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High-Dimensional Approximation

Part 3: Approximation from samples

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Consider the approximation of functions $f : \mathcal{X} \to \mathbb{R}$ from a set $K \subset X$ using *n* information

 $\ell_1(f),\ldots,\ell_n(f)$

that can be deterministic or random.

When $\ell_i : X \to \mathbb{R}$ are linear (or affine) maps, we talk about linear (or affine) information.

Type of information

A particular type of linear information is point evaluations (aka standard information)

 $\ell_i(f) = f(x_i)$

Another type of linear information is

$$\ell_i(f) = \int_{\mathcal{X}} \psi_i(x) f(x) d\mu(x)$$

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If f is known to satisfy an equation

$$B(f) = b$$

with given right-hand side $b \in Z' \subset \mathbb{R}^X$ and operator $B : X \to Z'$, we can have access to the information

$$\ell_i(f) = B(f)(x_i), \text{ or } \ell_i(f) = \langle \psi_i, B(f) \rangle$$

for some function $\psi_i \in Z$. For linear (resp. nonlinear) operator *B*, this corresponds to linear (resp. nonlinear) information. This is the framework of Galerkin or variational methods for PDEs, Physics-informed machine learning (Deep-Galerkin, Deep-Ritz, PINN, ...).

We distinguish two different settings

- information is given (passive learning)
- information can be freely generated (active learning), a typical setting in computer/physical experiments, numerical analysis of PDEs, or scientific machine learning.

Algorithm

Given information $\ell(f) = (\ell_1(f), \dots, \ell_n(f))$, an algorithm returns an approximation

 $A(\ell(f))$

in a subset of X, where the map A is related to the choice of a restricted model class (or approximation tool).

A linear algorithm, with A also a linear map, corresponds to linear approximation:

$$A(\ell(f)) = \sum_{i=1}^{n} a_i(\ell(f))\varphi_i$$

where the a_i are linear maps and $span\{\varphi_1, \ldots, \varphi_n\}$ is the range of A.

It could be a low-dimensional manifold V_m (model class) that is known to approximate well the set K, or a sequence of models with increasing complexity $(V_m)_{m\geq 1}$ (approximation tool) that is known to approximate the manifold with a good rate of convergence.

• For K a ball of Sobolev or Besov spaces: splines (with fixed or adaptive mesh) or wavelets (with or without sparsity)

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- For a larger class of sets K: neural networks or tensor networks
- For more general manifolds K, V_m can be obtained by manifold approximation (or dimension reduction) methods

Approximation in a given model class

For a given model class V_m and given information $z = \ell(f)$, an approximation $f_m = A(z) \in V_m$ may be defined by

$$\ell(f_m) = z. \tag{1}$$

If for any z there exists a unique element f_m in V_m satisfying (1), we say that ℓ is unisolvent for V_m . When information are point evaluations, this corresponds to interpolation. When information are linear functionals of an equation's residual, this corresponds to (Petrov-)Galerkin projection.

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More generally, $f_m = A(z)$ can be defined as a solution of

 $\min_{f_m \in V_m} d(\ell(f_m), z),$

and in particular

$$\min_{f_m \in V_m} \sum_{i=1}^n w_i (\ell_i(f_m) - z_i)^2$$

When information are point evaluations, this corresponds to (weighted) least-squares approximation. When information are linear functionals of an equation's residual, this corresponds to (Petrov-)Galerkin projection.

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- When the information can be generated (active learning), a fundamental question is how to generate a good information for a given model class V_m . Adaptive strategies play with a collection of model classes $(V_m)_{m\geq 1}$ and generate information adaptively. A question is then to recycle information in order to obtain a near-optimal performance in terms of complexity.

- Manifold approximation
- 2 Linear approximation from point evaluations
- 3 Tensor networks approximation with point evaluations

Outline

1 Manifold approximation

2 Linear approximation from point evaluations

3 Tensor networks approximation with point evaluations

Manifold approximation

Assume we want to approximate (or recover) functions from a general manifold K in a vector space X. If K can be sampled, a suitable low-dimensional model class V_m (or sequence of model classes) can be obtained by manifold approximation (or dimension reduction) methods using samples from K.

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Typical model classes V_m include

• Low-dimensional linear/affine spaces

$$V_m = \{g(a) : a \in \mathbb{R}^m\}, \quad \text{with } g : \mathbb{R}^m \to X \text{ linear/affine}$$



Manifold approximation

• Union of low-dimensional linear spaces

$$V_m = \bigcup_{k=1}^m W_k$$



• Manifold $V_m = \{g(a) : a \in \mathbb{R}^m\}$ with continuous parametrization map $g : \mathbb{R}^m \to X$.



A typical setting is when K is the set of trajectories of a random process or more generally the range of some function-valued random variable. A possible dimension reduction method is principal component analysis (for linear approximation).

Another setting is the solution of forward or inverse problems of parameter-dependent equations where $K = \{u(y) : y \in Y\}$ is the manifold of solutions. Manifold approximation is called model order reduction (reduced basis, POD, ...).

Let Y be equipped with a probability measure μ and X a Hilbert space, and $K = \{u(y) : y \in Y\}$ with u a map in the Bochner space $L^2(Y; X)$.

The optimal performance of a linear approximation of K is measured in mean-squared error by

$$d_{m}^{2,\mu}(K)_{X} = \inf_{\dim(V_{m})=m} \int_{Y} E(u(y); V_{m})_{X}^{2} d\mu(y) = \inf_{\dim(V_{m})=m} \mathbb{E}_{y \sim \mu}(\|u(y) - P_{V_{m}}u(y)\|_{X}^{2})$$

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An optimal subspace V_m is given by principal component analysis (PCA), where V_m is the dominant eigenspace of the self-adjoint compact operator $T : v \mapsto \mathbb{E}_{y \sim \mu}((u(y), v)_X u(y))$ and the error is

$$\mathbb{E}_{y\sim\mu}(\|u(y)-P_{V_m}u(y)\|_X^2)=\sum_{i>m}\lambda_i$$

where $(\lambda_i)_{i \ge 1}$ is the decreasing sequence of eigenvalues of T. This is related to singular value decomposition (or Karhunen-Loeve decomposition) of $u \in L^2(Y) \otimes X$,

$$u(y) = \sum_{i\geq 1} \sqrt{\lambda_i} \varphi_i a_i(y), \quad P_{V_m} u(y) = \sum_{i=1}^m \sqrt{\lambda_i} \varphi_i a_i(y)$$

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PCA even provides a hierarchical sequence of model classes $(V_m)_{m\geq 1}$.

An estimation of V_m is given by empirical PCA which consists in solving

$$\min_{\dim(V_m)=m} \frac{1}{n} \sum_{i=1}^n \|u(y_i) - P_{V_m} u(y_i)\|_X^2$$

where the y_i are samples in Y and the $u(y_i)$ are the corresponding samples in K. The solution is the dominant eigenspace of the operator

$$T_n: v \mapsto \frac{1}{n} \sum_{i=1}^n u(y_i)(u(y_i), v)_X.$$



For an analysis of empirical PCA, see e.g. [3, 4].

Assuming X is finite dimensional with orthonormal basis $(e_i)_{1 \le i \le N}$, $u(y) = \sum_{i=1}^N a_i(y)e_i$, and a basis of V_m is given by the dominant eigenvectors of the matrix

$$\frac{1}{n}\sum_{i=1}^n a(y_i)a(y_i)^T.$$

This is equivalent to obtain the dominant left singular vectors of the matrix

$$A = (a(y_1), \ldots a(y_n)) \in \mathbb{R}^{N \times n}$$

Optimal sampling strategy have been proposed for singular value decomposition of matrices. This requires an estimation of dominant right singular vectors.

Given a set K from a Banach space X, the optimal performance of linear approximation in worst case error is measured through the Kolmogorov width

$$d_n(K)_X = \inf_{\dim(V_m)=m} \sup_{u\in K} E(u, V_m) \quad \text{with} \quad E(u, V_m)_X := \inf_{v\in V_m} \|u-v\|_X$$

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Greedy algorithms can be used to the construction of a hierarchical sequence of spaces $(V_m)_{m\geq 1}$ using samples (snapshots) from K. Spaces are defined by $V_m = span\{u_1, \ldots, u_m\}$ where $(u_i)_{i\geq 1}$ is a sequence from K selected greedily.

