

# Fast Computation of RBF-PU Interpolants

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Joint work with

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# Introduction

Main target: interpolate (large) scattered data sets using efficient and accurate algorithms.

- [Allasia, Besenghi, Cavoretto, De Rossi (AMC, 2011)]: new 2D efficient implementation of the modified Shepard's algorithm using MLSAs and RBFs as nodal functions  $\Rightarrow$  novelty of the partition of the domain in *strips* for constructing a **strip-based searching procedure**.
- [Cavoretto, De Rossi (JCAM, 2010)]: extension of the previous idea to the spherical case using ZBFs (instead of RBFs)  $\Rightarrow$  partition of the domain in *spherical zones*  $\Rightarrow$  **spherical zone searching procedure**.
- [Cavoretto, De Rossi (AML, 2012)]: the spherical zone algorithm has been generalized using the **partition of unity method**.
- [Cavoretto, De Rossi (CAMWA, 2014)]: new 2D partition of unity algorithm which is based on the partition of the domain in *cells* (squares) using a double structure of crossed-strips  $\Rightarrow$  **cell-based searching procedure**.
- [Cavoretto, De Rossi (Submitted, 2014)]: extension of the 2D algorithm to the 3D case partitioning the domain in *cells* (cubes).
- [Cavoretto, De Rossi, Perracchione (Submitted, 2015)]: generalization for generic 2D and 3D domains using a new **block-based searching procedure**.



## BLOCK-BASED INTERPOLATION ALGORITHMS FOR GENERIC DOMAINS

# Scattered data interpolation problem

Since many applications either arise from a *function approximation problem*, or include *function interpolation* as a fundamental component, we consider the **scattered data interpolation problem** which requires the use of meshfree methods and algorithms.



- $\mathcal{X}_n = \{\mathbf{x}_i, i = 1, \dots, n\}$ , set of *data points or nodes*;
- $\mathcal{F}_n = \{f_i = f(\mathbf{x}_i), i = 1, \dots, n\}$ , set of *data values or function values*.

## Definition

Given a set  $\mathcal{X}_n = \{\mathbf{x}_i, i = 1, \dots, n\}$  of  $n$  distinct data points on the domain  $\Omega \subset \mathbb{R}^N$  and a set  $\mathcal{F}_n = \{f_i, i = 1, \dots, n\}$  of the corresponding data values of an (unknown) continuous function  $f : \Omega \rightarrow \mathbb{R}$ , the **interpolation problem** is to find a continuous function  $\mathcal{I} : \Omega \rightarrow \mathbb{R}$  which satisfies the interpolation conditions, i.e.

$$\mathcal{I}(\mathbf{x}_i) = f_i, \quad i = 1, \dots, n.$$

References: [Buhmann (2003), Fasshauer (2007), Wendland (2005)]

# RBF interpolation

- The standard *RBF interpolation problem* is to find an interpolant  $R : \Omega \rightarrow \mathbb{R}$  of the form

$$R(\mathbf{x}) = \sum_{i=1}^n c_i \phi(\|\mathbf{x} - \mathbf{x}_i\|_2), \quad \mathbf{x} \in \Omega, \quad (1)$$

where  $\|\cdot\|_2$  is the Euclidean norm, and  $\phi : [0, \infty) \rightarrow \mathbb{R}$  is a RBF. The coefficients  $\{c_i\}_{i=1}^n$  are determined by enforcing the interpolation conditions

$$R(\mathbf{x}_i) = f_i, \quad i = 1, \dots, n. \quad (2)$$

- Imposing the conditions (2) leads to a *symmetric* linear system of equations

$$\Phi \mathbf{c} = \mathbf{f}, \quad (3)$$

where  $\Phi_{ki} = \phi(\|\mathbf{x}_k - \mathbf{x}_i\|_2)$ ,  $k, i = 1, \dots, n$ ,  $\mathbf{c} = [c_1, \dots, c_n]^T$ , and  $\mathbf{f} = [f_1, \dots, f_n]^T$ . When  $\mathbf{c}$  is found by solving the *system* (3), we can evaluate the RBF interpolant at a point  $\mathbf{x}$  as

$$R(\mathbf{x}) = \bar{\phi}(\mathbf{x}) \mathbf{c},$$

where  $\bar{\phi}(\mathbf{x}) = [\phi(\|\mathbf{x} - \mathbf{x}_1\|_2), \dots, \phi(\|\mathbf{x} - \mathbf{x}_n\|_2)]$ .

