorithm SM4.1 in section SM4. Thus, we can ensure the condition number is small by making ||E|| and $\hat{\lambda}_{\ell}$ fall below some desired tolerance. The adaptive strategy proceeds to do this as follows. We compute a randomized Nyström approximation with initial sketch size ℓ_0 , and we estimate the error ||E|| using randomized powering. If ||E|| is smaller than a prescribed tolerance, we accept the rank- ℓ_0 approximation. If the tolerance is not met, then we double the sketch size, update the approximation, and estimate ||E|| again. The process repeats until the estimate for ||E|| falls below the tolerance or it breaches a threshold ℓ_{max} for the maximum sketch size. Algorithm SM4.2 usings the following stopping criterions

 $||E|| \leq \tau \mu$ and $\hat{\lambda}_{\ell} \leq \tau \mu/11$ for a modest constant τ . The stopping criterion on $\hat{\lambda}_{\ell}$ does not seem to be necessary in practice, as it is usually an order of magnitude small than ||E||, but it is needed for Theorem 5.5. Based on numerical experience, we recommend a choice $\tau \in [1, 100]$. For full algorithmic details of adaptive rank selection by estimating ||E||, see SM4.2 in SM4.

The following theorem shows that with high probability, Algorithm SM4.2 terminates with a modest sketch size in at most a logarithmic number of steps, and PCG with the resulting preconditioner converges rapidly.

THEOREM 5.5. Run Algorithm SM4.2 with initial sketch size ℓ_0 and tolerance $\tau \mu$ where $\tau \geq 1$, and let $\tilde{\ell} = 2 \lceil 2d_{\text{eff}} \left(\frac{\delta \tau \mu}{11} \right) \rceil + 1$. Then with probability at least $1 - \delta$:

- 1. Algorithm SM4.2 doubles the sketch size at most $\lceil \log_2\left(\frac{\tilde{\ell}}{\ell_0}\right) \rceil$ times.
- 2. The final sketch size ℓ satisfies

$$\ell \leq 4 \lceil 2d_{\text{eff}} \left(\frac{\delta \tau \mu}{11}\right) \rceil + 2.$$

3. With the preconditioner constructed from Algorithm SM4.2, Nyström PCG converges in at most $\lceil \frac{\log(2/\epsilon)}{\log(1/\tau_0)} \rceil$ iterations, where $\tau_0 = \frac{\sqrt{1+12\tau/11}-1}{\sqrt{1+12\tau/11}+1}$.

Theorem 5.5 immediately implies the following concrete guarantee.

COROLLARY 5.6. Set $\tau = 44$ and $\delta = 1/4$ in Algorithm SM4.2 then with probability at least 3/4:

- 1. Algorithm SM4.2 doubles the sketch size at most $\lceil \log_2\left(\frac{\tilde{\ell}}{\ell_0}\right) \rceil$ times.
- 2. The final sketch size ℓ satisfies

$$\ell \le 4\lceil 2d_{\text{eff}}(\mu) \rceil + 2$$

3. With the preconditioner constructed from Algorithm SM4.2, Nyström PCG converges in at most $\lceil 3.48 \log(2/\epsilon) \rceil$ iterations.

5.4.2. Strategy 2: Adaptive rank selection by monitoring $\hat{\lambda}_{\ell}/\mu$. The second strategy is almost identical to the first, except we monitor the ratio $\hat{\lambda}_{\ell}/\mu$ instead of $||E||/\mu$. Strategy 2 doubles the approximation rank until $\hat{\lambda}_{\ell}/\mu$ falls below some tolerance (say, 10) or the sample size reaches the threshold ℓ_{max} . The approach is justified by the following proposition which shows that once the rank ℓ is sufficiently large, with high probability, the exact condition number differs from the empirical condition number $(\hat{\lambda}_{\ell} + \mu)/\mu$ by at most a constant.

PROPOSITION 5.7. Let $\tau \geq 0$ denote the tolerance and $\delta > 0$ a given failure probability. Suppose the rank of the randomized Nyström approximation satisfies $\ell \geq 2\lceil 2d_{\text{eff}}(\tau\mu) \rangle \rceil + 1$. Then

(5.7)
$$\mathbb{P}\left\{\left(\kappa_2(P^{-1/2}A_\mu P^{-1/2}) - \frac{\hat{\lambda}_\ell + \mu}{\mu}\right)_+ \le \frac{\tau}{\delta}\right\} \ge 1 - \delta,$$

where $X_{+} = \max\{X, 0\}.$

This strategy has the benefit of saving a bit of computation and is preferable when the target precision is not too important, eg, in machine learning problems where training error only loosely predicts test error.

Dataset	n	d
CIFAR-10	50,000	3,072
Guillermo	20,000	4,297
smallNorb-rf	24,300	10,000
shuttle-rf	43,300	10,000
Higgs-rf	800,000	10,000
YearMSD-rf	463,715	15,000

m 11 0	D 1	•	1
Table 2	Ridge	regression	datasets.
Table T.	rorage	rogrobbion	aacaboob

5.4.3. Strategy 3: Choose ℓ as large as the user can afford. The third strategy is to choose the rank ℓ as large as the user can afford. This approach is coarse, and it does not yield any guarantees on the cost of the Nyström PCG method.

Nevertheless, once we have constructed a rank- ℓ Nyström approximation we can combine the posterior estimate of the condition number used in strategy 1 with the standard convergence theory of PCG to obtain an upper bound for the iteration count of Nyström PCG. This gives us advance warning about how long it may take to solve the regularized linear system. As in strategy 1 we compute the error ||E|| in the condition number bound inexpensively with the randomized power method.

