

# Random Projections and Dimension Reduction

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- Kernel PCA
- Kernel SVM

# Johnson-Lindenstrauss Lemma

If we have  $n$  data points in  $\mathbb{R}^d$ , there exists a linear map into  $\mathbb{R}^k$  such that pairwise distances between data points can be preserved up to an  $\epsilon$  tolerance, provided  $k > C\epsilon^{-2} \log n$ , where  $C$  is some constant. The proof follows three steps:

- Define a random linear map  $f : \mathbb{R}^d \rightarrow \mathbb{R}^k$  by  $f(u) = \frac{1}{\sqrt{k}} R \cdot u$ , where  $R \in \mathbb{R}^{k \times d}$  drawn elementwise from a standard normal distribution
- If  $u \in \mathbb{R}^d$ , show  $\mathbb{E}[\|f(u)\|_2^2] = \|u\|_2^2$
- Show that the random variable  $\|f(u)\|_2^2$  concentrates around  $\|u\|_2^2$ , and construct a union bound over all pairwise distances.



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# Deterministic Interpolative Decomposition

Given a matrix  $A \in \mathbb{R}^{m \times n}$  we can come up with a low-rank matrix approximation that uses  $A$ 's own columns. The ID can be computed using the column-pivoted  $QR$  factorization:

$$AP = QR.$$

To obtain our low-rank approximation we form the submatrix  $Q_k$  formed by the first  $k$  columns of  $Q$ . Thus we have the approximation:

$$A \approx Q_k Q_k^* A$$

which gives us a particular rank  $k$  projection of  $A$ .

# Randomized Interpolative Decomposition

This method constructs a subset  $S$  of randomly selected distinct  $p > k$  columns from the  $n$  columns of  $A$ . The algorithm then performs the column-pivoted  $QR$  factorization on the  $p$  columns of  $A$ :

$$A_{(:,S)}P = QR$$

Accordingly we have the following rank  $k$  projection of  $A$ :

$$A \approx Q_k Q_k^* A,$$

where  $Q_k$  is the submatrix formed by the first  $k$  columns of  $Q$ .



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# Deterministic Singular Value Decomposition

- Recall the singular value decomposition of a matrix:

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

where  $U$  and  $V$  are orthogonal matrices, and  $\Sigma$  is a diagonal matrix with positive diagonal entries  $\sigma_1 \geq \sigma_2 \geq \cdots \geq \sigma_r$ , where  $r$  is the rank of the matrix  $A$ .

- The  $\sigma_i$ s are called the singular values of  $A$ .

# Randomized Singular Value Decomposition

Utilizing ideas from [HMT09], our algorithm executes the following steps to compute the randomized SVD:

- 1 Construct a  $n \times k$  random Gaussian matrix  $\Omega$
- 2 Form  $Y = A\Omega$
- 3 Construct a matrix  $Q$  whose columns form an orthonormal basis for the column space of  $Y$
- 4 Set  $B = Q^*A$
- 5 Compute the SVD:  $B = U'\Sigma V^*$ ,
- 6 Construct the SVD approximation:  $A \approx QQ^*A = QB = QU'\Sigma V^*$

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# Results - Testing $620 \times 187500$ Matrix