# Examples of standard RBFs

RBF	$\phi(r)$
Gaussian $C^\infty$ (G)	$e^{-\alpha^2 r^2}, \alpha > 0$
Inverse MultiQuadric $C^\infty$ (IMQ)	$(1 + \gamma^2 r^2)^{-1/2}, \gamma > 0$
MultiQuadric $C^\infty$ (MQ)	$(1 + \gamma^2 r^2)^{1/2}, \gamma > 0$
Matérn $C^4$ (M4)	$e^{-\epsilon r}(\epsilon^2 r^2 + 3\epsilon r + 3), \epsilon > 0$
Wendland $C^4$ (W4)	$(1 - cr)_+^6 (35c^2 r^2 + 18cr + 3), c > 0$
Wendland $C^2$ (W2)	$(1 - cr)_+^4 (4cr + 1), c > 0$

# Partition of unity interpolation

## Definition

Given a partition of the open and bounded domain  $\Omega \subseteq \mathbb{R}^N$  into  $d$  subdomains  $\Omega_j$  such that  $\Omega \subseteq \bigcup_{j=1}^d \Omega_j$  with some mild overlap among the subdomains, the *partition of unity method* is obtained selecting a family of compactly supported, non-negative, continuous functions  $W_j$  with  $\text{supp}(W_j) \subseteq \Omega_j$  such that

$$\sum_{j=1}^d W_j(\mathbf{x}) = 1, \quad \mathbf{x} \in \Omega.$$

The *global approximant*  $\mathcal{I} : \Omega \rightarrow \mathbb{R}$  takes the form

$$\mathcal{I}(\mathbf{x}) = \sum_{j=1}^d R_j(\mathbf{x}) W_j(\mathbf{x}), \quad \mathbf{x} \in \Omega, \quad (4)$$

where  $R_j$  are local approximants satisfying the interpolation conditions at nodes  $\mathbf{x}_i$ ,  $i = 1, \dots, n$ .

- *Shepard's weights:*

$$W_j = \bar{W}_j / \sum_k \bar{W}_k.$$

# Local RBF interpolants

- Here  $R_j : \Omega_j \rightarrow \mathbb{R}$  defines a *RBF interpolant* of the form

$$R_j(\mathbf{x}) = \sum_{i=1}^{\bar{n}_j} c_i^{(j)} \phi(||\mathbf{x} - \mathbf{x}_i^{(j)}||_2),$$

where  $\phi : [0, \infty) \rightarrow \mathbb{R}$  represents a *RBF*,  $||\cdot||_2$  denotes the Euclidean norm, and  $\bar{n}_j$  indicates the number of data points in  $\Omega_j$ , i.e. the points  $\mathbf{x}_i^{(j)} \in \mathcal{X}_j = \mathcal{X}_n \cap \Omega_j$ .

- Furthermore,  $R_j$  satisfies the *interpolation conditions*

$$R_j(\mathbf{x}_i^{(j)}) = f_i^{(j)}, \quad i = 1, \dots, \bar{n}_j. \quad (5)$$

Note: if the *local approximants* satisfy the interpolation conditions (5), then the global approximant also interpolates at this node, i.e.

$$\mathcal{I}(\mathbf{x}_i^{(j)}) = f_i^{(j)}, \quad i = 1, \dots, \bar{n}_j.$$

Solving the *j-th interpolation problem* (5) leads to a system of linear equations of the form

$$\begin{bmatrix} \phi(\|\mathbf{x}_1^{(j)} - \mathbf{x}_1^{(j)}\|_2) & \phi(\|\mathbf{x}_1^{(j)} - \mathbf{x}_2^{(j)}\|_2) & \cdots & \phi(\|\mathbf{x}_1^{(j)} - \mathbf{x}_{\bar{n}_j}^{(j)}\|_2) \\ \phi(\|\mathbf{x}_2^{(j)} - \mathbf{x}_1^{(j)}\|_2) & \phi(\|\mathbf{x}_2^{(j)} - \mathbf{x}_2^{(j)}\|_2) & \cdots & \phi(\|\mathbf{x}_2^{(j)} - \mathbf{x}_{\bar{n}_j}^{(j)}\|_2) \\ \vdots & \vdots & \vdots & \vdots \\ \phi(\|\mathbf{x}_{\bar{n}_j}^{(j)} - \mathbf{x}_1^{(j)}\|_2) & \phi(\|\mathbf{x}_{\bar{n}_j}^{(j)} - \mathbf{x}_2^{(j)}\|_2) & \cdots & \phi(\|\mathbf{x}_{\bar{n}_j}^{(j)} - \mathbf{x}_{\bar{n}_j}^{(j)}\|_2) \end{bmatrix} \begin{bmatrix} c_1^{(j)} \\ c_2^{(j)} \\ \vdots \\ c_{\bar{n}_j}^{(j)} \end{bmatrix} = \begin{bmatrix} f_1^{(j)} \\ f_2^{(j)} \\ \vdots \\ f_{\bar{n}_j}^{(j)} \end{bmatrix},$$

or simply

$$\Phi^{(j)} \mathbf{c}^{(j)} = \mathbf{f}^{(j)}. \quad (6)$$

- In particular, the interpolation problem is *well-posed*, i.e., a solution to the problem exists and is unique, if and only if the matrix  $\Phi^{(j)}$  is nonsingular.
- A sufficient condition to have nonsingularity is that the corresponding matrix is *positive definite*. In fact, if the matrix  $\Phi^{(j)}$  is positive definite, then all its eigenvalues are positive and therefore  $\Phi^{(j)}$  is nonsingular.