6. Applications and experiments. In this section, we study the performance of Nyström PCG on real world data from three different applications: ridge regression, kernel ridge regression, and approximate cross-validation. The experiments demonstrate the effectiveness of the preconditioner and our strategies for choosing the rank ℓ compared to other algorithms in the literature: on large datasets, we find that our method outperforms competitors by a factor of 5–10 (Table 3 and Table 10).

6.1. Preliminaries. We implemented all experiments in MATLAB R2019a and MATLAB R2021a on a server with 128 Intel Xeon E7-4850 v4 2.10GHz CPU cores and 1056 GB. Except for the very large scale datasets $(n \ge 10^5)$, every numerical experiment in this section was repeated twenty times; tables report the mean over the twenty runs, and the standard deviation (in parentheses) when it is non-zero. We highlight the best-performing method in a table in bold.

We select hyperparameters of competing methods by grid search to optimize performance. This procedure tends to be very charitable to the competitors, and it may not be representative of their real-world performance. Indeed, grid search is computationally expensive, and it cannot be used as part of a practical implementation. A detailed overview of the experimental setup for each application may be found in the appropriate section of section SM2, and additional numerical results in section SM3.

6.2. Ridge regression. In this section, we solve the ridge regression problem (1.7) described in subsection 1.3 on some standard machine learning data sets (Table 2) from OpenML [34] and LIBSVM [6]. We compare Nyström PCG to standard CG and two state-of-the-art randomized preconditioning methods, the sketch-and-precondition method of Rokhlin and Tygert (R&T) [25] and the Adaptive Iterative Hessian Sketch (AdaIHS) [17].

6.2.1. Experimental overview. We perform two sets of experiments: computing regularization paths on CIFAR-10 and Guillermo, and random features regression [23, 24] on shuttle, smallNORB, Higgs and YearMSD with specified values of μ . The values of μ may be found in subsection SM2.1. We use Euclidean norm $||r||_2$ of the residual as our stopping criteria, with convergence declared when $||r||_2 \leq 10^{-10}$. For

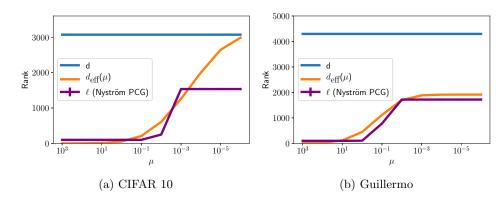


Fig. 2: Ridge regression: Adaptive sketch size selection. Nyström PCG with adaptive rank selection (Algorithm SM4.2) selects a preconditioner whose rank is less than or equal to the effective dimension. We report error bars for the rank selected by the adaptive algorithm, however the variation is so small that the error bars aren't visible. Hence despite being inherently random, the adaptive algorithm's behavior is practically deterministic across runs. See subsection 6.2.2.

both sets of experiments, we use Nyström PCG with adaptive rank selection, Algorithm SM4.2 in section SM4. For complete details of the experimental methodology, see subsection SM2.1.

The regularization path experiments solve (1.7) over a regularization path $\mu = 10^{j}$ where $j = 3, \dots, -6$. We begin by solving the problem for the largest μ (initialized at zero), and solve for progressively smaller μ with warm starting. For each value of μ , every method is allowed at most 500 iterations to reach the desired tolerance.

6.2.2. Computing the regularization path. Figure 2 shows the evolution of $d_{\text{eff}}(\mu)$ along the regularization path. CIFAR-10 and Guillermo are both small, so we compute the exact effective dimension as a point of reference. We see that we reach the sketch size cap of $\ell_{\text{max}} = 0.5d$ for CIFAR-10 and $\ell_{\text{max}} = 0.4d$ for Guillermo when μ is small enough. For CIFAR-10, Nyström PCG chooses a rank much smaller than the effective dimension for small values of μ . Nevertheless, the method still performs well (Figure 3).

Figure 3 show the effectiveness of each method for computing the entire regularization path. Nyström PCG is the fastest of the methods along most of the regularization path. For CIFAR-10, Nyström PCG is faster than R&T until the very end of the regularization path, when $d_{\text{eff}}(\mu) \approx d$. That is, the $O(d^3)$ cost of forming the R&T preconditioner is not worthwhile unless $d_{\text{eff}}(\mu) \approx d$. We expect Nyström PCG to perform better on ridge regression problems with non-vanishing regularization.

AdaIHS is rather slow. It increases the sketch size parameter several times along the regularization path. Each time, AdaIHS must form a new sketch of the matrix, approximate the Hessian, and compute a Cholesky factorization.

6.2.3. Random features regression. Tables 3 and 4 compare the performance of Nyström PCG, AdaIHS, and R&T PCG for random features regression. Table 3 shows that Nyström PCG performs best on all datasets for all metrics. The most striking feature is the difference between sketch sizes: AdaIHS and R&T require much

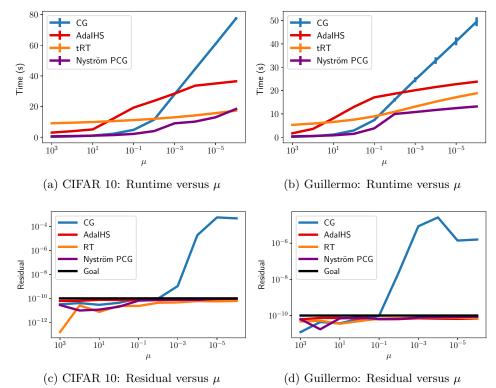


Fig. 3: Ridge regression: Runtime and residual. Nyström PCG is either the fastest method, or it is competitive with the fastest method for all values of the regularization parameter μ . CG is generally the slowest method. All the methods reliably achieve the target residual along the entire regularization path, except for ordinary CG at small values of μ . See subsection 6.2.2.

larger sketch sizes than Nyström PCG, leading to greater computation time and higher storage costs. Table 4 contains estimates for ||E|| and the condition number of the preconditioned system, which explain the fast convergence.