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Given V_m , u_{m+1} is the element which provides the highest error of approximation by V_m

$$E(u_{m+1}, V_m)_X = \max_{u \in K} E(u, V_m)_X$$



When $K = \{u(y) : y \in Y\}$, $u_{m+1} = u(y_{m+1})$ where the parameter value y_{m+1} is such that $y_{m+1} \in \arg \max_{y \in Y} E(u(y), V_m)_X$

In practice, for a computationally feasible algorithm, $E(u(y), V_m)_X$ is replaced by some error estimate $\Delta(u(y), V_m)$, and the maximum is taken over a finite training set in Y (possibly random [Cohen et al 2020]).

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A typical setting is when $K = \{u(y) : y \in Y\} \subset X$ is the solution of some parameter dependent equation

$$R(u(y);y)=0$$

Here $\Delta(u(y), V_m)$ is typically defined as some residual norm

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Randomized linear algebra can be used for an efficient and stable estimation of residual norms [Balabanov and Nouy 2021a], and for the construction of preconditioners [Balabanov and Nouy 2021b].

This yield a suboptimal selection of u_{m+1} satisfying

$$E(u_{m+1}, V_m)_X \geq \gamma \max_{u \in K} E(u, V_m)_X, \quad \gamma \leq 1.$$

This algorithm therefore generates a suboptimal sequence of spaces yielding a worst case error

$$\sigma_m(K)_X := \sup_{u \in K} E(u, V_m)_X \ge d_m(K)_X$$

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Assuming $\gamma \geq 1$ is independent of *m*, the algorithm is a weak greedy algorithm for which results have been obtained in [DeVore et al 2013].

For X a Hilbert space, it holds

•
$$\sigma_{2m}(K)_X \leq \sqrt{2}\gamma^{-1}\sqrt{d_m(K)_X}$$

- If $d_m(K)_X \leq C_0 m^{-lpha}$ then $\sigma_m(K)_X \leq C_1 m^{-lpha}$
- If $d_m(K)_X \leq C_0 e^{-c_0 m^{lpha}}$ then $\sigma_m(K)_X \leq C_1 e^{-c_1 m^{lpha}}$

For X a Banach space, similar but slightly worse results hold.
Multi-space approximation

h or h-p reduced basis methods [Eftang et al 2010] are multi-space approximation methods that consist is partitioning the manifold K (or corresponding parameter set Y) into subsets K_k , and approximating each subset by a linear space W_k of fixed dimension (*h* method) or variable dimension (h-p method).



These methods requires a partitioning (or clustering) strategy.

Dictionary-based multi-space approximation

Multiple spaces can be extracted from a dictionary $\mathcal{D} = \{u_1, \ldots, u_N\}$ of samples from K. By considering subspaces with dimension less than m, this yields the model class

$$V_m := V_m(\mathcal{D}) = \bigcup_{\alpha \in \{1, \dots, N\}^m} W_\alpha(\mathcal{D}), \quad W_\alpha(\mathcal{D}) = span\{u_{\alpha_1}, \dots, u_{\alpha_m}\}$$



This is equivalent to *m*-term approximation

$$V_m = \{g(a) := \sum_{i=1}^N a_i u_i : a \in \mathbb{R}^N, \|a\|_0 \leq m\}.$$

The dictionary (samples) can be taken arbitrarily or generated with a greedy procedure proposed in [Balabanov and Nouy 2021a], using randomized linear algebra for handling large dictionaries.

Several approaches exist for the approximation of a set ${\cal K}$ by a parametrized nonlinear manifold of the form

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Neural networks are popular tools for this task.

For $X = \mathbb{R}^N$, a neural network representation can be used for $g : \mathbb{R}^m \to \mathbb{R}^N$.



Learning a map g from samples from K can be done (offline) by learning a compositional function (or autoencoder) $g \circ h$, where both functions $h : \mathbb{R}^N \to \mathbb{R}^m$ (the encoder) and $g : \mathbb{R}^m \to \mathbb{R}^N$ (the decoder) can be represented by neural networks.



Given samples $\{u_1, \ldots, u_n\} \subset K$, h and g can be obtained by minimizing

$$\sum_{i=1}^{n} \|u_i - g \circ h(u_i)\|_X^2$$
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This methodology is not restricted to the use of neural networks for h and g.

For *h*, one can use a linear map (a matrix of size $N \times m$), so that $g \circ h$ corresponds to a ridge approximation.

Note that if h and g are restricted to be linear maps (or matrices of size $N \times m$ and $m \times N$ respectively), it boils down to linear approximation learned by PCA.

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Note that if *h* and *g* are restricted to be linear maps (or matrices of size $N \times m$ and $m \times N$ respectively), it boils down to linear approximation learned by PCA.

A two-step strategy can be used, by first learning a composition of linear maps $\tilde{g} \circ h$ by PCA, or another algorithm for linear approximation, and then learning $g \circ h$ with a fixed h by solving (2).

If we know that $K = \{u(y) : y \in Y\}$ the image through a map u of a low-dimensional space Y, we can learn the map g from samples in Y by learning a compositional function

 $g \circ h$

where $h: Y \to \mathbb{R}^m$. Given samples y_1, \ldots, y_n in Y, this can be done by minimizing

$$\sum_{i=1}^{n} \|u(y_i) - g \circ h(y_i)\|_X^2$$



If K in an infinite dimensional space X, a discretization is required.

A discretization can be represented by some encoder-decoder pair (E, R) with $E: X \to \mathbb{R}^N$ and $D: \mathbb{R}^N \to X$ (e.g. *E* could provide the values E(u) of a function at the nodes of a mesh, and D(E(u)) a spline interpolation), and the functions *g* and *h* can be learned by solving



The map $D \circ g \circ h \circ E$ is called an Operator Network that aims at approximating the identity map from K to X.

For K a set of functions defined on a domain $\mathcal X,$ with values in $\mathbb R,$ an alternative is to consider

$$V_m = \{g(\cdot, a) : x \mapsto g(x, a) : a \in \mathbb{R}^m\}$$

with $g : \mathcal{X} \times \mathbb{R}^m \to \mathbb{R}$ in some high-dimensional approximation format (e.g. neural or tensor networks).

Function g can be learned (offline) from samples in K by solving

$$\min_{h,g} \sum_{i=1}^{n} \|u_i - g(\cdot, h(E(u_i)))\|_X^2$$

where $E: K \to \mathbb{R}^N$ is some fixed discretization map (encoder) and $h: \mathbb{R}^N \to \mathbb{R}^m$. Here, no explicit decoder is used.

1 Manifold approximation

2 Linear approximation from point evaluations

3 Tensor networks approximation with point evaluations

We consider the approximation of functions from a set

$$K \subset X \subset \mathbb{R}^{\mathcal{X}}$$

using point evaluations (standard information) and linear algorithms (linear approximation).

The best we can expect for the linear approximation of functions from a set K is characterized by sampling numbers $\rho_n(K)_X$ (for deterministic setting) or $\rho_n^{rand}(K)_X$ (for randomized setting) (see Part 1).

We assume that we are given a *m*-dimensional linear space V_m that is supposed to approximate well the set K.

The question is how to generate good points in \mathcal{X} that allow to obtain an approximation in V_m with an error close to the best approximation error.

Interpolation

For a set of points $\mathbf{x} = (x_1, \ldots, x_m)$ unisolvent for V_m , we let $\mathcal{I}_{V_m} : X \to V_m$ be the corresponding interpolation (linear) operator.

We have

$$\|f-\mathcal{I}_{V_m}f\|_X\leq \left(1+\|\mathcal{I}_{V_m}\|
ight)\inf_{v\in V_m}\|f-v\|_X$$

For $(X, \|\cdot\|_{\infty})$ the set of functions with bounded norm $\|f\|_{\infty} := \sup_{x \in \mathcal{X}} |f(x)|, \|\mathcal{I}_{V_m}\|$ is the Lebesgue constant, with

$$\|\mathcal{I}_{V_m}\| = \sup_{x \in \mathcal{X}} \sum_{i=1}^m |L_i(x)|$$

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For univariate functions and classical spaces V_m (polynomials, splines), the theory is well established and suitable choices of points are available.