## Definition

Let  $\Omega \subseteq \mathbb{R}^N$  be a bounded set. Let  $\{\Omega_j\}_{j=1}^d$  be an open and bounded covering of  $\Omega$ . This means that all  $\Omega_j$  are open and bounded and that  $\Omega \subseteq \bigcup_{j=1}^d \Omega_j$ . Set

$\delta_j = \text{diam}(\Omega_j) = \sup_{\mathbf{x}, \mathbf{y} \in \Omega_j} \|\mathbf{x} - \mathbf{y}\|_2$ . We call a family of nonnegative functions  $\{W_j\}_{j=1}^d$  with  $W_j \in C^k(\mathbb{R}^N)$  a  $k$ -stable partition of unity with respect to the covering  $\{\Omega_j\}_{j=1}^d$  if

- 1)  $\text{supp}(W_j) \subseteq \Omega_j$ ;
- 2)  $\sum_{j=1}^d W_j(\mathbf{x}) \equiv 1$  on  $\Omega$ ;
- 3) for every  $\beta \in \mathbb{N}_0^N$  with  $|\beta| \leq k$  there exists a constant  $C_\beta > 0$  such that

$$\|D^\beta W_j\|_{L_\infty(\Omega_j)} \leq \frac{C_\beta}{\delta_j^{|\beta|}},$$

for all  $1 \leq j \leq d$ .

We require additional regularity assumptions on the *covering*  $\{\Omega_j\}_{j=1}^d$ .

### Definition

Suppose that  $\Omega \subseteq \mathbb{R}^N$  is bounded and  $\mathcal{X}_n = \{\mathbf{x}_i, i = 1, \dots, n\} \subseteq \Omega$  are given. An open and bounded covering  $\{\Omega_j\}_{j=1}^d$  is called regular for  $(\Omega, \mathcal{X}_n)$  if the following properties are satisfied:

- (a) for each  $\mathbf{x} \in \Omega$ , the number of subdomains  $\Omega_j$  with  $\mathbf{x} \in \Omega_j$  is bounded by a global constant  $K$ ;
- (b) each subdomain  $\Omega_j$  satisfies an interior cone condition;
- (c) the local fill distances  $h_{\mathcal{X}_j, \Omega_j}$ , where  $\mathcal{X}_j = \mathcal{X}_n \cap \Omega_j$ , are uniformly bounded by the global fill distance  $h_{\mathcal{X}_n, \Omega}$ , i.e.

$$h_{\mathcal{X}_n, \Omega} = \sup_{\mathbf{x} \in \Omega} \min_{\mathbf{x}_i \in \mathcal{X}_n} \|\mathbf{x} - \mathbf{x}_i\|_2.$$

# Theoretical result

## Theorem

Let  $\phi \in C_\nu^k(\mathbb{R}^N)$  be a strictly conditionally positive definite function of order  $m$ . Let  $\{\Omega_j\}_{j=1}^d$  be a regular covering for  $(\Omega, \mathcal{X}_n)$  and let  $\{W_j\}_{j=1}^d$  be  $k$ -stable for  $\{\Omega_j\}_{j=1}^d$ . Then the error between  $f \in \mathcal{N}_\phi(\Omega)$  and its partition of unity interpolant (4) can be bounded by

$$|D^\alpha f(\mathbf{x}) - D^\alpha F(\mathbf{x})| \leq Ch_{\mathcal{X}_n, \Omega}^{(k+\nu)/2 - |\alpha|} |f|_{\mathcal{N}_\phi(\Omega)},$$

for all  $\mathbf{x} \in \Omega$  and all  $|\alpha| \leq k/2$ .

[Wendland (2005)]

## Remark

- If we compare this result with the global error estimates, we can see that the **partition of unity** preserves the **local approximation order** for the global fit.
- This means that we can efficiently compute **large RBF interpolants** by solving **small RBF interpolation problems** and then glue them together with the global partition of unity  $\{W_j\}_{j=1}^d$ .
- The partition of unity approach is a simple and effective technique to **decompose a large problem** into **many small problems** while at the same time ensuring that the accuracy obtained for the local fits is carried over to the global one.

