

# Data Representation

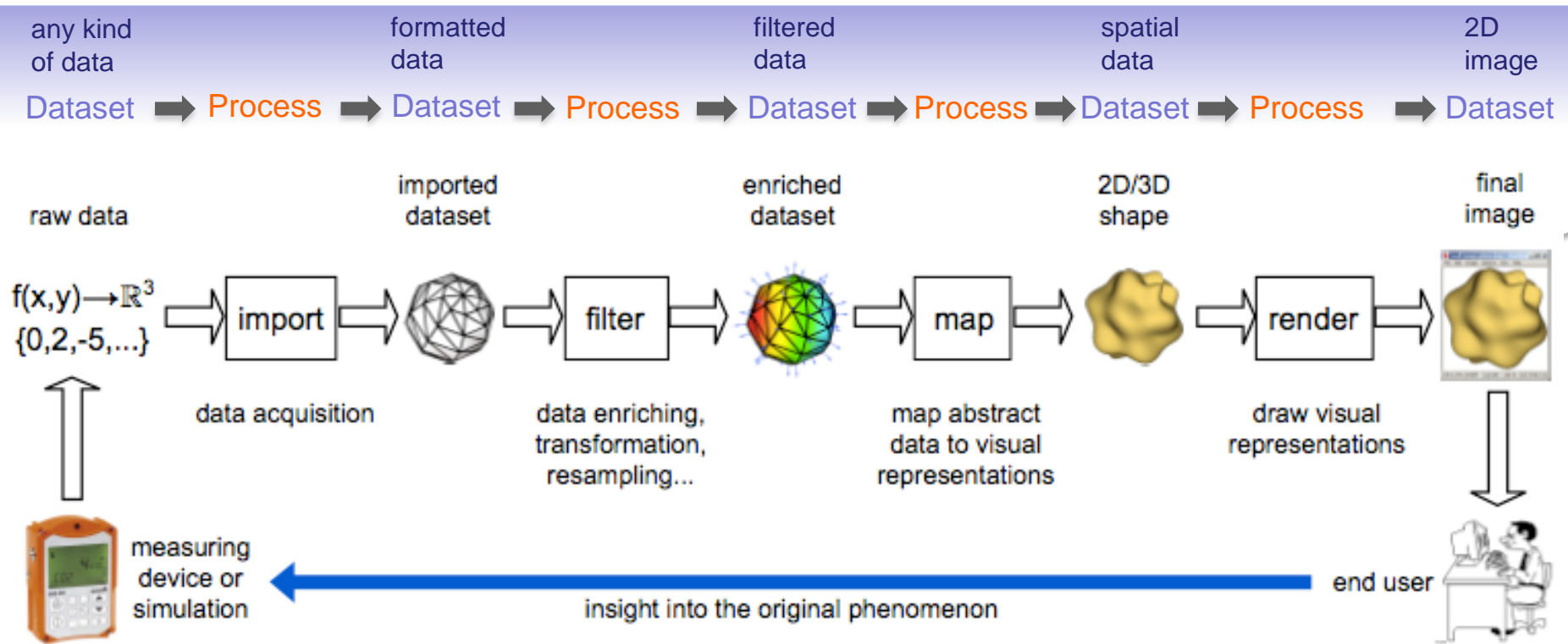
Cmpt 767 Visualization

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[based on slides by A. C. Telea]

# The Visualization Pipeline - Recall



# The Visualization Pipeline - Recall



## 1. Input data

- your primary “raw” source of information
- can be anything (measurements, simulations, databases, ...)

## 2. Formatted data

- converted to points, cells, attributes (discussed next in this module)
- Ready to use for visualization algorithms

## 3. Filtered data

- eliminates the unneeded **data**, adds the needed **information**
- read and written by visualization algorithms