Except in very specific cases (e.g. piecewise constant or linear approximation), $\|\mathcal{I}_{V_m}\|$ grows with m. The question is to find good points such that $\|\mathcal{I}_{V_m}\|$ grows not too fast with m.

Given a space V_m with basis $\varphi_1, \ldots, \varphi_m$, a general greedy algorithm has been proposed in [Maday et al 2009] to construct interpolation points, called magic points.

The idea is to construct a good sequence of spaces $W_k = span\{\psi_1, \ldots, \psi_k\}$ for the approximation of the discrete set $\{\varphi_i : 1 \le i \le m\}$ in $(X, \|\cdot\|_{\infty})$, and associated interpolation points.

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Starting from $V_0 = \{0\}$, we define

$$i_k \in \arg \max_{1 \le i \le m} \|\varphi_i - \mathcal{I}_{W_{k-1}}\varphi_i\|_{\infty}, \quad \psi_k = \varphi_{i_k} - \mathcal{I}_{W_{k-1}}\varphi_{i_k}$$

where $\mathcal{I}_{W_{k-1}}$ is the interpolation onto W_{k-1} using points (x_1, \ldots, x_{k-1}) , and define

$$x_k \in \arg \max_{x \in \mathcal{X}} |\psi_k(x)|.$$

Empirical interpolation



Figure: Polynomial space $V_m = \mathbb{P}_9$ on [-1, 1]. Function $|\psi_k(x)|$ and corresponding interpolation point $x_k = \arg \max_x |\psi_k(x)|$

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Figure: Polynomial space $V_m = \mathbb{P}_9$ on [-1, 1]. Function $|\psi_k(x)|$ and corresponding interpolation point $x_k = \arg \max_x |\psi_k(x)|$

In the context of adaptive approximation in a sequence of spaces $V_1 \subset \ldots \subset V_m \subset \ldots$, and in order to recycle interpolation points, we modify the algorithm by simply taking $W_k = V_k$.

Letting $V_0 = \{0\}$, we define

$$\psi_k = \varphi_k - \mathcal{I}_{V_{k-1}}\varphi_k$$

where $\mathcal{I}_{V_{k-1}}$ is the interpolation onto V_{k-1} using points (x_1, \ldots, x_{k-1}) , and define

$$x_k \in \arg \max_{x \in \mathcal{X}} |\psi_k(x)|.$$

Empirical interpolation — adaptive setting



Figure: Polynomial space $V_m = \mathbb{P}_9$ on [-1, 1]. Function $|\psi_k(x)|$ and corresponding interpolation point $x_k = \arg \max_x |\psi_k(x)|$

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Another strategy can be defined as follows. Let $\varphi(x) = (\varphi_1(x), \dots, \varphi_m(x)) \in \mathbb{R}^m$, where $\varphi : \mathcal{X} \to \mathbb{R}^m$ is the feature map associated with V_m . The feature space \mathbb{R}^m is equipped with the Euclidian norm $\|\cdot\|$.

The idea is to construct an increasing sequence of spaces

$$U_k = span\{arphi(x_1), \ldots, arphi(x_k)\} \subset \mathbb{R}^m$$

for the approximation of the manifold $\{\varphi(x) : x \in \mathcal{X}\}$.

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for the approximation of the manifold $\{\varphi(x) : x \in \mathcal{X}\}$.

Starting from $U_0 = \{0\}$, we define

$$x_k \in rg\max_{x\in\mathcal{X}} \Lambda_k(x), \quad \Lambda_k(x) = \|arphi(x) - P_{U_{k-1}}arphi(x)\|_2^2$$

where $P_{U_{k-1}}$ is the orthogonal projection from \mathbb{R}^m to U_{k-1} .

Let (e_1, \ldots, e_m) be the orthonormal basis of \mathbb{R}^m defined by

$$\boldsymbol{e}_k \propto \boldsymbol{\varphi}(x_k) - P_{U_{k-1}} \boldsymbol{\varphi}(x_k), \quad \|\boldsymbol{e}_k\|_2 = 1.$$

If V_m is a Hilbert space and the functions φ_i form an orthonormal basis of V_m , then the functions $\psi_i(x) = \varphi(x)^T \boldsymbol{e}_i$ also form an orthonormal basis of V_m and

$$\Lambda_k(x) = \sum_{i=k}^m \psi_i(x)^2 = \|\varphi(x)\|_2^2 - \sum_{i=1}^{k-1} \psi_i(x)^2$$



Figure: Polynomial space $V_m = \mathbb{P}_9$ on [-1, 1]. Function $\Lambda_k(x)$ and corresponding interpolation point $x_k = \arg \max_x \Lambda_k(x)$



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Figure: Haar wavelets space V_m on [0, 1], with resolution 5. Function $\Lambda_k(x)$ and corresponding interpolation point $x_k = \arg \max_x \Lambda_k(x)$.



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(c) k = 3

(d) k = 4

Figure: Bivariate polynomial space $V_m = \mathbb{P}_4$ on $[-1, 1]^2$. Function $\Lambda_k(x)$ and corresponding interpolation point $x_k = \arg \max_x \Lambda_k(x)$.



Figure: Bivariate polynomial space $V_m = \mathbb{P}_4$ on $[-1,1]^2$. Function $\Lambda_k(x)$ and corresponding interpolation point $x_k = \arg \max_x \Lambda_k(x)$.

In the context of adaptive approximation in a sequence of spaces $V_1 \subset \ldots \subset V_m \subset \ldots$, and in order to recycle interpolation points, we modify the algorithm by considering at step k the feature map φ associated with the basis of V_k .

Empirical interpolation based on feature map — adaptive setting



Figure: Polynomial space $V_m = \mathbb{P}_9$ on [-1,1]. Function $\Lambda_k(x)$ and corresponding interpolation point $x_k = \arg \max_x \Lambda_k(x)$.

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Interpolation in RKHS

A reproducing kernel Hilbert space (RKHS) H is a Hilbert space of functions defined on \mathcal{X} such that the point evaluation $\delta_x : f : x \mapsto f(x)$ is a continuous linear map. There is a so called reproducing kernel k such that $k(x, \cdot)$ is the Riesz representer of δ_x , that is

$$f(x) = (f, k(x, \cdot))_{H},$$

where $(\cdot, \cdot)_H$ is the inner product on *H*.

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For given points $\mathbf{x} = (x_1, \dots, x_k)$, the interpolation operator \mathcal{I}_{W_k} onto the space $W_k = span\{k(\cdot, x_1), \dots, k(\cdot, x_k)\}$ is defined by

$$\mathcal{I}_{W_k}f(x) = k(x, \mathbf{x})k(\mathbf{x}, \mathbf{x})^{-1}f(\mathbf{x})$$

where $k(x, y) = (k(x_i, y_j))_{i,j}$ and $f(x) = (f(x_j))_j$.
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where $k(\mathbf{x}, \mathbf{y}) = (k(x_i, y_j))_{i,j}$ and $f(\mathbf{x}) = (f(x_j))_j$. The operator \mathcal{I}_{W_k} is the

H-orthogonal projection onto W_k , which provides the element of best approximation of a function in W_k . Indeed, for $f \in H$, the interpolation conditions

$$\mathcal{I}_{W_k}f(x_i) = f(x_i), \quad 1 \leq i \leq k,$$

are equivalent to

$$(k(\cdot, x_i), \mathcal{I}_{W_k}f - f)_H = 0, \quad 1 \le i \le k,$$

that is $\mathcal{I}_{W_k}f - f$ is orthogonal to W_k .