# PART I: block-based interpolation algorithms

# Outline of block algorithms

The *interpolation algorithms* can be briefly described as follows:

- 1 Partition the domain  $\Omega$  into a finite/suitable number of blocks.
- 2 Consider a *block-based searching procedure* that establishes the minimal number of blocks to be examined, in order to localize the set of nodes for each subdomain.
- 3 Apply the *Partition of Unity Method (PUM)* which uses RBFs as local approximants.



*Properties:*

- efficiency  $\rightarrow$  optimal searching procedure;
- accuracy  $\rightarrow$  RBFs;
- high parallelism  $\rightarrow$  PUM + block-based partition process.

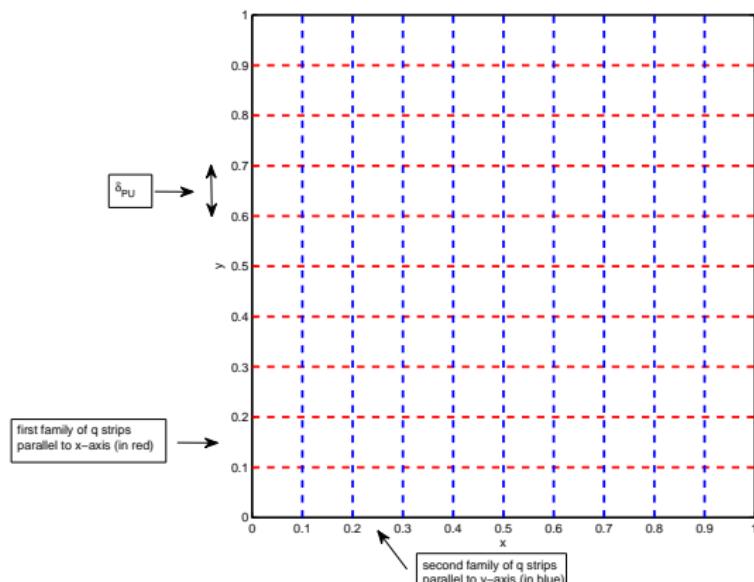
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REMARK: in the following, for simplicity, we restrict our focus on the unit square (domain), BUT our software works with generic (convex) domains!!!

## Basic idea - 2D case

- The basic idea in the construction of this searching procedure comes from the **repeated use** of a *quicksort routine* with respect to **different directions** (essentially, along the  $y$ -axis and the  $x$ -axis), enabling us to pass **from unordered to ordered data** structures.
- This process is strictly related to the construction of a partition of the domain  $\Omega$  in **square blocks**, which consists in generating **two orthogonal families of parallel strips**, where the original data set is suitably split up in ordered and well-organized data subsets.
- More precisely, to obtain the block-based partition structure/procedure, we act as follows:
  - 1 we organize all the data by a *quicksort<sub>y</sub> procedure* applied along the  $y$ -axis;
  - 2 we consider a **first family** of  $q$  strips, parallel to the  $x$ -axis and order the points of each strip by using a *quicksort<sub>x</sub> procedure*;
  - 3 we create a **second family** of  $q$  strips, parallel to the  $y$ -axis, which orthogonally intersect the first strip family  $\Rightarrow$  partition of  $\Omega$  in square blocks.

# Families of crossed-strips

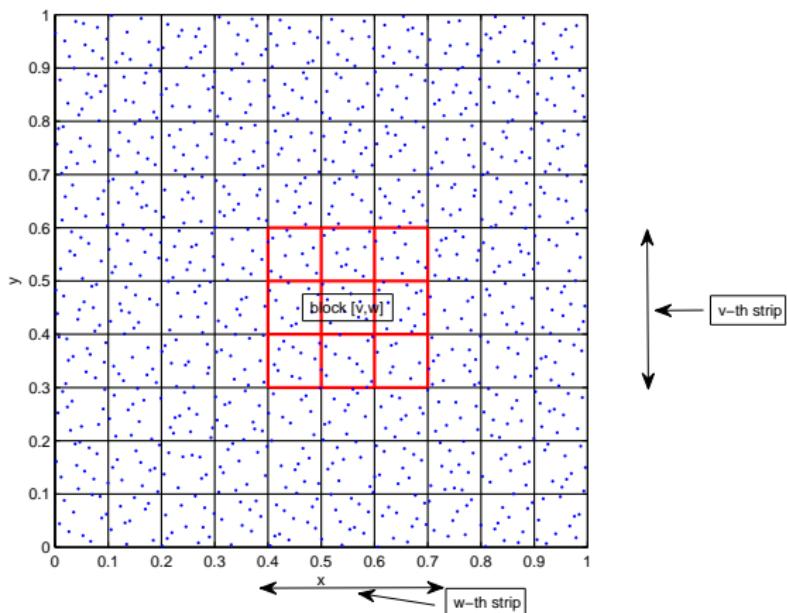


# Block-based searching procedure

- The aim is to construct an **efficient searching procedure** to be used in the **localization of points**, exploiting the data structure and the domain partition we have earlier described.
- An effective way to obtain an efficient searching technique is to connect the partition of unity method with the block-based partition structure, assuming that the **block width/side**  $\delta_{block}$  is **equal** to the **subdomain radius**  $\delta_{PU}$ , i.e.