6.3. Approximate cross-validation. In this subsection we use our preconditioner to compute approximate leave-one-out cross-validation (ALOOCV), which requires solving a large linear system with multiple right-hand sides.

6.3.1. Background. Cross-validation is an important machine-learning technique to assess and select models and hyperparameters. Generally, it requires re-fitting a model on many subsets of the data, so can take quite a long time. The worst culprit is leave-one-out cross-validation (LOOCV), which requires running an expensive training algorithm n times. Recent work has developed approximate leave-one-out cross-validation (ALOOCV), a faster alternative that replaces model retraining by a linear system solve [12, 22, 36]. In particular, these techniques yield accurate and computationally tractable approximations to LOOCV.

To present the approach, we consider the infinitesimal jacknife (IJ) approximation

Dataset	Method	Final sketch size	Number of iterations	Total runtime (s)
	AdaIHS	10,000	66.9(0.933)	66.9(5.27)
shuttle-rf	R&T PCG	20,000	60.15	242.6(12.24)
	Nyström PCG	800	13.1(1.47)	9.78 (0.943)
	AdaIHS	12,800	38.7(1.42)	41.0(2.46)
smallNORB-rf	R&T PCG	20,000	34.5(1.31)	181.5(6.53)
	Nyström PCG	800	31.5(0.489)	6.67 (0.372)
	AdaIHS	30,000	44	1,327.3
YearMSD-rf	R&T PCG	30,000	49	766.5
	Nyström PCG	2,000	22	209.7
	AdaIHS	6,400	55	1,052.7
Higgs-rf	R&T PCG	20,000	53	607.4
	Nyström PCG	800	28	91.26

Table 3: **Ridge regression: Nyström PCG versus AdaIHS and R&T PCG.** Nyström PCG outperforms AdaIHS and R&T PCG in iteration and runtime complexity for both datasets. Additionally, Nyström PCG requires much less storage.

Table 4: **Ridge regression: Quality of Nyström preconditioner.** We use Nyström PCG as outlined in subsection 5.4.3. The first column gives an estimate for the error ||E|| in the randomized Nyström approximation. The second column gives an estimate for the condition number $\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})$.

Dataset	Estimate of $ E $	Estimated condition number of preconditioned system
shuttle-rf	$7.29e{-13}$	4.17(0.161)
smallNorb-rf	9.90e - 3	18.5(0.753)
YearMSD-rf	$2.07e{-4}$	22.7
Higgs-rf	2.21e - 3	23.8

to LOOCV [12, 30]. The IJ approximation computes

(6.1)
$$\tilde{\theta}_{\mathrm{IJ}}^{n/j} = \hat{\theta} + \frac{1}{n} H^{-1}(\hat{\theta}) \nabla_{\theta} l(\hat{\theta}, a_j),$$

where $H(\hat{\theta}) \in \mathbb{R}^{d \times d}$ is the Hessian of the loss function at the solution $\hat{\theta}$, for each datapoint a_j . The main computational challenge is computing the inverse Hessian vector product $H^{-1}(\hat{\theta})\nabla_{\theta}l(\hat{\theta}, a_j)$. When n is very large, we can also subsample the data and average (6.1) over the subsample to estimate ALOOCV. Since ALOOCV solves the same problem with several right-hand sides, blocked PCG methods (here, Nyström blocked PCG) are the tool of choice to efficiently solve for multiple right-hand sides at once. To demonstrate the idea, we perform numerical experiments on ALOOCV for logistic regression. The datasets we use are all from LIBSVM [6]; see Table 5.

6.3.2. Experimental overview. We perform two sets of experiments in this section. The first set of experiments uses Gisette and SVHN to test the efficacy of Nyström sketch-and-solve. These datasets are small enough that we can factor $H(\theta)$ using a direct method. We also compare to block CG and block PCG with the computed Nyström approximation as a preconditioner. To assess the error due to an inexact solve for datapoint a_j , let $x_*(a_j) = H^{-1}(\theta) \nabla_{\theta} l(\hat{\theta}, a_j)$. For any putative

Dataset	n	d	%nz(A)	μ	Initial sketch size
Gisette	6,000	5,000	99.1%	1, 1e-4	850
real-sim	72,308	20,958	0.245%	1e-4, 1e-8	500
rcv1.binary	20,242	47,236	0.157%	1e-4, 1e-8	500
SVHN	73,257	3,072	100%	1, 1e-4	850

Table 5: ALOOCV datasets and experimental parameters.

Table 6: ALOOCV: Small datasets. The error for a given value of μ is the maximum relative error on 100 randomly sampled datapoints, averaged over 20 trials.

Datase	et μ	Nyström sketch-and-solve	Block CG	Block Nyström PCG
Gisette	e 1	4.99e-2	$2.68e{-11}$	$2.58e{-}12$
Gisette	e 1e-4	1.22	$1.19e{-11}$	$6.59e{-}12$
SVHN	1	9.12e-5	2.80e - 13	1.26e - 13
SVHN	1e-4	3.42e-1	$2.01e{-10}$	1.41e-11

solution $\hat{x}(a_j)$, we compute the relative error $\|\hat{x}(a_j) - x_{\star}(a_j)\|_2 / \|x_{\star}(a_j)\|_2$. We average the relative error over 100 data-points a_j .

The second set of experiments uses the larger datasets real-sim and rcv1.binary and small values of μ , the most challenging setting for ALOOCV. We restrict our comparison to block Nyström PCG versus the block CG algorithm, as Nyström sketchand-solve is so inaccurate in this regime. We employ Algorithm SM4.2 to construct the preconditioner for block Nyström PCG.