The error of interpolation at point $x \in \mathcal{X}$ is such that

$$|f(x) - \mathcal{I}_{W_{k}}f(x)| = |(k(x, \cdot), \mathcal{I}_{W_{k}}f - f)_{H}| = |(k(x, \cdot) - \mathcal{I}_{W_{k}}k(x, \cdot), \mathcal{I}_{W_{k}}f - f)_{H}| \leq ||k(x, \cdot) - \mathcal{I}_{W_{k}}k(x, \cdot)||_{H}||f||_{H}$$

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A natural definition of a new basis function $k(x_{k+1}, \cdot)$ is to consider a point x_{k+1} where the error bound is maximum, that is

$$x_{k+1} \in \arg \max_{x \in \mathcal{X}} \Lambda_k(x),$$

with

$$\Lambda_k(x) = \|k(x,\cdot) - \mathcal{I}_{W_k}k(x,\cdot)\|_H^2 = k(x,x) - k(x,x)k(x,x)^{-1}k(x,x).$$

A finite dimensional space V_m with basis $\varphi_1, \ldots, \varphi_m$ defines a RKHS with kernel

$$k(x,y) = \varphi(x)^T \varphi(y), \quad \varphi(x) := (\varphi_1(x), \dots, \varphi_m(x))$$

A sequential interpolation method consists in defining a sequence of points $(x_k)_{k\geq 1}$ and corresponding spaces $W_k = span\{k(x_1, \cdot), \ldots, k(x_k, \cdot)\}$ such that

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with $x = (x_1, \dots, x_k)$ and $\varphi(x) = (\varphi_i(x_j))_{1 \le i,j \le k}$.

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In bayesian regression with gaussian processes (with noisy-free observations), the function $\Lambda_k(x)$ is the variance of the conditional gaussian process given observations at points $\mathbf{x} = (x_1, \dots, x_k)$.

Note that the obtained sequence of points only depends on the space V_m .

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Note that the obtained sequence of points only depends on the space V_m .

Letting $U_k = span\{\varphi(x_1), \ldots, \varphi(x_k)\} \subset \mathbb{R}^m$, we note that

$$\Lambda_k(x) = \|\varphi(x) - P_{U_{k-1}}\varphi(x)\|_2^2$$

This is equivalent to the previously presented empirical interpolation based on feature map.

Consider the approximation of a function f in $X = L^2_{\mu}(\mathcal{X})$ equipped with the norm

$$\left\|f\right\|^{2} = \int f(x)^{2} d\mu(x)$$

Given a *m*-dimensional space V_m in $L^2_\mu(\mathcal{X})$, a weighted least-squares approximation $\hat{f}_m \in V_m$ is defined by minimizing

$$\frac{1}{n}\sum_{i=1}^{n}w_{i}(v(x_{i})-f(x_{i}))^{2}$$

over $v \in V_m$, for some suitably chosen points $\mathbf{x} = (x_1, \dots, x_n)$ and corresponding weights $\mathbf{w} = (w_1, \dots, w_n)$.

Least squares approximation

This is equivalent to minimize

$$\|f-v\|_n^2$$

where $\|\cdot\|_n^2$ is a semi-norm defined by

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Assuming that the x_i are i.i.d. samples from a distribution ν defined by

$$d\nu(x) = w(x)^{-1}d\mu(x),$$

and the weights $w_i = w(x_i)$, then for all $f \in L^2_\mu$

$$\mathbb{E}(\|f\|_{n}^{2}) = \mathbb{E}_{x \sim \nu}(w(x)f(x)^{2}) = \mathbb{E}_{x \sim \mu}(f(x)^{2}) = \|f\|^{2}$$

Least squares approximation

Given an L^2_{μ} -orthonormal basis $\varphi_1(x), ..., \varphi_m(x)$ of V_m , and letting $\varphi(x) = (\varphi_1(x), ..., \varphi_m(x))^T \in \mathbb{R}^m$, a function $v \in V_m$ can be written

$$\mathbf{v}(\mathbf{x}) = \sum_{i=1}^{m} a_i \varphi_i(\mathbf{x}) = \boldsymbol{\varphi}(\mathbf{x})^T \boldsymbol{a}$$

We have

$$||v||^2 = ||a||_2^2$$

and

$$\|v\|_n^2 = a^T G a$$

where G is the empirical Gram matrix (or weighted information matrix) given by

$$\boldsymbol{G} := \boldsymbol{G}(\boldsymbol{x}) = \frac{1}{n} \sum_{i=1}^{n} w_i \varphi(x_i) \varphi(x_i)^T.$$

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We have

$$\lambda_{\textit{min}}(\boldsymbol{G}) \| v \|^2 \leq \| v \|_n^2 \leq \lambda_{\textit{max}}(\boldsymbol{G}) \| v \|^2 \quad orall v \in V_m.$$

The quality of least-squares projection is related to how much ${\pmb G}$ deviates from the identity.

Optimal design of experiments

Consider the model

$$Y = f(X) + \epsilon$$

where $X \sim \mu$ and $\epsilon \sim \mathcal{N}(0, \lambda)$ is independent of X, that corresponds to noisy evaluations of a function f.

For given points $\mathbf{x} = (x_1, \dots, x_n)$ we have access to $\mathbf{y} = (y_1, \dots, y_n)$ such that

$$y_i = f(x_i) + \epsilon_i$$

with $\boldsymbol{\epsilon} = (\epsilon_1, \dots, \epsilon_n) \sim \mathcal{N}(0, \Lambda)$ independent of \boldsymbol{x} .

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with $\boldsymbol{\epsilon} = (\epsilon_1, \dots, \epsilon_n) \sim \mathcal{N}(0, \Lambda)$ independent of \boldsymbol{x} .

A weighted least-squares estimate \hat{f}_m is then obtained by solving

$$\min_{v\in V_m}\frac{1}{n}\sum_{i=1}^n w_i(v(x_i)-y_i)^2$$

Letting $\Phi := \Phi(x) = (\varphi_j(x_i))_{1 \le i \le n, 1 \le j \le m}$ (the design matrix) and W = diag(w) the weight matrix, we have

$$\hat{f}_m(x) = \boldsymbol{\varphi}(x)^T \hat{\boldsymbol{a}}, \quad \hat{\boldsymbol{a}} = \boldsymbol{G}^{-1} \boldsymbol{\Phi}^T \boldsymbol{W} \boldsymbol{y}$$

with

$$\boldsymbol{G} := \boldsymbol{G}(\boldsymbol{x}, \boldsymbol{w}) = \boldsymbol{\Phi}^{\mathsf{T}} \boldsymbol{W} \boldsymbol{\Phi}$$

For fixed x, the covariance of \hat{a} is

$$\textit{Cov}(\hat{\boldsymbol{a}}) = (\boldsymbol{\Phi}^{\mathsf{T}} \boldsymbol{W} \boldsymbol{\Phi})^{-1} \boldsymbol{\Phi}^{\mathsf{T}} \boldsymbol{W} \boldsymbol{\Lambda} \boldsymbol{W} \boldsymbol{\Phi} (\boldsymbol{\Phi}^{\mathsf{T}} \boldsymbol{W} \boldsymbol{\Phi})^{-1}$$

For $\Lambda = \lambda W^{-1}$, we obtain

$$\mathcal{C}ov(\hat{\pmb{a}}) = \lambda \pmb{G}^{-1}$$

and the variance of the prediction $\hat{f}_m(x)$ at some point x is

1

$$\mathbb{V}(\hat{f}_m(x)) = \lambda \varphi(x)^T \boldsymbol{G}^{-1} \varphi(x)$$

In order to minimize the variance for any $x \in \mathcal{X}$, that is for any $\varphi(x) \in \mathbb{R}^m$, we would like to minimize \mathbf{G}^{-1} over $x \in \mathcal{X}^n$ and $\mathbf{w} \in \mathbb{R}^n_+$, in the sense of the Loewner order, over the space S^+_m of symmetric positive semi-definite matrices. However, a global optimum does not necessarily exist since Loewner order is only a partial order.

A common approach is to consider as a proxy the minimization of a decreasing convex function $h: S_m^+ \to \mathbb{R}$, i.e. such that

$$h(\mathbf{A}) \leq h(\mathbf{B})$$
 for $\mathbf{A} \succcurlyeq \mathbf{B}$,

and solve

$$\min_{\boldsymbol{x},\boldsymbol{w}} h(\boldsymbol{G}(\boldsymbol{x},\boldsymbol{w}))$$

- E-optimal design corresponds $h(G) = \lambda_{max}(G^{-1}) = \lambda_{min}(G)^{-1}$
- A-optimal design corresponds to $h(\mathbf{G}) = Tr(\mathbf{G}^{-1})$
- D-optimal design corresponds to $h(\boldsymbol{G}) = det(\boldsymbol{G}^{-1}) = det(\boldsymbol{G})^{-1}$
- c-optimal design correspond to $h(\mathbf{G}) = \mathbf{c}^T \mathbf{G}^{-1} \mathbf{c}$ for some vector $\mathbf{c} \in \mathbb{R}^m$.