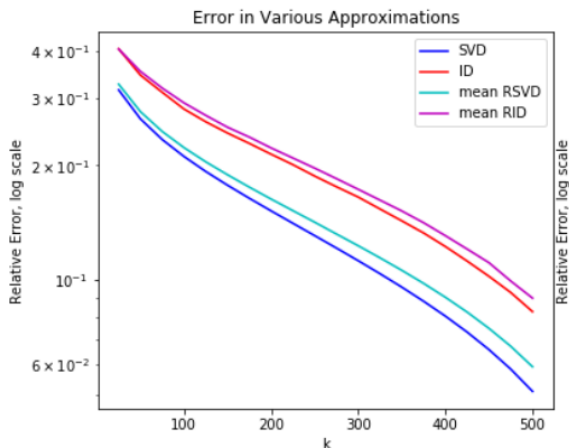


Figure: Error Relative to Original Data

# Results - Testing $620 \times 187500$ Matrix

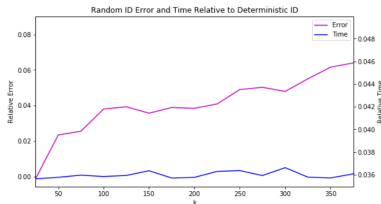


Figure: Random ID Error and Time Relative to Deterministic ID

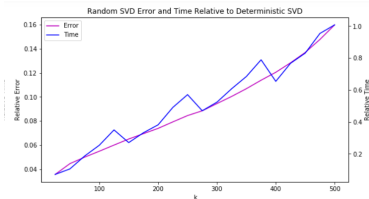


Figure: Random SVD Error and Time Relative to Deterministic SVD

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# Eigenfaces

- Using ideas from [BKP15] our eigenfaces experiment tests the LFW dataset [Hua+07]. This dataset contains more than 13,000 RGB images of faces where each image is a  $250 \times 250$ .
- We can flatten each image to represent it as vector of length  $250 \cdot 250 \cdot 3 = 187500$ .
- In our experiment we will only use 620 images from the LFW dataset. This gives us a data matrix  $A$  of size  $187500 \times 620$ .
- We then can perform SVD on the mean-subtracted columns of  $A$ .

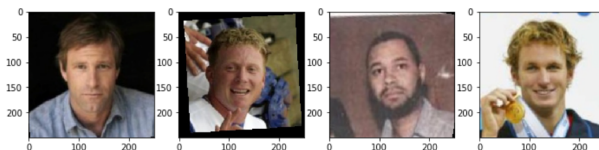


Figure: Original LFW Images



# Image Results

We obtain the following eigenfaces from the columns of the matrix  $U$ :

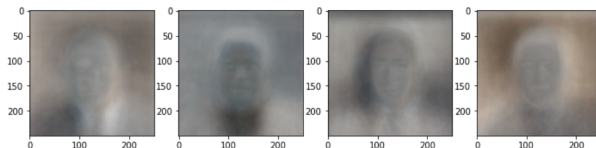


Figure: Eigenfaces Obtained using Deterministic SVD

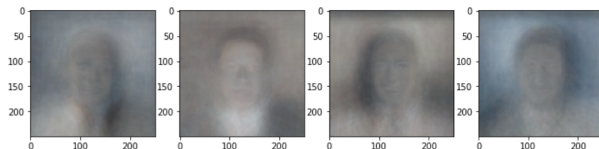


Figure: Eigenfaces Obtained using Randomized SVD

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- Kernel methods describe mapping the data into a high-dimensional space to add more structure and encourage linear separability.

$$\phi : \mathbb{R}^n \rightarrow \mathbb{R}^m, \quad m > n$$

- The 'kernel trick' allows for only the inner products to be computed, as opposed to the explicit high-dimensional mappings.

$$k(\mathbf{x}, \mathbf{y}) = \langle \phi(\mathbf{x}), \phi(\mathbf{y}) \rangle$$

- Kernel methods include kernel PCA, kernel SVM, and more.

# Randomized Fourier Features Kernel

Introduced in [RR08], we can sample random Fourier features to approximate a kernel. Let  $k(\mathbf{x}, \mathbf{y})$  denote our kernel, and  $p(w)$  the probability distribution corresponding to the inverse Fourier transform of  $k$ .

$$\begin{aligned} k(\mathbf{x}, \mathbf{y}) &= \int_{\mathbb{R}^d} p(w) e^{-jw^T(\mathbf{x}-\mathbf{y})} dw \\ &\approx \frac{1}{m} \sum_{i=1}^m \cos(w_i^T \mathbf{x} + b_i) \cos(w_i^T \mathbf{y} + b_i) \end{aligned}$$

where  $w_i \sim p(w)$ ,  $b_i \sim \text{Uniform}(0, 2\pi)$ . For a given  $m$ , define

$$z(\mathbf{x}) = \sum_{i=1}^m \cos(w_i^T \mathbf{x} + b_i)$$

to yield the approximation  $k(\mathbf{x}, \mathbf{y}) \approx \frac{1}{m} z(\mathbf{x}) z(\mathbf{y})^T$ .