6.3.3. Nyström sketch-and-solve. As predicted, Nyström sketch-and-solve works poorly (Table 6). When $\mu = 1$, the approximate solutions are modestly accurate, and the accuracy degrades as μ decreases to 10^{-4} . The experimental results agree with the theoretical analysis presented in section 4, which indicate that sketch-and-solve degrades as μ decreases. In contrast, block CG and block Nyström PCG both provide high-quality solutions for each datapoint for both values of the regularization parameter.

6.4. Large scale ALOOCV experiments. Table 7 summarizes results for block Nyström PCG and block CG on the larger datasets. When $\mu = 10^{-4}$, block Nyström PCG offers little or no benefit over block CG because the data matrices are very sparse (see Table 5) and the rcv1 problem is well-conditioned (see Table SM2).

For $\mu = 10^{-8}$, block Nyström PCG reduces the number of iterations substantially, but the speedup is negligible. The data matrix A is sparse, which reduces the benefit of the Nyström method. Block CG also benefits from the presence of multiple righthand sides just as block Nyström PCG. Indeed, O'Leary proved that the convergence of block CG depends on the ratio $(\lambda_s + \mu)/(\lambda_n + \mu)$, where s is is the number of right-hand sides [21]. Consequently, multiple right-hand sides precondition block CG and accelerate convergence. We expect bigger gains over block CG when A is dense.

6.5. Kernel ridge regression. Our last application is kernel ridge regression (KRR), a supervised learning technique that uses a kernel to model nonlinearity in the data. KRR leads to large dense linear systems that are challenging to solve.

Dataset	μ	Method	Number of iterations	Runtime (s)
rcv1	1e-4	Block CG	12	$11.06\ (0.874)$
1011	1e-4	Block Nyström PCG	10	11.87(0.767)
rcv1	1e-8	Block CG	52	39.03(2.97)
1011	1e-8	Block Nyström PCG	15	24.1(1.79)
realsim	1e-4	Block CG	12	23.04(2.04)
Teaisiin	1e-4	Block Nyström PCG	8	19.05(1.10)
realsim	1e-8	Block CG	90	163.7(12.3)
Tealshii	1e-8	Block Nyström PCG	32	68.9 (5.30)

Table 7: **ALOOCV: Large datasets**. Block Nyström PCG outperforms block CG as μ becomes small.

6.5.1. Background. We briefly review KRR [28]. Given a dataset of inputs $x_i \in \mathcal{D}$ and their corresponding outputs $b_i \in \mathbb{R}$ for $i = 1, \ldots, n$ and a kernel function $\mathcal{K}(x, y)$, KRR finds a function $f_\star : \mathcal{D} \to \mathbb{R}$ in the associated reproducing kernel Hilbert space \mathcal{H} that best predicts the outputs for the given inputs. The solution $f_\star \in \mathcal{H}$ minimizes the square error subject to a complexity penalty:

(6.2)
$$f_{\star} = \underset{f \in \mathcal{H}}{\operatorname{argmin}} \quad \frac{1}{2n} \sum_{i=1}^{n} (f(x_i) - b_i)^2 + \frac{\mu}{2} \|f\|_{\mathcal{H}}^2,$$

where $\|\cdot\|_{\mathcal{H}}$ denotes the norm on \mathcal{H} . Define the kernel matrix $K \in \mathbb{R}^{n \times n}$ with entries $K_{ij} = \mathcal{K}(x_i, x_j)$. The representer theorem [29] states the solution to (6.2) is

$$f_{\star}(x) = \sum_{i=1}^{n} \alpha_i \mathcal{K}(x, x_i)$$

where $\alpha = (\alpha_1, \ldots, \alpha_n)$ solves the linear system

$$(6.3) (K+n\mu I)\alpha = b$$

Solving the linear system (6.3) is the computational bottleneck of KRR. Direct factorization methods to solve (6.3) are prohibitive for large n as their costs grow as n^3 ; for $n > 10^4$ or so, iterative methods are generally preferred. However, K is often extremely ill-conditioned, even with the regularization term $n\mu I$. As a result, CG for Problem (6.3) converges slowly.

6.5.2. Experimental overview. We use Nyström PCG to solve several KRR problems derived from classification problems on real world datasets from [6, 34]. For all experiments, we use the Gaussian kernel $\mathcal{K}(x,y) = \exp(-||x-y||^2/(2\sigma^2))$. We compare our method to random features PCG, proposed in [2]. We do not compare to vanilla CG as it is much slower than Nyström PCG and random features PCG.

All datasets either come with specified test sets, or we create one from a random 80-20 split. The PCG tolerance, σ , and μ were all chosen to achieve good performance on the test sets (see Table 11 below). Both methods were allowed to run for a maximum of 500 iterations. The statistics for each dataset and the experimental parameters are given in Table 8.

We run two sets of experiments. For the datasets with $n < 10^5$ we run oracle random features PCG against two versions of the Nyström PCG algorithm. The first

Dataset	n	d	n _{classes}	μ	σ	PCG tolerance
ijcnn1	49,990	49	2	1e-6	0.5	1e-3
MNIST	60,000	784	10	1e-7	5	1e-4
Sensorless	48,509	48	11	1e-8	0.8	1e-4
SensIT	78,823	100	3	1e-8	3	1e-3
MiniBooNE	104,052	50	2	1e-7	5	1e-4
EMNIST-Balanced	105,280	784	47	1e-6	8	1e-3
Santander	160,000	200	2	1e-6	7	1e-3

Table 8: Kernel ridge regression datasets and experimental parameters.

Table 9: Kernel ridge regression: Iteration count and runtime. The adaptive and oracle Nyström PCG algorithms outperform oracle random features PCG in both time and iteration complexity.