Least-squares approximation with i.i.d. sampling

Assume that the x_i are i.i.d. samples from a distribution $d\nu(x) = w(x)^{-1}d\mu(x)$ for some weight function w, and $w_i = w(x_i)$. We have

$$\boldsymbol{G} = rac{1}{n}\sum_{i=1}^{n} \boldsymbol{A}_{i}, \quad \boldsymbol{A}_{i} = w(x_{i})\boldsymbol{\varphi}(x_{i})\boldsymbol{\varphi}(x_{i})^{T},$$

where the A_i are i.i.d. rank-one matrices with expectation

$$\mathbb{E}(\boldsymbol{A}_{i}) = \mathbb{E}_{x \sim \nu}(w(x)\varphi(x)\varphi(x)^{T}) = \mathbb{E}_{x \sim \mu}(\varphi(x)\varphi(x)^{T}) = \boldsymbol{I}$$

and spectral norm

$$\|\boldsymbol{A}_i\| = w(x_i)\|\boldsymbol{\varphi}(x_i)\|_2^2 \leq K_{w,m}$$

with

$$K_{w,m} = \sup_{x \in \mathcal{X}} w(x) \| \varphi(x) \|_2^2.$$

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and spectral norm

$$\|\boldsymbol{A}_i\| = w(x_i)\|\varphi(x_i)\|_2^2 \leq K_{w,m},$$

with

$$\mathcal{K}_{w,m} = \sup_{x \in \mathcal{X}} w(x) \| \varphi(x) \|_2^2.$$

Based on matrix Chernoff concentration inequality, it can be shown that for any $0<\delta<1,$

$$\mathbb{P}(\lambda_{\textit{max}}(\boldsymbol{G}) > 1 + \delta) \land \mathbb{P}(\lambda_{\textit{min}}(\boldsymbol{G}) < 1 - \delta) \leq m \exp(-\frac{n\delta^2}{\mathcal{K}_{w,m}})$$

and

$$\mathbb{P}(\|\boldsymbol{G} - \boldsymbol{I}\| > \delta) = \mathbb{P}(\lambda_{max}(\boldsymbol{G}) > 1 + \delta \text{ or } \lambda_{min}(\boldsymbol{G}) < 1 - \delta) \leq 2m \exp(-\frac{n\delta^2}{K_{w,m}})$$

Least-squares approximation with i.i.d. sampling

We obtain that

$$\mathbb{P}(\|\boldsymbol{G}-\boldsymbol{I}\| > \delta) \leq \eta$$

provided that

$$n \geq K_{w,m} \delta^{-2} \log(2m\eta^{-1}).$$

We note that

$$\mathcal{K}_{w,m} = \sup_{x \in \mathcal{X}} w(x) \|\varphi(x)\|_2^2 \geq \mathbb{E}_{x \sim \nu}(w(x)\|\varphi(x)\|_2^2) = \mathbb{E}_{x \sim \mu}(\sum_{j=1}^m \varphi_j(x)^2)$$

so that

 $K_{w,m} \ge m$

For classical least-squares, w = 1 ($\nu = \mu$).

- For V_m piecewise constant functions on a uniform partition of (0, 1) and μ the uniform measure, $K_{1,m} = m$.
- For V_m trigonometric polynomials of degree (m-1)/2 on $(0, 2\pi)$ and μ the uniform measure, $\kappa_{1,m} = m$.
- For polynomial spaces $V_m = \mathbb{P}_{m-1}$ and μ the uniform measure, $|K_{1,m} = m^2|$.
- For polynomial spaces $V_m = \mathbb{P}_{m-1}$ and μ the gaussian measure on \mathbb{R} , $|K_{1,m} = \infty|$.

With i.i.d. sampling, an optimal sampling measure ν_m is given by $d\nu_m(x) = w_m(x)^{-1} d\mu(x)$ with density

$$w_m(x)^{-1} = \frac{1}{m} \sum_{j=1}^m \varphi_j(x)^2$$

that minimizes $K_{w,m}$ over all densities, and yields

$$K_{w_m,m} = m$$

For polynomial approximation, $\sum_{j=1}^{m} \varphi_j(x)^2$ is the inverse of the Christoffel function.

Under the condition

$$n \ge m\delta^{-2}\log(2m\eta^{-1})$$

we have

$$\mathbb{P}(\|\boldsymbol{G}-\boldsymbol{I}\| > \delta) \le \eta$$

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- For polynomial spaces $V_m = \mathbb{P}_{m-1}$ and μ the uniform measure on (-1, 1)



Figure: Polynomials and uniform measure: density of ν_m

• For polynomial spaces $V_m = \mathbb{P}_{m-1}$ and μ the gaussian measure on \mathbb{R}



Figure: Polynomials and Gaussian measure: density of ν_m

• For *d*-variate polynomials,

$$V_m = \mathbb{P}_{\Lambda} := span\{x^{\alpha} = x_1^{\alpha_1} \dots x_d^{\alpha_d} : \nu \in \Lambda \subset \mathbb{N}^d\}$$

$$\begin{split} &\Lambda = \Lambda_{1,p} := \{ \alpha : \|\alpha\|_1 \leq p \} \text{ corresponds to polynomials with total degree} \leq p. \\ &\Lambda = \Lambda_{\infty,p} := \{ \alpha : \|\alpha\|_\infty \leq p \} \text{ corresponds to polynomials with partial degree} \leq p. \end{split}$$



(a) A_{1,4}

(b) Λ_{∞,4}

Figure: Polynomials and uniform measure on $[-1, 1]^2$: density w_m for polynomials with total (left) or partial (right) degree less than 4.

We have to sample from the optimal measure

$$d\nu_m = w_m^{-1} d\mu, \quad w_m(x)^{-1} = \frac{1}{m} \sum_{j=1}^m \varphi_j(x)^2$$

Standard sampling technique can be used: inverse transform, rejection, Markov Chain Monte-Carlo...

However, for general spaces V_m , sampling may be a non trivial task.

We observe that ν_m is a mixture of measures

$$d\nu^{(j)}(x) = \varphi_j(x)^2 d\mu(x)$$

with equal weights 1/m. We can first sample j uniformly at random in $\{1, \ldots, m\}$ and then sample from $\nu^{(j)}$.

- In adaptive approximation, we construct approximations from a sequence of spaces $(V_m)_{m\geq 1}$.
- To each space V_m is associated a specific optimal sampling measure $\nu_m = w_m^{-1} \mu$.

When functions evaluations are costly, we would like to exploit samples generated at previous iterations.

Recycling samples for adaptive approximation: hierarchical spaces

Consider the adaptive approximation in a sequence of nested spaces

 $V_1 \subset \ldots \subset V_m \subset V_{m+1} \subset \ldots$

Let $(\varphi_j)_{j\geq 1}$ be such that $V_m = span\{\varphi_1, \ldots, \varphi_m\}$. Then

$$V_{m+1} = V_m \oplus span\{\varphi_{m+1}\}$$

and the optimal sampling measure ν_{m+1} associated to V_{m+1} is such that

$$d
u_{m+1}(x) = rac{1}{m+1} \sum_{j=1}^{m+1} \varphi_j(x)^2 d\mu(x) = rac{m}{m+1} d
u_m(x) + rac{1}{m+1} \varphi_{m+1}^2 d\mu(x)$$

that corresponds to a mixture between ν_m and $\varphi_{m+1}^2 \mu$, with respective weights $\frac{m}{m+1}$ and $\frac{1}{m+1}$.

Recycling samples for adaptive approximation: hierarchical spaces

Consider the adaptive approximation in a sequence of nested spaces

 $V_1 \subset \ldots \subset V_m \subset V_{m+1} \subset \ldots$

Let $(\varphi_j)_{j\geq 1}$ be such that $V_m = span\{\varphi_1, \ldots, \varphi_m\}$. Then

$$V_{m+1} = V_m \oplus span\{\varphi_{m+1}\}$$

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that corresponds to a mixture between ν_m and $\varphi_{m+1}^2 \mu$, with respective weights $\frac{m}{m+1}$ and $\frac{1}{m+1}$.

