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## HPC for numerical methods and data analysis

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### Randomized low rank approximation on MNIST data set

The Radial Basis Function (RBF) applications can be found in neural networks, data visualization, surface reconstruction, etc. These techniques are based on collocation in a set of scattered nodes, the **computational cost of these techniques increase** with the number of points in the given dataset with the dimensionality of the data.

For RBF approximation we assume that we have an unordered dataset  $\{x_i\}_1^n$ , each point associated with a given  $f_i \in \mathbb{R}^p$ . We are going to consider  $f_i \in \mathbb{R}$  i.e. each point in the dataset is associated with a label. The approximation scheme can be written as follows:

$$s(x) = \sum_{i=1}^n \lambda_i \phi(\|x - x_i\|_2),$$

where:

- $x_i$  are the data points
- $x$  is a free variable at which we wish to evaluate the approximation
- $\phi$  is the RBF
- $\lambda_i$  are the scalar parameters

The  $\lambda_i$ 's are chosen so that  $s$  approximates  $f$  in a desired way. One of the simplest ways of computing these parameters is by forcing the interpolation to be exact at  $x_i$  i.e.  $s(x_i) = f(x_i) = f_i$ . Define a matrix  $A \in \mathbb{R}^{n \times n}$  such that  $A_{ij} = \phi(\|x_i - x_j\|_2)$ , let  $\lambda = [\lambda_1, \dots, \lambda_n]^\top \in \mathbb{R}^n$  and  $f = [f_1, \dots, f_n]^\top \in \mathbb{R}^n$  (both column vectors). In order to compute the scalar parameters we need to solve the following linear system:

$$A\lambda = f. \tag{1}$$

Before computing  $A$ , answer the following questions:

- a) How does the computational cost of solving (1) scale in both the number of data points and the dimension of such points?
- b) What would it mean if  $A$  is nearly singular?
- c) What would be the effect on  $A$  if  $\phi$  has compact support? What would be the disadvantage of using such RBF?

The *MNIST data set* contains pictures of handwritten digits. It contains 60'000 training images and 10'000 testing images. You can download this database from here: <https://www.csie.ntu.edu.tw/~cjlin/libsvmtools/datasets/>. You can also download the labels for the training and testing images (these are going to be our  $f_i$ 's. We are going to use the following RBF:

$$\phi(\|x_i - x_j\|_2) = e^{-\|x_i - x_j\|_2/c},$$

with  $c > 0$ .

- d) We are going to start by taking a relatively small sample of the training set (i.e.  $n$  being "small"). Download the data set (both the test and training sets). From the training set (and labels) pick the  $n$  top rows.
- e) Write a Python scrip that computes  $A$  using the subsampled data set and optionally saves it to memory (try using the pickle package on Python). Test different values of  $c$  to solve (1). (Optional: write a parallel implementation of the function to build  $A$ )
- f) Explain Nyström approximation and why it would be useful in this setting.
- g) Given a sketching matrix  $\Omega$  use your code from last week for different values of  $l$  and compute  $A_{\text{Nyst}} = (A\Omega)(\Omega^\top A\Omega)^\dagger(\Omega^\top A)$ .
- h) Test the accuracy of the previously computed Nyström approximation. Provide graphs that show the error of the approximation using the nuclear norm. Compare these errors with the best rank  $k$  approximation of  $A$ .
- i) (Optional) Try solving (1) using  $A_{\text{Nyst}}$