Dataset	Method	Number of iterations	Total runtime (s)
	Oracle random features PCG	63.8(2.66)	38.3(2.33)
icjnn1	Adaptive Nyström PCG	43.7(1.77)	32.0(1.47)
	Oracle Nyström PCG	31.8(0.835)	33.3(1.60)
	Oracle random features PCG	314.5(2.88)	254.7(6.93)
MNIST	Adaptive Nyström PCG	78.5(17.65)	148.1(46.39)
	Oracle Nyström PCG	77.9(2.08)	91.7(2.08)
	Oracle random features PCG	55.4(2.35)	39.9(3.96)
Sensorless	Adaptive Nyström PCG	22.0(0.510)	24.3(1.26)
	Oracle Nyström PCG	21.7 (0.571)	22.7(1.63)
	Oracle random features PCG	68.0(4.31)	95.2(6.19)
SensIT	Adaptive Nyström PCG	47.8(1.72)	70.1(2.43)
	Oracle Nyström PCG	48.7(3.40)	61.6(6.41)

version uses the oracle best value of ℓ found by grid search (from the same grid used to select $m_{\rm rf}$) to minimize the total runtime, and the second is the adaptive algorithm described in subsection 5.4.2. The grid for ℓ and $m_{\rm rf}$ is restriced to less than 10,000 to keep the preconditioners cheap to apply and store. The adaptive algorithm for each dataset was initialized at $\ell = 2,000$, which is smaller than 0.05*n* for all datasets. For the datasets with $n \ge 10^5$, we restricted both ℓ and $m_{\rm rf}$ to 1,000, which corresponds to less than 0.01*n*. We then run both algorithms till they reach the desired tolerance or the maximum number of iterations are reached.

We use column sampling to construct the Nyström preconditioner for all KRR problems: on these problems, random projections take longer and yield similar performance (with somewhat lower variance).

6.5.3. Experimental results. Tables 9 to 11 summarize the results for the KRR experiments. Table 9 shows that both versions of Nyström PCG deliver better performance than random features preconditioning on all the datasets considered. Nyström PCG also uses less storage. Table 10 shows that Nyström PCG yields better performance than random features PCG on the larger scale datasets when both are restricted to ranks of 1,000. Table 11 shows the adaptive strategy proposed in subsection 5.4.2 to select ℓ works very well. In contrast, it is difficult to choose $m_{\rm rf}$ for random features preconditioning: the authors of [2] provide no guidance except for the polynomial kernel.

Table 10: Kernel ridge regression regression: Nyström PCG fixed rank versus random features PCG fixed rank. Nyström PCG again outperforms random features PCG in iteration and time complexity.

Dataset	Method	Number of iterations	Total runtime (s)
MiniBooNE	Random Features PCG	92	154.2
	Nyström PCG	72	137.4
EMNIST-Balanced	Random Features PCG	154	635.2
	Nyström PCG	32	268.4
Santander	Random Features PCG	160	810.4
	Nyström PCG	31	164.8

Table 11: Kernel ridge regression: Oracle parameters and test errors. The final ranks selected by the adaptive algorithm are almost in full agreement with the oracle ranks. Furthermore, the adaptive and oracle ranks for Nyström PCG are never larger than $m_{\rm rf}$.

Dataset	Adaptive Nyström final rank	Oracle Nyström rank	$\mathbf{Oracle} \ \mathbf{m}_{\mathrm{rf}}$	Test set error
ijcnn1	2,000	3,000	3,000	1.25%
MNIST	6,000(1,716)	4,000	9,000	1.22%
Sensorless	2,000	2,000	5,000	2.01%
SensIT	2,000	2,000	7,000	12.83%
MiniBooNE	NA	NA	NA	7.93%
EMNIST-Balanced	NA	NA	NA	15%
Santander	NA	NA	NA	8.90%

7. Conclusion. We have shown that Nyström PCG delivers a strong benefit over standard CG both in the theory and in practice, thanks to the ease of parameter selection, on a range of interesting large-scale computational problems including ridge regression, kernel ridge regression, and ALOOCV. In our experience, Nyström PCG outperforms all generic methods for solving large-scale dense linear systems with spectral decay. It is our hope that this paper motivates further research on randomized preconditioning for solving large scale linear systems and offers a useful speedup to practitioners.

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FRANGELLA, TROPP, AND UDELL

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NYSTRÖM PCG

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Appendix A. Proofs of main results. This appendix contains full proofs of the main results that are substantially novel (Theorem 4.2, Proposition 5.3, , and Corollary 5.2). The supplement contains proofs of results that are similar to existing work, but do not appear explicitly in the literature.

A.1. Proof Theorem 4.2. This result contains the analysis of the Nyström sketch-and-solve method. We begin with (4.2), which provides an error bound that compares the regularized inverse of a psd matrix A with the regularized inverse of the randomized Nyström approximation \hat{A}_{nys} . Since $0 \leq \hat{A}_{nys} \leq A$, we can apply Proposition 3.1 to obtain a deterministic bound for the discrepancy:

$$\|(\hat{A}_{nys} + \mu I)^{-1} - (A + \mu I)^{-1}\| \le \frac{1}{\mu} \frac{\|E\|}{\|E\| + \mu}$$
 where $E = A - \hat{A}_{nys}$.

The function $f(t) = t/(t + \mu)$ is concave, so we can take expectations and invoke Jensen's inequality to obtain

$$\mathbb{E}\|(\hat{A}_{nys} + \mu I)^{-1} - (A + \mu I)^{-1}\| \le \frac{1}{\mu} \frac{\mathbb{E}\|E\|}{\mathbb{E}\|E\| + \mu}$$

Inserting the bound (2.3) on $\mathbb{E}||E||$ from Corollary 2.3 gives

$$\mathbb{E}\|(\hat{A}_{nys} + \mu I)^{-1} - (A + \mu I)^{-1}\| \le \frac{1}{\mu} \cdot \frac{(3 + 4e^{2}sr_{p}(A)/p)\lambda_{p}}{\mu + (3 + 4e^{2}sr_{p}(A)/p)\lambda_{p}}$$

To conclude, observe that the denominator of the second fraction exceeds $\mu + \lambda_p$.