$$\delta_{block} \equiv \delta_{PU}.$$

- Though this choice might seem to be trivial, in practice such an imposition means that the search of the nearby points is limited at most to **nine blocks**: the block on which the considered point lies, and the eight neighbouring blocks.
- The combination between block and subdomain sizes is an **optimal** choice, since it allows us to search the closest points only considering a very small number of them (that is only those points belonging to one of the nine blocks) and *a priori* ignoring all the other points of  $\Omega$ .
- Obviously, for all those points belonging to the **first and last blocks**, i.e. the ones close to the boundary of  $\Omega$ , a **reduction** of the total number of blocks to be examined will be required.



# Description of the 2D algorithm

The algorithm consists of three stages:

## 1. Distribution phase

- The nodes in the domain  $\Omega$  are ordered with respect to a common direction (e.g. the  $y$ -axis), by applying a *quicksort<sub>y</sub> procedure*.
- For each subdomain point  $(\bar{x}_i, \bar{y}_i)$ ,  $i = 1, \dots, d$ , a local circular subdomain is constructed, whose half-size (the radius) depends on the subdomain number  $d$ , that is

$$\delta_{PU} = \sqrt{\frac{2}{d}}.$$

- A double structure of crossed strips is constructed as follows:
  - i) a first family of  $q$  strips, parallel to the  $x$ -axis, is considered taking

$$q = \left\lceil \frac{1}{\delta_{PU}} \right\rceil,$$

and a *quicksort<sub>x</sub> procedure* is applied to order the nodes of each strip;

- ii) a second family of  $q$  strips, parallel to the  $y$ -axis, is considered.

Note that each of the two strip structures are ordered and numbered from 1 to  $q$ .

## 2. Localization phase

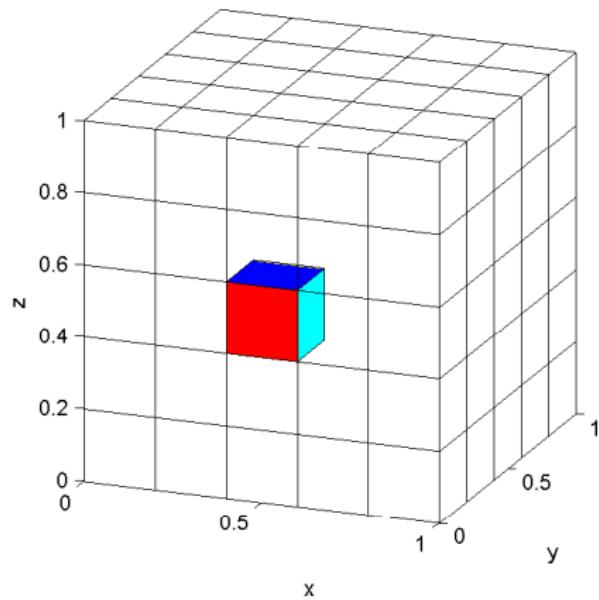
- The domain (unit square) is partitioned by a block-based structure consisted of  $q^2$  square blocks, whose length of the sides is given by  $\delta_{block} \equiv \delta_{PU}$ . Then, the following structure is considered:
  - the sets  $\mathcal{X}_n$  and  $\mathcal{C}_d$  are partitioned by the block structure into  $q^2$  subsets  $\mathcal{X}_{m_k}$  and  $\mathcal{C}_{d_k}$ ,  $k = 1, \dots, q^2$ ,  
where  $m_k$  and  $d_k$  are the number of points in the  $k$ -th block.
- After defining which and how many blocks are to be examined, a *block-based searching procedure* is applied for each subdomain point of  $\mathcal{C}_{d_k}$ ,  $k = 1, \dots, q^2$ , to determine all nodes belonging to a subdomain. The number of nodes of the  $j$ -th subdomain is counted and stored in  $n_j$ ,  $j = 1, \dots, d$ .
- Taking the  $\bar{n}_j$  nodes of the  $j$ -th subdomain, a *local interpolant*  $R_j$ ,  $j = 1, \dots, d$ , is found for each subdomain point.