To sample the mixture, draw a Bernoulli variable $B(\frac{1}{m+1})$. If 1 is obtained, generate a new sample from $\varphi_{m+1}^2\mu$. If 0 is obtain, then either pick without replacement a sample from previously generated samples from ν_m , or generate a new sample from ν_m .

Different strategies can be found in [Arras et al 2019, Migliorati 2019].

Optimal weighted least-squares: error analysis

Let $f_m = P_{V_m} f$ be the orthogonal projection of f onto V_m w.r.t. the norm $\|\cdot\|$, that is the element of best approximation of f in V_m . We have

$$\begin{split} \|f - \hat{f}_m\|^2 &\leq \|f - f_m\|^2 + \|f_m - \hat{f}_m\|^2 \\ &\leq \|f - f_m\|^2 + \lambda_{\min}(\boldsymbol{G})^{-1} \|f_m - \hat{f}_m\|_n^2 \\ &\leq \|f - f_m\|^2 + \lambda_{\min}(\boldsymbol{G})^{-1} \|f_m - f\|_n^2 \end{split}$$

where we have used the fact that \hat{f}_m is the orthogonal projection of f onto V_m w.r.t. the semi-norm $\|\cdot\|_n$.

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where we have used the fact that \hat{f}_m is the orthogonal projection of f onto V_m w.r.t. the semi-norm $\|\cdot\|_n$.

If $\|\boldsymbol{G} - \boldsymbol{I}\| \leq \delta$, then $\lambda_{\min}(\boldsymbol{G}) \geq 1 - \delta$ and $\|\boldsymbol{f} - \hat{f}_m\|^2 \leq \|\boldsymbol{f} - f_m\|^2 + (1 - \delta)^{-1} \|\boldsymbol{f} - f_m\|_n^2$

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In order to control of the approximation when $\|\boldsymbol{G} - \boldsymbol{I}\| > \delta$, different alternatives:

- assuming $||f||_{\infty} \leq \tau$, define a truncated estimator $\hat{f}_m^{\tau} = T_{\tau} \circ \hat{f}_m$ with $T_{\tau}(t) = sign(t) \min\{|t|, \tau\},$
- define a conditional estimator $\hat{f}_m^C = f_m$ if $\|\boldsymbol{G} \boldsymbol{I}\| \leq \delta$ or 0 if $\|\boldsymbol{G} \boldsymbol{I}\| > \delta$,
- condition the samples to guarantee stability $\|\boldsymbol{G} \boldsymbol{I}\|$.

Optimal weighted least-squares with conditioning

Assume that $\mathbf{x} = (x_1, \dots, x_n)$ are drawn from $\nu^{\otimes n}$ conditioned to satisfy the event $S = \{ \| \mathbf{G}(\mathbf{x}) - \mathbf{I} \| \leq \delta \}$. This can be obtained by sampling \mathbf{x} from $\nu^{\otimes n}$ until S is satisfied (rejection).

Under the condition

$$n \ge m\delta^{-2}\log(2m\eta^{-1}) \tag{3}$$

we have

$$\mathbb{P}(S) \geq 1 - \eta$$

For $\eta < 1$, the random number N of samples from $\nu^{\otimes n}$ generated before acceptation follows a geometric distribution with parameter $\mathbb{P}(S)$, is almost surely finite, and with expectation $\mathbb{E}(N) = \mathbb{P}(S)^{-1} \leq (1 - \eta)^{-1}$.

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The least-squares estimator satisfies

$$\begin{split} \mathbb{E}(\|f - \hat{f}_m\|^2) &\leq \|f - f_m\|^2 + (1 - \delta)^{-1} \mathbb{E}(\|f - f_m\|_n^2) \\ &\leq \|f - f_m\|^2 + (1 - \delta)^{-1} (1 - \eta)^{-1} \mathbb{E}_{\mathbf{x} \sim \nu^{\otimes n}} (\|f - f_m\|_n^2) \\ &= (1 + (1 - \delta)^{-1} (1 - \eta)^{-1}) \|f - f_m\|^2 \end{split}$$

Optimal weighted least-squares with conditioning

Therefore, we deduce a quasi-optimality in expectation

$$\mathbb{E}(\|f-\hat{f}_m\|^2)^{1/2} \leq C \inf_{v \in V_m} \|f-v\|,$$

with $C = (1 + (1 - \delta)^{-1}(1 - \eta)^{-1})^{1/2}$.
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For a compact set K of functions in L^2_{μ} , using the previous result with an optimal subspace V_m of dimension m such that

$$\inf_{v\in V_m}\|f-v\|=d_m(K)_{L^2_{\mu}},$$

we deduce that for $n \gtrsim cm \log(m)$, for some universal constant c, there exists a distribution over \mathcal{X}^n and a linear recovery map A such that

$$\mathbb{E}(\|f - A(f(x_1), \dots, f(x_n))\|^2)^{1/2} \le Cd_m(K)_{L^2_{\mu}}$$

which proves

$$\rho_{cm\log(m)}^{rand}(K)_{L^2_{\mu}} \leq Cd_m(K)_{L^2_{\mu}}$$

Optimal weighted least-squares with conditioning and subsampling

By conditioning, we obtain $n \ge cm \log(m)$ samples that guarantee almost surely

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However, the number of samples n may be large compared to m, and a fundamental question is whether the log(m) factor can be removed.

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In [Haberstich, Nouy and Perrin 2022], a subsampling approach is proposed, which consists in removing samples until the stability condition is violated. More precisely, for $I \subset \{1, ..., n\}$, we let $\boldsymbol{G}_{I} = \frac{1}{|I|} \sum_{i \in I} \boldsymbol{A}_{i}$. Starting from the set $I = \{1, ..., n\}$, we successively remove from the current set I an index i such that

$$i \in \min_{j \in I} \|\boldsymbol{G}_{I \setminus \{j\}} - \boldsymbol{I}\|$$

If $\|\boldsymbol{G}_{I\setminus\{i\}} - \boldsymbol{I}\| > \delta$, we stop and return *I*. Otherwise, we continue removing samples.

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However, the number of samples n may be large compared to m, and a fundamental question is whether the log(m) factor can be removed.

We observe in many applications that the algorithm returns a number of samples close to or even equal to m, without any theoretical guaranty.

Optimal weighted least-squares with conditioning and subsampling

In [Cohen and Dolbeault 2021], it is proposed a subsampling strategy, based on successive random partitioning of the set of samples, which yields a number of samples in O(m) while preserving stability.¹ Note that

$$oldsymbol{G} = \sum_{i=1}^n oldsymbol{a}_i oldsymbol{a}_i^T \quad ext{with} \quad oldsymbol{a}_i = \sqrt{rac{w(x_i)}{n}} oldsymbol{arphi}(x_i) \in \mathbb{R}^m.$$

We have

$$(1-\delta)\boldsymbol{I} \preccurlyeq \sum_{i=1}^n \boldsymbol{a}_i \boldsymbol{a}_i^{\mathsf{T}} \preccurlyeq (1+\delta)\boldsymbol{I} \quad \text{and} \quad \|\boldsymbol{a}_i\|_2^2 = m/n.$$

¹It relies on results from [Markus, Spielman and Srivastava 2015][Nitzan, Olevskii and Olevskii 2016] that provide a solution to the Kadinson-Singer problem.

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A procedure is introduced which provides a partition of $\{1, \ldots, n\}$ into sets J_1, \ldots, J_{2^L} with cardinal $|J_k| \leq cm$, and such that for all $1 \leq k \leq 2^L$

$$c_0 \mathbf{I} \preccurlyeq \frac{n}{m} \sum_{i \in J_k} \mathbf{a}_i \mathbf{a}_i^T \preccurlyeq C_0 \mathbf{I}$$

with universal constants c_0 and C_0 . Then pick k at random in $\{1, \ldots, 2^L\}$ with probability $p_k = |J_k|/m$.

¹It relies on results from [Markus, Spielman and Srivastava 2015][Nitzan, Olevskii and Olevskii 2016] that provide a solution to the Kadinson-Singer problem.

This proves that

$$ho_{cm}^{rand}(K)_{L^2_{\mu}} \leq Cd_m(K)_{L^2_{\mu}}$$

for some universal constants c and C.

However, the subsampling strategy is not computationally feasible.