Now, let us establish (4.3), the error bound for Nyström sketch-and-solve. Introduce the solution \hat{x} to the Nyström sketch-and-solve problem and the solution x_{\star} to the regularized linear system:

$$(\hat{A}_{nys} + \mu I)\hat{x} = b$$
 and $(A + \mu I)x_{\star} = b$.

We may decompose the regularized matrix as $A + \mu I = \hat{A}_{nys} + \mu I + E$. Subtract the two equations in the last display to obtain

$$(\hat{A}_{nys} + \mu I)(\hat{x} - x_{\star}) - Ex_{\star} = 0.$$

Rearranging to isolate the error in the solution, we have

$$\hat{x} - x_\star = (\hat{A}_{nys} + \mu I)^{-1} E x_\star$$

Take the norm, apply the operator norm inequality, and use the elementary bound $\|(\hat{A}_{nys} + \mu I)^{-1}\| \leq \mu^{-1}$. We obtain

$$\frac{\|\hat{x} - x_\star\|_2}{\|x_\star\|_2} \le \frac{\|E\|}{\mu}.$$

Finally, take the expectation and repeat the argument used to control $\mathbb{E}||E||/\mu$ in the proof of Theorem 5.1.

A.1.1. Proof of Proposition 5.3. Let $\hat{A} = U\hat{\Lambda}U^T$ be an arbitrary rank- ℓ Nyström approximation whose ℓ th eigenvalue is $\hat{\lambda}_{\ell}$. Proposition 5.3 provides a deterministic bound on the condition number of the regularized matrix A_{μ} after preconditioning with

$$P = \frac{1}{\hat{\lambda}_{\ell} + \mu} U(\hat{\Lambda} + \mu I)U^T + (I - UU^T)$$

We remind the reader that this argument is completely deterministic.

First, note that the preconditioned matrix $P^{-1/2}A_{\mu}P^{-1/2}$ is psd, so

$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) = \frac{\lambda_1(P^{-1/2}A_\mu P^{-1/2})}{\lambda_n(P^{-1/2}A_\mu P^{-1/2})}$$

Let us begin with the upper bound on the condition number. We have the decomposition

(A.1)
$$P^{-1/2}A_{\mu}P^{-1/2} = P^{-1/2}(\hat{A} + \mu I)P^{-1/2} + P^{-1/2}EP^{-1/2},$$

owing to the relation $A_{\mu} = \hat{A} + \mu I + E$. Recall that the error matrix E is psd, so the matrix $P^{-1/2}EP^{-1/2}$ is also psd.

First, we bound the maximum eigenvalue. Weyl's inequalities imply that

$$\lambda_1(P^{-1/2}A_{\mu}P^{-1/2}) \le \lambda_1(P^{-1/2}(\hat{A} + \mu I)P^{-1/2}) + \lambda_1(P^{-1/2}EP^{-1/2}).$$

A short calculation shows that $\lambda_1(P^{-1/2}(\hat{A} + \mu I)P^{-1/2}) = \hat{\lambda}_\ell + \mu$. When $\ell < n$, we have $\lambda_1(P^{-1}) = 1$. Therefore,

$$\lambda_1(P^{-1/2}EP^{-1/2}) = \lambda_1(P^{-1}E) \le \lambda_1(P^{-1})\lambda_1(E) = \lambda_1(E) = ||E||.$$

In summary,

(A.2)
$$\lambda_1(P^{-1/2}A_\mu P^{-1/2}) \le \hat{\lambda}_\ell + \mu + ||E||$$

For the minimum eigenvalue, we first assume that $\mu > 0$. Apply Weyl's inequality to (A.1) to obtain to obtain

(A.3)
$$\lambda_n(P^{-1/2}A_\mu P^{-1/2}) \ge \lambda_n(P^{-1/2}(\hat{A} + \mu I)P^{-1/2}) + \lambda_n(P^{-1/2}EP^{-1/2}) \\ \ge \lambda_n(P^{-1/2}(\hat{A} + \mu I)P^{-1/2}) = \mu.$$

Combining (A.2) and (A.3), we reach

$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le \frac{\hat{\lambda}_\ell + \mu + \|E\|}{\mu}.$$

This gives a bound for the maximum in case $\mu > 0$.

If we only have $\mu \geq 0$, then a different argument is required for the smallest eigenvalue. Assume that A is positive definite, in which case $\hat{\lambda}_{\ell} > 0$. As $P^{-1/2}A_{\mu}P^{-1/2}$ is symmetric positive definite we have

$$\lambda_n(P^{-1/2}A_\mu P^{-1/2}) = \frac{1}{\lambda_1(P^{1/2}A_\mu^{-1}P^{1/2})}.$$

Conjugating by $A_{\mu}^{1/2}P^{-1/2}$ and using similarity, we obtain the equality

$$\lambda_1(P^{1/2}A_{\mu}^{-1}P^{1/2}) = \lambda_1(A_{\mu}^{-1/2}PA_{\mu}^{-1/2}).$$

Hence it suffices to produce an upper bound for $\lambda_1(A_{\mu}^{-1/2}PA_{\mu}^{-1/2})$. To that end, we expand

$$\begin{split} \lambda_1(A_{\mu}^{-1/2}PA_{\mu}^{-1/2}) &= \lambda_1 \left(A_{\mu}^{-1/2} \left(\frac{1}{\hat{\lambda}_{\ell} + \mu} (\hat{A} + \mu U U^T) + (I - U U)^T \right) A_{\mu}^{-1/2} \right) \\ &\leq \frac{1}{\hat{\lambda}_{\ell} + \mu} \lambda_1 \left(A_{\mu}^{-1/2} (\hat{A} + \mu U U^T) A_{\mu}^{-1/2} \right) \\ &+ \lambda_1 \left(A_{\mu}^{-1/2} \left(I - U U^T \right) A_{\mu}^{-1/2} \right). \end{split}$$