### 3. Evaluation phase

- The evaluation points are ordered with respect to a common direction (e.g. the  $y$ -axis), by applying a *quicksorty procedure*.
- Then, the set  $\mathcal{E}_s$  is partitioned into  $q^2$  subsets  $\mathcal{E}_{p_k}$ ,  $k = 1, \dots, q^2$ , so that the evaluation points of  $\mathcal{E}_{p_k}$  belong to the  $k$ -th block.
- A block-based searching procedure is applied for each evaluation point of  $\mathcal{E}_s$ , in order to find all those points belonging to a subdomain of centre  $(\bar{x}_i, \bar{y}_i)$  and radius  $\delta_{PU}$ . The number of subdomains containing the  $i$ -th evaluation point is counted and stored in  $r_i$ ,  $i = 1, \dots, s$ .
- A **local approximant**  $R_j(x, y)$  and a **weight function**  $W_j(x, y)$ ,  $j = 1, \dots, d$ , are found for each evaluation point.
- Applying the **PUM** (4), the surface can be approximated at any evaluation point  $(x, y) \in \mathcal{E}_s$ .

# Basic idea - 3D case

- The basic idea in constructing this searching procedure comes from the **repeated use** of a *quicksort* routine with respect to **different directions** (here, along the  $z$ -axis, the  $y$ -axis and the  $x$ -axis), passing **from unordered to ordered data structures**.
- This process is strictly related to the construction of a partition of the domain (cube)  $\Omega$  in smaller **cubes**, which are obtained generating **three orthogonal families of parallelepipeds**, where the original data set is suitably split up in ordered and well-organized data subsets.
- More precisely, to obtain the cube-based structure/procedure, we act as follows:
  - 1 organize all the data by a *quicksort<sub>z</sub> procedure* applied along the  $z$ -axis;
  - 2 consider a **first family** of  $q$  parallelepipeds, parallel to the  $x$ -axis, and order the points of each parallelepiped by using a *quicksort<sub>x</sub> procedure*;
  - 3 create a **second family** of  $q$  parallelepipeds, parallel to the  $y$ -axis, which orthogonally intersect the first family, and order the points of each parallelepiped by using a *quicksort<sub>y</sub> procedure*;
  - 4 construct a **third family** of  $q$  parallelepipeds, parallel to the  $z$ -axis, which orthogonally intersect the two previous families  $\Rightarrow$  partition of  $\Omega$  in cubes.



# Extension to the 3D case

- Following the same idea described in the 2D case we obtain that the search of the nearby points is limited at most to **twenty-seven** ( $3^3$ ) cubes:
  - the **cube** on which the considered point lies,
  - and the **twenty-six** neighboring cubes.
- The combination between cube and subdomain sizes provides also here an **optimal** choice, allowing us to search the closest points only considering a very small number of them (only those belonging to one of the twenty-seven cubes) and *a priori* ignoring all the other points of  $\Omega$ .
- For all those points belonging to cubes close to the boundary of  $\Omega$ , it will be required a **reduction** of the total number of cubes to be examined.



**block-based searching procedure  
for 3D interpolation**

# Complexity of block algorithms

**Distribution phase:** to build the data structure  $\Rightarrow$  computational cost of order  $\mathcal{O}(M \log M)$  ( $M$  = number of nodes to be sorted) due to the [quicksort routine](#).

- $\mathcal{O}(n \log n)$  for the first sorting of all  $n$  nodes.

**Localization phase:** solution of  $d$  linear systems of size  $\bar{n}_j$  to compute the RBF coefficients:

- $\mathcal{O}(\bar{n}_j^3)$  arithmetic operations to compute the local RBF interpolants.

**Evaluation phase:** computational cost of order

- $r_i \cdot \mathcal{O}(\bar{n}_j)$  to evaluate the global interpolant at the  $i$ -th evaluation point.

**Storage locations:**

- $Nn$ ,  $Nd$  and  $Ns$  for the data, and  $\bar{n}_j$  for the coefficients of each local RBF interpolant.