Other subsampling strategy have been proposed in [Bartel et al 2022], with theoretical guarantees and feasible implementations.

Note that the samples x_1, \ldots, x_n obtained by conditioning (and possibly subsampling) are no more independent and follows a distribution which is not explicit.

In adaptive setting, we can no more recycle samples using mixture sampling.

An alternative recycling method has been proposed in [Haberstich 2020].

We would like to obtain quasi-optimality guarantees with high probability, or even almost surely, for the approximation of functions from a space X continuously embedded in L^2_{μ} , that is such that $||f|| \leq C_X ||f||_X$ for all $f \in X$.

For that, the sampling should depend on both X and V_m .

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We can consider a mixture between the optimal distribution $w_m^{-1}d\mu$ and a distribution $hd\mu$, with density

$$w(x)^{-1} = \frac{1}{2}w_m(x)^{-1} + \frac{1}{2}h(x)$$

where h is a related to X.

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The empirical Gram matrix **G** remains an unbiased estimator of **I** and

$$K_{w,m} = \sup_{x \in \mathcal{X}} w(x) \| \varphi(x) \|_2^2 \le 2K_{w_m,m} = 2m$$

Therefore, only a factor 2 is lost in the number of samples required to ensure $\|\boldsymbol{G} - \boldsymbol{I}\| \le \delta$ with nonzero probability. By conditioning we obtain almost surely the error bound

$$\|f - \hat{f}_m\| \leq \|f - g\| + (1 - \delta)^{-1/2} \|f - g\|_n \quad orall g \in V_m.$$

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$$\|f - \hat{f}_m\| \le \|f - g\| + (1 - \delta)^{-1/2} \|f - g\|_n \quad \forall g \in V_m.$$

If the function h is chosen such that for all $f \in X$, $||f||_n \leq C ||f||_X$, we obtain

$$\|f - \hat{f}_m\| \le (C_X + (1 - \delta)^{-1/2}C) \inf_{g \in V_m} \|f - g\|_X$$

For $X = L^{\infty}_{\mu}(\mathcal{X})$ equipped with its natural norm $\|\cdot\|_{\infty}$, we can take

h(x) = 1

so that $w(x)^{-1} \geq 1/2$. For all $f \in X$, we then have $\|f\| \leq \|f\|_\infty$ and

$$\|f\|_n^2 = \frac{1}{n} \sum_{i=1}^n w(x_i) f(x_i)^2 \le \frac{2}{n} \sum_{i=1}^n f(x_i)^2 \le 2 \|f\|_{\infty}^2$$

This yields

$$\|f - \hat{f}_m\| \le (1 + (1 - \delta)^{-1/2} \sqrt{2}) \inf_{g \in V_m} \|f - g\|_{\infty}$$

Consider for X a RKHS with a kernel k in $L^2_{\mu\otimes\mu}(\mathcal{X}\times\mathcal{X})$ that admits a decomposition

$$k(x,y) = \sum_{i \ge 1} \lambda_i \psi_i(x) \psi_i(y)$$

where the ψ_i form an orthonormal basis of $L^2_{\mu}(\mathcal{X})$ and where $(\lambda_i)_{i\geq 1}$ is a decreasing sequence of strictly positive numbers such that

$$\sum_{i\geq 1}\lambda_i^2 = \|k\|_{L^2}^2 < \infty.$$

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The (ψ_i, λ_i) are the eigenpairs of the Hilbert-Schmidt integral operator T_k with kernel k. The norm on X is given by

$$\|f\|_X^2 = \sum_{i\geq 1} (f, \psi_i)_{L^2_{\mu}}^2 / \lambda_i,$$

and

$$\|f\|^{2} = \sum_{i \ge 1} (f, \psi_{i})^{2}_{L^{2}_{\mu}} = \sum_{i \ge 1} \lambda_{i} (f, \psi_{i})^{2}_{L^{2}_{\mu}} / \lambda_{i} \le \lambda_{1} \|f\|^{2}_{X}.$$

Therefore, X is continuously embedded in L^2_{μ} with embedding constant $C_X = \lambda_1^{1/2}$.

We further assume (up to a rescaling) that

$$\sum_{i\geq 1}\lambda_i=\int k(x,x)d\mu(x)=1<\infty$$

that is T_k is nuclear (trace class) with unit nuclear norm.

Therefore, k(x, x) defines a density and we can take

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We have $w(x)^{-1} \ge k(x,x)/2$, so that

$$\|f\|_n^2 \leq \frac{2}{n} \sum_{i=1}^n k(x_i, x_i)^{-1} f(x_i)^2 = \frac{2}{n} \sum_{i=1}^n k(x_i, x_i)^{-1} (k(x_i, \cdot), f)_X^2 \leq 2 \|f\|_X^2$$

We finally deduce

$$\|f - \hat{f}_m\| \le (\lambda_1 + (1 - \delta)^{-1/2} \sqrt{2}) \inf_{g \in V_m} \|f - g\|_X$$

Using subsampling techniques from [Cohen and Dolbeault 2021], we then prove that for $X = L^{\infty}$ or X a RKHS associated with a trace class operator, there exists a set of $n \leq cm$ points and a linear algorithm such that for all $f \in X$, the produced approximation $\hat{f}_m = A(f(x_1), \ldots, f(x_n))$ is such that

 $\|f-\hat{f}_m\| \leq CE(f;V_m)_X$

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Consider a compact set $K \subset X$ and an optimal approximating subspace V_m in the sense that $\sup_{f \in K} E(f; V_m)_X = d_m(K)_X$. We then have proven that

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For K the unit ball of a RKHS (with the trace class assumption), a refined analysis (see [1]) yields

$$\rho_{cm}(K)_{L^2} \leq \sqrt{\frac{1}{m}\sum_{k\geq m}d_k(K)_{L^2}^2}$$

for some universal constant c, which is known as a sharp bound.

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$$\rho_{cm}(K)_{L^2} \leq \sqrt{\frac{1}{m}\sum_{k\geq m}d_k(K)_{L^2}^2}$$

for some universal constant c, which is known as a sharp bound. For a larger class of spaces including the space of bounded functions equipped with the supremum norm, they show

$$\rho_{cm}(\mathcal{K})_{L^2} \leq \left(\frac{1}{m}\sum_{k \geq m} d_k(\mathcal{K})_{L^2}^p\right)^{1/p} \quad \text{for any } 0$$

- 1 Manifold approximation
- 2 Linear approximation from point evaluations
- 3 Tensor networks approximation with point evaluations

For the approximation of tensors (or functions) using tensor networks, different contexts depending on the available information:

- all entries of the tensor,
- equations satisfied by the tensor,
- some entries, either arbitrary or structured,
- more general functionals of the tensor.

- tensap. A Python package for the approximation of functions and tensors. (link to GitHub page).
- ApproximationToolbox. An object-oriented MATLAB toolbox for the approximation of functions and tensors. (link to GitHub page).

For the approximation of a multivariate function with tree tensor networks using point evaluations, different strategies have been proposed, either based on cross approximation [Oseledets'10, Ballani'13] or principal component analysis [Nouy'19, Haberstich'21].

These methods rely on structured evaluations

$$u(x^i_{\alpha}, x^j_{\alpha^c})$$

where x_{α}^{i} are samples of the variables x_{α} , and $x_{\alpha c}^{j}$ samples of the variables $x_{\alpha c}$.