The second inequality is Weyl's. Since $\hat{A} \leq A$, we have $\hat{A} + \mu U U^T \leq A_{\mu}$. The last display simplifies to

$$\lambda_1(A_{\mu}^{-1/2}PA_{\mu}^{-1/2}) \leq \frac{1}{\hat{\lambda}_{\ell} + \mu} + \frac{1}{\lambda_n + \mu}.$$

Putting the pieces together with (A.2), we obtain

$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le (\hat{\lambda}_\ell + \mu + ||E||) \left(\frac{1}{\hat{\lambda}_\ell + \mu} + \frac{1}{\lambda_n + \mu}\right).$$

Thus,

$$\kappa_2(P^{-1/2}A_{\mu}P^{-1/2}) \le \left(\hat{\lambda}_{\ell} + \mu + \|E\|\right) \min\left\{\frac{1}{\mu}, \ \frac{\hat{\lambda}_{\ell} + \lambda_n + 2\mu}{(\hat{\lambda}_{\ell} + \mu)(\lambda_n + \mu)}\right\}.$$

This formula is valid when A is positive definite or when $\mu > 0$.

We now prove the lower bound on $\kappa_2(P^{-1/2}A_{\mu}P^{-1/2})$. Returning to (A.1) and invoking Weyl's inequalities yields

$$\lambda_1(P^{-1/2}A_{\mu}P^{-1/2}) \ge \lambda_1(P^{-1/2}(\hat{A} + \mu I)P^{-1/2}) + \lambda_n(P^{-1/2}EP^{-1/2}) \ge \hat{\lambda}_{\ell} + \mu.$$

For the smallest eigenvalue we observe that

$$\lambda_n(P^{-1/2}A_{\mu}P^{-1/2}) = \lambda_n(A_{\mu}P^{-1}) \le \lambda_n(A_{\mu})\lambda_1(P^{-1}) = \lambda_n + \mu.$$

Combining the last two displays, we obtain

$$\frac{\hat{\lambda}_{\ell} + \mu}{\lambda_n + \mu} \le \kappa_2 (P^{-1/2} A_{\mu} P^{-1/2}).$$

Condition numbers always exceed one, so

$$\max\left\{\frac{\hat{\lambda}_{\ell}+\mu}{\lambda_n+\mu},1\right\} \le \kappa_2(P^{-1/2}A_{\mu}P^{-1/2}).$$

This point concludes the argument.

A.1.2. Proof of Lemma 5.4. Lemma 5.4 establishes the central facts about the effective dimension. First, we prove Item 1. Fix a parameter $\gamma \geq 1$, and set $j_{\star} = \max\{1 \leq j \leq n : \lambda_j > \gamma\mu\}$. We can bound the effective dimension below by the following mechanism.

$$d_{\text{eff}}(\mu) = \sum_{j=1}^{n} \frac{\lambda_j}{\lambda_j + \mu} \ge \sum_{j=1}^{j_\star} \frac{\lambda_j}{\lambda_j + \mu} \ge j_\star \cdot \frac{\lambda_{j_\star}}{\lambda_{j_\star} + \mu}.$$

We have used the fact that $t \mapsto t/(1+t)$ is increasing for $t \ge 0$, Solving for j_{\star} , we determine that

$$j_{\star} \leq (1 + \mu/\lambda_{j_{\star}}) d_{\text{eff}}(\mu) < (1 + \gamma^{-1}) d_{\text{eff}}(\mu)$$

The last inequality depends on the definition of j_{\star} . This is the required result.

Item 2 follows from a short calculation:

$$\frac{1}{k}\sum_{j>k}\lambda_j = \frac{\lambda_k + \mu}{k}\sum_{j>k}\frac{\lambda_j}{\lambda_k + \mu} \le \frac{\lambda_k + \mu}{k}\sum_{j>k}\frac{\lambda_j}{\lambda_j + \mu}$$
$$= \frac{\lambda_k + \mu}{k}\left(d_{\text{eff}}(\mu) - \sum_{j=1}^k\frac{\lambda_j}{\lambda_j + \mu}\right) \le \frac{\lambda_k + \mu}{k}\left(d_{\text{eff}}(\mu) - \frac{k\lambda_k}{\lambda_k + \mu}\right)$$
$$= \frac{\mu d_{\text{eff}}(\mu)}{k} + \lambda_k\left(\frac{d_{\text{eff}}(\mu)}{k} - 1\right) \le \frac{\mu d_{\text{eff}}(\mu)}{k}.$$

The last inequality depends on the assumption that $k \ge d_{\text{eff}}(\mu)$.

A.2. Proof of Corollary 5.2. This result gives a bound for the relative error δ_t in the iterates of PCG. Recall the standard convergence bound for CG [32, Theorem 38.5]:

$$\delta_t \le 2 \left(\frac{\sqrt{\kappa_2 (P^{-1/2} A_\mu P^{-1/2})} - 1}{\sqrt{\kappa_2 (P^{-1/2} A_\mu P^{-1/2})} + 1} \right)^t.$$

We conditioned on the event that $\{\kappa(P^{-1/2}A_{\mu}P^{-1/2}) \leq 56\}$. On this event, the relative error must satisfy

$$\delta_t < 2 \left(\frac{\sqrt{56} - 1}{\sqrt{56} + 1}\right)^t \le 2 \cdot (0.77)^t.$$

Solving for t, we see that $\delta_t < \epsilon$ when $t \ge \lfloor 3.8 \log(1/\epsilon) \rfloor$. This concludes the proof.