# Comparison: block v.s. kd-tree

$N$	Block-based structure	kd-tree structure	Block-based search	kd-tree search
2	$\mathcal{O}(3/2n \log n) + \mathcal{O}(3/2s \log s)$	$\mathcal{O}(2n \log n) + \mathcal{O}(2s \log s)$	$\mathcal{O}(1)$	$\mathcal{O}(\log n) + \mathcal{O}(\log s)$
3	$\mathcal{O}(2n \log n) + \mathcal{O}(2s \log s)$	$\mathcal{O}(3n \log n) + \mathcal{O}(3s \log s)$	$\mathcal{O}(1)$	$\mathcal{O}(\log n) + \mathcal{O}(\log s)$

# Numerical experiments I

- Tests using the 2D Franke's function:

$$f_1(x, y) = \frac{3}{4} \exp \left[ -\frac{(9x-2)^2 + (9y-2)^2}{4} \right] + \frac{3}{4} \exp \left[ -\frac{(9x+1)^2}{49} - \frac{9y+1}{10} \right] \\ + \frac{1}{2} \exp \left[ -\frac{(9x-7)^2 + (9y-3)^2}{4} \right] - \frac{1}{5} \exp [-(9x-4)^2 - (9y-7)^2].$$

- RBFs with shape parameter  $\epsilon > 0$ :

$$\phi(r) = (1 - \epsilon r)_+^4 (4\epsilon r + 1), \quad \text{Wendland } C^2 \text{ function (W2),}$$

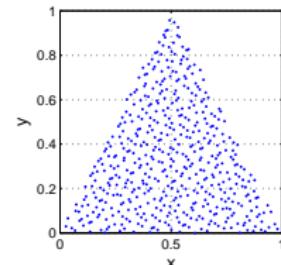
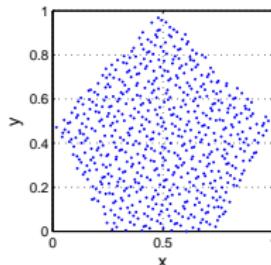
- Maximum Absolute Error (MAE) and Root Mean Square Error (RMSE):

$$MAE = \max_{1 \leq i \leq s} |f(\tilde{\mathbf{x}}_i) - \mathcal{I}(\tilde{\mathbf{x}}_i)|,$$

$$RMSE = \sqrt{\frac{1}{s} \sum_{i=1}^s |f(\tilde{\mathbf{x}}_i) - \mathcal{I}(\tilde{\mathbf{x}}_i)|^2}.$$

# Errors and CPU times

- Interpolation nodes: sets of Halton points (scattered data) in convex domains like a polygon  $\Omega \subseteq \mathbb{R}^2$  (e.g., triangle, hexagon, etc.).

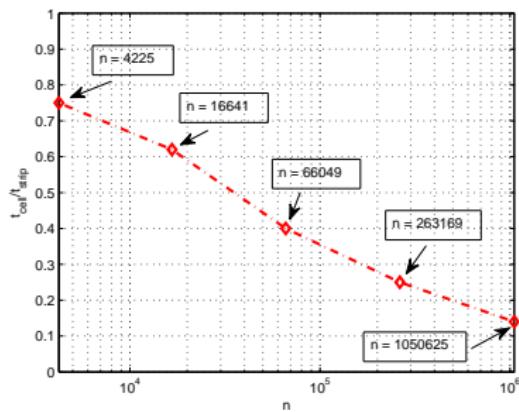


<i>n</i>	MAE	RMSE	<i>t<sub>block</sub></i>	<i>t<sub>kdtree</sub></i>
622	1.65E – 03	1.40E – 04	1.0	15.3
2499	5.02E – 04	3.30E – 05	3.7	42.3
9999	4.33E – 05	6.33E – 06	9.1	134.0
39991	9.86E – 06	1.25E – 06	34.1	494.1
159994	1.67E – 06	3.05E – 07	142.3	2013.88

Table: Errors and CPU times for pentagon,  $\varepsilon = 0.5$ .

# Comparison of CPU times (in seconds)

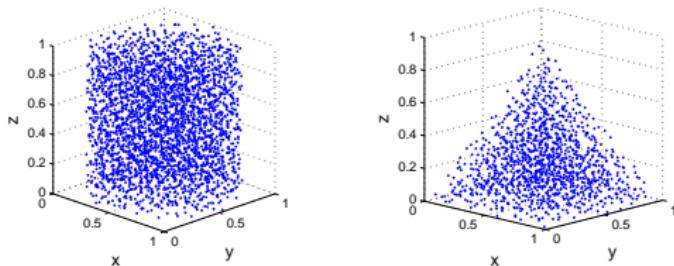
$n$	$t_{cell}$	$t_{strip}$	$t_{basic}$
4225	<b>0.3</b>	0.4	1.8
16641	<b>0.8</b>	1.3	14.2
66049	<b>2.6</b>	6.5	166.4
263169	<b>10.2</b>	41.2	2662.4



## Numerical experiments II

- Tests using the 3D Franke's function.
- W2-RBF with shape parameter  $\epsilon > 0$ .
- Interpolation nodes: sets of Halton points (scattered data) in convex domains like a polyhedron  $\Omega \subseteq \mathbb{R}^3$  (e.g., pyramid, cylinder, etc.).