Consider a multivariate function $u \in L^2_{\mu}(\mathcal{X})$ where $\mathcal{X} = \mathcal{X}_1 \times \ldots \times \mathcal{X}_d$ is equipped with a probability measure $\mu = \mu_1 \otimes \ldots \otimes \mu_d$. Let $X = (X_1, \ldots, X_d)$ be a random vector with distribution μ , such that the L^2_{μ} -norm is given by

$$||u||^{2} = \int u(x)^{2} d\mu(x) = \mathbb{E}(u(X)^{2}).$$

For each a subset of variables α and its complementary subset $\alpha^c = D \setminus \alpha$, u is identified with a bivariate function defined on $\mathcal{X}_{\alpha} \times \mathcal{X}_{\alpha^c}$ which admits a singular value decomposition

$$u(x_{\alpha}, x_{\alpha^{c}}) = \sum_{k=1}^{\operatorname{rank}_{\alpha}(u)} \sigma_{k}^{\alpha} v_{k}^{\alpha}(x_{\alpha}) v_{k}^{\alpha^{c}}(x_{\alpha^{c}})$$

Learning from principal component analysis

The subspace of α -principal components

$$U_{lpha} = span\{v_1^{lpha}, \dots, v_{r_{lpha}}^{lpha}\}$$

is such that

$$u_{\mathbf{r}_{\alpha}}(\cdot, x_{\alpha^{c}}) = \mathcal{P}_{U_{\alpha}} u(\cdot, x_{\alpha^{c}})$$

It is solution of

$$\min_{\dim(U_{\alpha})=r_{\alpha}} \|u-\mathcal{P}_{U_{\alpha}}u\|^{2}$$

that is for $\|\cdot\|$ the $L^2_\mu(\mathcal{X})$ -norm,

$$\min_{\dim(U_{\alpha})=r_{\alpha}} \mathbb{E}\left(\|u(\cdot, X_{\alpha^{c}}) - \mathcal{P}_{U_{\alpha}} u(\cdot, X_{\alpha^{c}})\|_{L^{2}_{\mu_{\alpha}}(\mathcal{X}_{\alpha})}^{2} \right)$$

where u is seen as a function-valued random variable

$$u(\cdot, X_{\alpha^c}) \in L^2_{\mu_{\alpha}}(\mathcal{X}_{\alpha}).$$

 U_{α} is the optimal *m*-dimensional space for the approximation of the manifold $\{u(\cdot, x_{\alpha^c}) : x_{\alpha^c} \in \mathcal{X}_{\alpha^c}\}$ in mean-squared error.

For tree tensor networks

$$\mathcal{T}_r^{\mathsf{T}}(\mathsf{V}) = \{\mathsf{v} \in \mathsf{V} : \mathsf{rank}_{\alpha}(\mathsf{v}) \leq \mathsf{r}_{\alpha}, \alpha \in \mathsf{T}\},\$$

where T is a dimension partition tree over $D = \{1, ..., d\}$, different variants of higher order singular value decomposition (also called hierarchical singular value decomposition) can be defined from singular value decompositions of bivariate functions.



Higher-order principal component analysis for Tucker format

Tucker format corresponds a trivial tree with d + 1 nodes (the root and the d leaves).



For each leaf $\nu \in \{1, ..., d\}$, we determine a $\{\nu\}$ -principal subspace $U_{r_{\nu}}^{\nu}$ of dimension r_{ν} (a space of functions of the variable x_{ν}).

Then, we obtain an approximation in Tucker format (with ranks r_1, \ldots, r_d) by a projection of the function u onto the linear tensor product space

 $U_1 \otimes \ldots \otimes U_d$

Leaves to root strategy for general tree tensor networks

For each leaf node α , let $U_{r_{\alpha}}^{\alpha}$ be the r_{α} -dimensional α -principal subspace of u.



For each interior node $\alpha \in T \setminus \{D\}$ with children $S(\alpha)$, define a tensor space

$$V_{\alpha} = \bigotimes_{\beta \in \mathcal{S}(\alpha)} U_{r_{\beta}}^{\beta}$$

and let $U_{r_{\alpha}}^{\alpha} \subset V_{\alpha}$ be the r_{α} -dimensional α -principal subspace of the function u_{α} defined by

$$u_{\alpha}(\cdot, x_{\alpha^{c}}) = \mathcal{P}_{V_{\alpha}} u(\cdot, x_{\alpha^{c}})$$



Finally define an approximation u_r as a projection of u onto the tensor space $V_D = \bigotimes_{\alpha \in S(D)} U_{\alpha}$.

We can prove that the resulting approximation u_r is a tree tensor network with ranks r_{α} , $\alpha \in \mathcal{T}$.



Leaves to root truncation scheme for tree-based tensor formats

Provided we use orthogonal projections, the obtained approximation u_r is such that

$$\|u-u_r\|^2 \leq \sum_{\alpha \in T \setminus D} \min_{\operatorname{rank}_{\alpha}(v) \leq r_{\alpha}} \|u-v\|^2 = \sum_{\alpha \in T \setminus D} \sum_{k_{\alpha} > r_{\alpha}} (\sigma_{k_{\alpha}}^{\alpha})^2,$$

from which we deduce that u_r is a quasi-optimal approximation of u in $\mathcal{T}_r^{\mathsf{T}}$ such that

$$\|u-u_r\|\leq C(T)\min_{v\in\mathcal{T}_r^{\mathcal{T}}}\|u-v\|,$$

where $C(T) = \sqrt{\#T - 1}$ is the square root of the number of projections applied to the tensor. The number of nodes of a dimension partition tree T being bounded by 2d - 1,

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Also, if we select the ranks $(r_{\alpha})_{\alpha \in T \setminus D}$ such that for all α

$$\sum_{k_{\alpha}>r_{\alpha}}(\sigma_{k_{\alpha}}^{\alpha})^{2}\leq \frac{\epsilon^{2}}{C(T)^{2}}\sum_{k_{\alpha}\geq 1}(\sigma_{k_{\alpha}}^{\alpha})^{2}=\frac{\epsilon^{2}}{C(T)^{2}}\|u\|^{2},$$

we finally obtain an approximation u_r with relative precision ϵ ,

$$\|u-u_r\|\leq \epsilon\|u\|.$$

Leaves to root truncation scheme for tree-based tensor formats

Given a finite dimensional tensor space $V = V_1 \otimes \ldots \otimes V_d$, an approximation in the tensor format $\mathcal{T}_r^T(V)$ can be obtained by modifying the procedure for the leaves.

For each leaf node α , $U^{\alpha}_{r_{\alpha}}$ is defined as a α -principal subspace of $u_{\alpha} = \mathcal{P}_{V_{\alpha}}u$.

Theorem (Fixed rank)

For a given T-rank, we obtain an approximation $u_r \in \mathcal{T}_r^{\mathsf{T}}(\mathsf{V})$ such that

$$\|u_r - u\|^2 \leq C(T)^2 \min_{v \in \mathcal{T}_r^T} \|v - u\|^2 + \sum_{l \in aves \ \alpha} \|u - \mathcal{P}_{V_\alpha} u\|^2$$

Theorem (Fixed precision)

For a desired precision ϵ , if the α -ranks are determined such that

$$\|\mathcal{P}_{U_{r_{\alpha}}^{\alpha}}u_{\alpha}-u_{\alpha}\|\leq \frac{\epsilon}{C(T)}\|u_{\alpha}\|,$$

we obtain an approximation u_r such that

$$\|u_r - u\|^2 \leq \epsilon^2 \|u\|^2 + \sum_{\text{leaves } \alpha} \|u - \mathcal{P}_{V_{\alpha}} u\|^2.$$

For a feasible algorithm using samples:

- Replacement of orthogonal projections by sampled-based projections, based on interpolation [Nouy 2019] or optimal least-squares projections [Haberstich 2021].
- Statistical estimation of principal subspaces U_{α} by empirical PCA, using samples $u(\cdot, x_{\alpha^c}^j)$

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The estimation of principal subspaces requires the evaluation of u on a structured set of points

$$\{(\mathbf{x}_{\alpha}^{i}, \mathbf{x}_{\alpha^{c}}^{j}) : 1 \leq i \leq M_{\alpha}, 1 \leq j \leq N_{\alpha}\}$$

where N_{α} is the number of samples $x_{\alpha^c}^j$ used for the estimation of U_{α} by empirical PCA, and M_{α} is the number of points x_{α}^j used for the projections onto the space V_{α} .

The sampling strategy is adaptive to the function.
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The sampling strategy is adaptive to the function.

Some guarantees can be obtained under additional assumptions on the function to approximate [Haberstich 2021].

But yet not guaranty of quasi-optimality in a general setting.

Development of near optimal learning algorithms.

- Theory well established for least-squares approximation in linear spaces
- Mainly an open problem for linear approximation in other spaces than L^2
- Only partial results on optimal sampling for least-squares approximation with tensor networks, and mainly open problem for neural networks.
- Optimal sampling for manifold approximation ? Some results for linear manifold approximation (PCA, Reduced basis), but mainly an open problem for general nonlinear approximation of manifolds.

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