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# Data for Kernel PCA Experiments

To test kernel PCA methods, we used a dataset that is not linearly separable— a cloud of points surrounded by a circle:

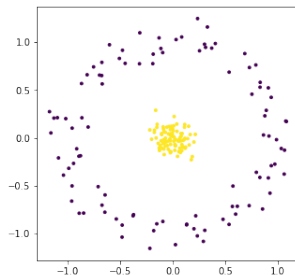


Figure: Data used to test kernel PCA methods

# Randomized Kernel PCA Results

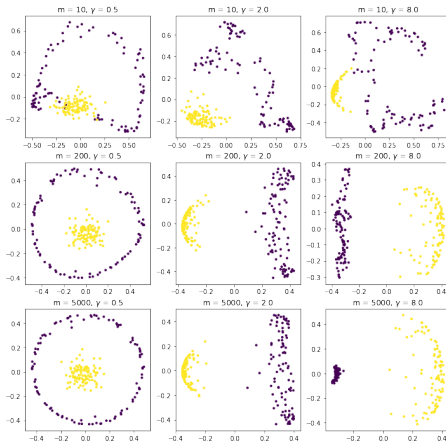


Figure: Random Fourier features KPCA results

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- We may also use kernel methods for support vector machines (SVM).
- The goal of an SVM is to find the  $(d - 1)$ -hyperplane that best separates two clusters of  $d$ -dimensional data points.
- In two dimensions, this is a line separating two clusters of points in a plane.
- Using the kernel trick, we can project inseparable points into a higher dimension and run an SVM algorithm on the resulting points.

# Randomized Kernel SVM

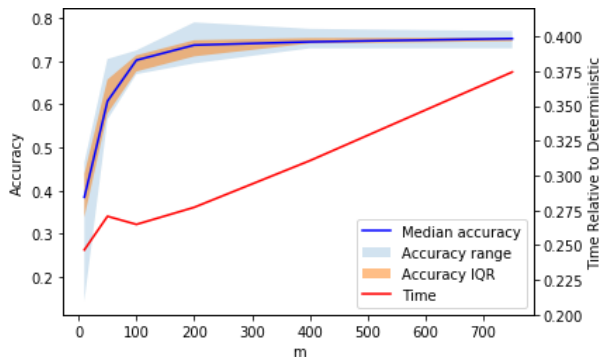


Figure: Randomized Kernel SVM Accuracy and time results as  $m$  varies

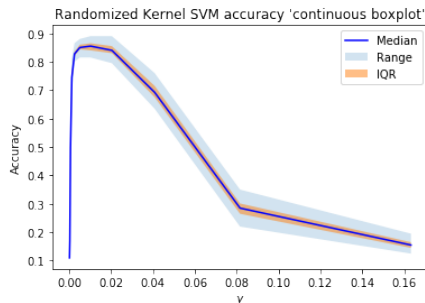
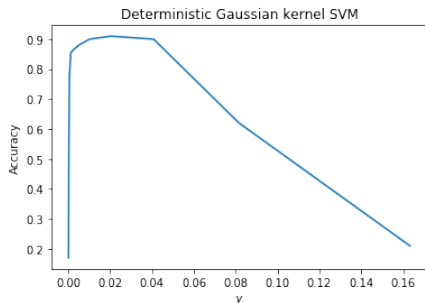
# Comparison of Deterministic and Randomized Kernel SVM

Using the MNIST dataset [LC10] we test 10000 images (784 features), for a **fixed**  $\gamma$ :

- Deterministic Kernel
  - Accuracy: 0.9195
  - Time: 37.99s
- Randomized Kernel
  - Accuracy: Mean: 0.891, St. dev. 0.0042  
Min: 0.881, Max: 0.9005
  - Mean Time: 2.14s

# Comparison of Deterministic and Randomized Kernel SVM

On 1000 MNIST images, we plot the accuracies of the deterministic and random kernel SVMs as  $\gamma$  varies:



# Application of Randomized Kernel SVM: Grid Search

$$\hat{\mathbf{K}} = \frac{1}{m} \mathbf{z}(\mathbf{X}) \mathbf{z}(\mathbf{X})^T$$

Testing 100  $\gamma$  values to identify the best one:

- Deterministic Kernel, Series: 133.03s
- Randomized Kernel, Series: 78.97s
- Randomize Kernel, Parallel: 41.18s
- Best  $\gamma$  value obtained from randomized method corresponds with either best or second best deterministic  $\gamma$  (3 trials)

# References



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