$n$	$d$	$q^3$	$t_{cube}$	$t_{no-cube}$
4913	512	$6^3$	<b>2.2</b>	2.6
35937	4096	$12^3$	<b>16.6</b>	35.6
274625	32768	$23^3$	<b>138.2</b>	1241.0



$n$	MAE	RMSE	$t_{block}$	$t_{kdtree}$
3134	5.94E – 03	2.71E – 04	14.8	266.9
12551	1.67E – 03	6.00E – 05	53.1	892.7
50184	4.67E – 04	2.27E – 05	184.5	3141.4
200734	1.22E – 04	7.49E – 06	1758.1	14693.4
802865	3.81E – 05	2.91E – 06	-	-

**Table:** Errors and CPU times for cylinder,  $\varepsilon = 0.5$ .

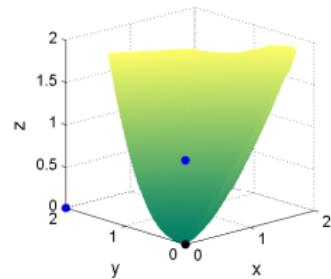
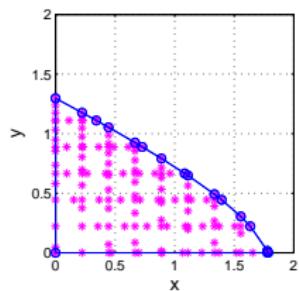
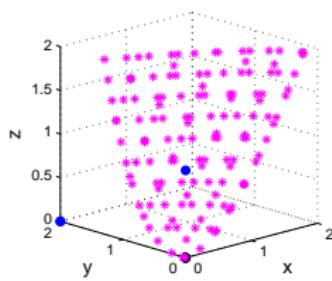
# Applications to geometric modelling

- Surface approximation from biomathematics:

- In **dynamical systems** saddle points partition the domain into basins of attraction of the remaining locally stable equilibria.
- This situation is rather common especially in population dynamics models, like competition systems. Trajectories with different initial conditions will possibly **converge toward different equilibria**, depending on the locations of their respective initial conditions.
- The set of all points that taken as initial conditions will have trajectories all tending to the same equilibrium is called the **basin of attraction** of that equilibrium point.

Example of competition model:

$$\begin{aligned}\frac{dx}{dt} &= p\left(1 - \frac{x}{u}\right)x - axy - bxz, \\ \frac{dy}{dt} &= q\left(1 - \frac{y}{v}\right)y - cxy - eyz, \\ \frac{dz}{dt} &= r\left(1 - \frac{z}{w}\right)z - fxz - gyz.\end{aligned}\tag{7}$$



- Reconstruction of 3D objects:

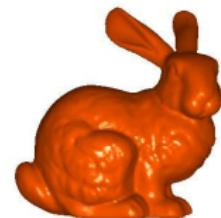
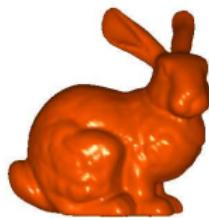


Figure: The Stanford Bunny with 8171 (left) and 35947 (right) data points.

## PART II: software

# MATLAB software

MATLAB codes free downloadable at:

<http://hdl.handle.net/2318/158790>

R. CAVORETTO, A. DE ROSSI, E. PERRACCHIONE, *Fast computation of partition of unity interpolants through block-based data structures*, submitted (2015).

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Computational issues:

- (i) **Range Search:** Given a set of data points  $\mathbf{x}_i \in \mathcal{X}_n$  and a subdomain  $\Omega_j$ , find all points situated in that subdomain, i.e.  $\mathbf{x}_i \in \mathcal{X}_j = \mathcal{X}_n \cap \Omega_j$ .
- (ii) **Containing Query:** Given  $\mathbf{x}_i \in \Omega$ , return all subdomains  $\Omega_j$  such that  $\mathbf{x}_i \in \Omega_j$ .

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PUM_2D_CSRBF.m	scripts performing the partition of unity using CSRBFs
PUM_3D_CSRBF.m	
BlockBased2D_Structure.m	scripts that store points into the different neighbourhoods
BlockBased3D_Structure.m	
BlockBased2D_ContainingQuery.m	scripts performing the containing query procedure
BlockBased3D_ContainingQuery.m	
BlockBased2D_RangeSearch.m	scripts that perform the range search procedure
BlockBased3D_RangeSearch.m	
BlockBased2D_DistanceMatrix.m	scripts that form the distance matrix of two sets of points for CSRBFs
BlockBased3D_DistanceMatrix.m	
inhull.m	script that tests if a point belongs to the convex hull
countingsort.m	script that performs a sorting routine for integers
haltonseq.m	script that generates Halton data

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Table: The MATLAB codes for the block-based partition of unity algorithms.

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Thank you!