A.3. Proof of Theorem 5.5. Theorem 5.5 establishes that with high probability Algorithm SM4.2 terminates in a logarithmic number of steps, the sketch size remains $O(d_{\text{eff}}(\delta \tau \mu))$, and PCG with the preconditioner constructed from the output converges fast.

Proof. We first recall that Algorithm SM4.2 terminates when the event

$$\mathcal{E} = \{ \|E\| \le \tau\mu \} \cap \{ \hat{\lambda}_{\ell} \le \frac{\tau\mu}{11} \}$$

holds. Observe that conditioned on \mathcal{E} , Proposition 5.3 yields

$$\kappa_2(P^{-1/2}A_\mu P^{-1/2}) \le \frac{\hat{\lambda}_\ell + \mu + \|E\|}{\mu} \le 1 + (1 + \frac{1}{11})\tau = 1 + \frac{12}{11}\tau.$$

Statement 3 now follows from the above display and the standard convergence theorem for CG.

Now, if Algorithm SM4.2 terminates with $N \leq \lceil \log_2 \left(\tilde{\ell}/\ell_0 \right) \rceil - 1$ steps of sketch size doubling, then \mathcal{E} holds with probability 1. Statement 3 then follows by our initial observation, while statements 1 and 2 hold trivially. Hence statements 1-3 all hold if the algorithm terminates in $N \leq \lceil \log_2 \left(\tilde{\ell}/\ell_0 \right) \rceil - 1$ steps.

Thus to conclude the proof, it suffices to show that if $N \geq \lceil \log_2 \left(\tilde{\ell}/\ell_0 \right) \rceil$, then \mathcal{E} holds with probability at least $1 - \delta$, which implies that statements 1-3 hold with probability at least $1 - \delta$, as above.

We now show that \mathcal{E} holds with probability at least $1-\delta$ when $N = \lceil \log_2(\tilde{\ell}/\ell_0) \rceil$. To see this note that when $N = \lceil \log_2(\tilde{\ell}/\ell_0) \rceil$, we have $\ell \geq \tilde{\ell}$. Consequently, we may invoke Proposition 2.2 with $p = \lceil 2d_{\text{eff}}(\frac{\delta\tau\mu}{11}) \rceil + 1$ and Lemma 5.4 to show

$$\mathbb{E}[\|E\|] \stackrel{(1)}{\leq} 3\lambda_p + \frac{2e^2}{p} \left(\sum_{j=p}^n \lambda_j\right)$$
$$\stackrel{(2)}{\leq} 3\frac{\delta\tau\mu}{11} + 2e^2 \frac{d_{\text{eff}}(\delta\tau\mu/11)}{p} \frac{\delta\tau\mu}{11}$$
$$\stackrel{(3)}{\leq} \frac{3}{11}\delta\tau\mu + \frac{2e^2}{2} \frac{\delta\tau\mu}{11} = \left(\frac{3+e^2}{11}\right)\delta\tau\mu \leq \delta\tau\mu.$$

Where step (1) uses Proposition 2.2, step (2) uses items 1 and 2 of Lemma 5.4 with $\gamma = 1$, and step (3) follows from $p \geq 2d_{\text{eff}}(\delta \tau \mu)$. Thus,

$$\mathbb{E}[\|E\|] \le \delta \tau \mu.$$

By Markov's inequality,

$$\mathbb{P}\{\|E\| > \tau\mu\} \le \delta$$

Hence $\{||E|| \leq \tau \mu\}$ holds with probability at least $1 - \delta$. Furthermore, by Lemma 5.4 we have $\{\hat{\lambda}_{\ell} \leq \delta \tau \mu/11\}$ with probability 1 as $\hat{\lambda}_{\ell} \leq \lambda_{\ell} \leq \lambda_{p}$. Thus when $N = \lceil \log_2(\tilde{\ell}/\ell_0) \rceil$, \mathcal{E} holds with probability at least $1 - \delta$, this immediately implies statements 1 and 3. Statement 2 follows as

$$\ell = 2^N \ell_0 \le 2^{\log_2(\tilde{\ell}/\ell_0)+1} \ell_0 = 2\tilde{\ell} = 4\lceil 2d_{\text{eff}}\left(\frac{\delta\tau\mu}{11}\right)\rceil + 2,$$

where in the first inequality we used $[x] \leq x + 1$, this completes the proof.

A.4. Proof of Proposition 5.7. Proposition 5.7 shows once $\ell = \Omega(d_{\text{eff}}(\tau\mu))$, then with high probability $\kappa_2(P^{-1/2}A_\mu P^{-1/2})$ differs from $(\hat{\lambda}_\ell + \mu)$ by at most a constant.

Proof. Proposition 5.3 implies that

$$\left(\kappa_2(P^{-1/2}A_{\mu}P^{-1/2}) - \frac{\hat{\lambda}_{\ell} + \mu}{\mu}\right)_+ \le \frac{\|E\|}{\mu}.$$

Combining the previous display with Markov's inequality yields

$$\mathbb{P}\left\{\left(\kappa_2(P^{-1/2}A_{\mu}P^{-1/2}) - \frac{\hat{\lambda}_{\ell} + \mu}{\mu}\right)_+ > \frac{\tau}{\delta}\right\} \le \frac{\delta}{\tau} \frac{\mathbb{E}[\|E\|]}{\mu}.$$

Now, our choice of ℓ combined with Proposition 2.2 and Lemma 5.4 implies that $\mathbb{E}[||E||] \leq \tau \mu$. Hence we have

$$\mathbb{P}\left\{\left(\kappa_2(P^{-1/2}A_{\mu}P^{-1/2}) - \frac{\hat{\lambda}_{\ell} + \mu}{\mu}\right)_+ > \frac{\tau}{\delta}\right\} \le \delta,$$

FRANGELLA, TROPP, AND UDELL

which implies the desired claim.