## 4. Spatial (mapped) data

- has spatial embedding → can be **drawn**

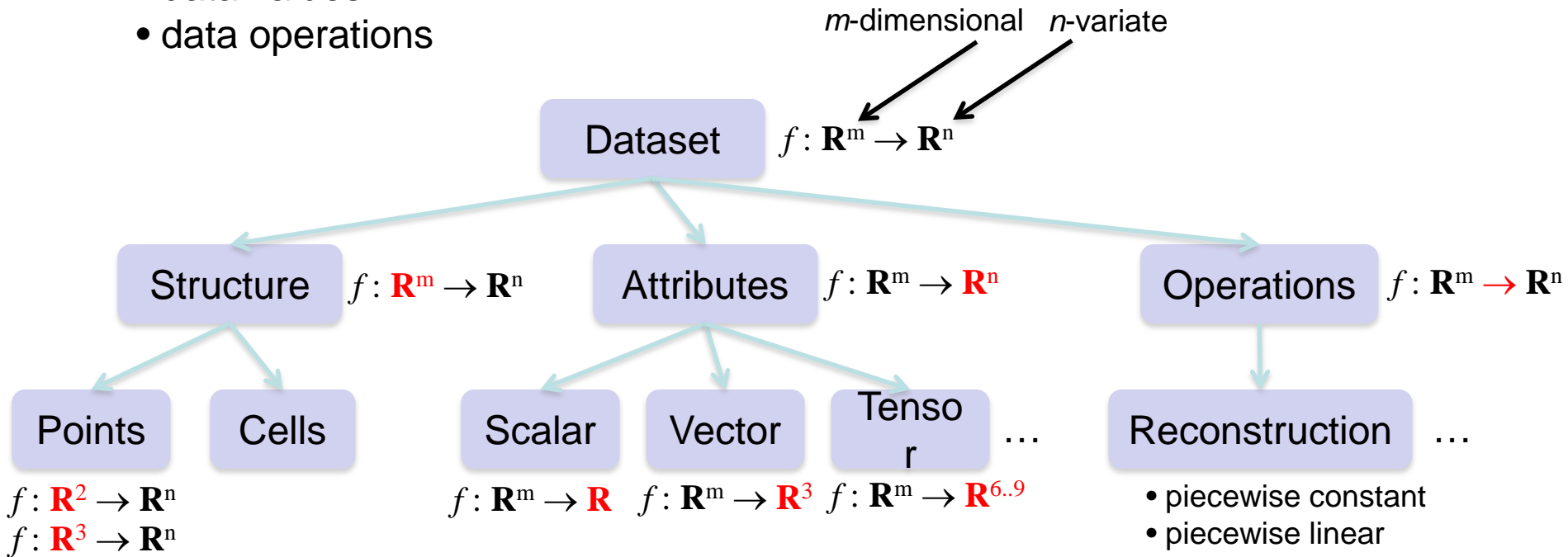
## 5. 2D Image

- final image you look at to get your answers
-

# Scientific Visualization - The Dataset

## Dataset

- key notion in visualization (SciVis, InfoVis, SoftVis)
- captures all relevant characteristics of a data collection
  - structure
  - data values
  - data operations



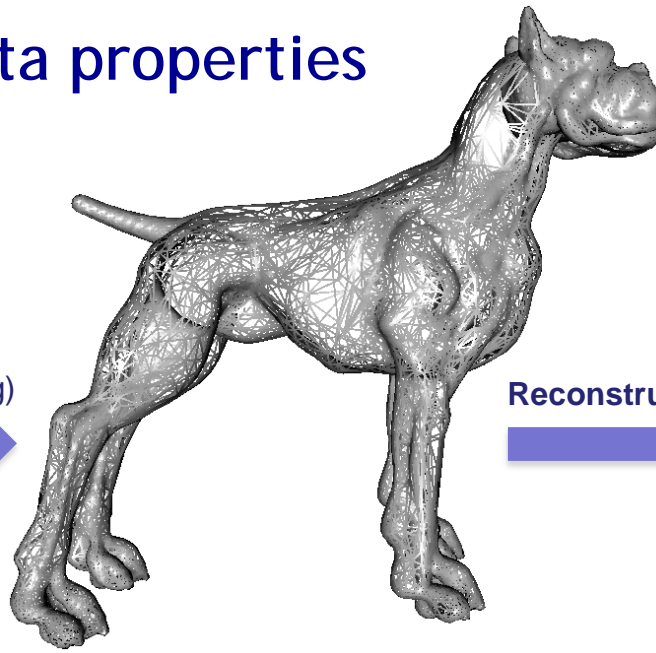
We'll detail all these next

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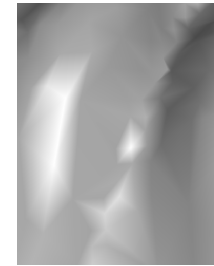
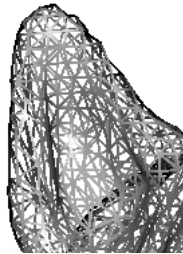
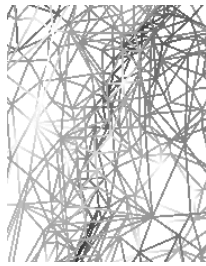
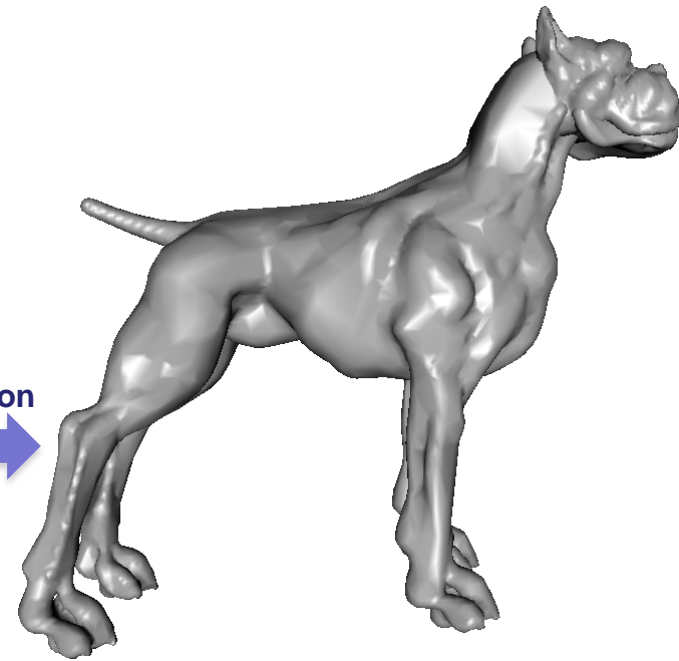
# Visualization data properties



Sampling  
(data importing)



Reconstruction



$$f : D \rightarrow C$$

Continuous data

Sampling



$$\{p_i, f_i\} \quad \begin{matrix} p_i \in D \\ f_i \in C \end{matrix}$$

Measurements (samples)  
at discrete set of points

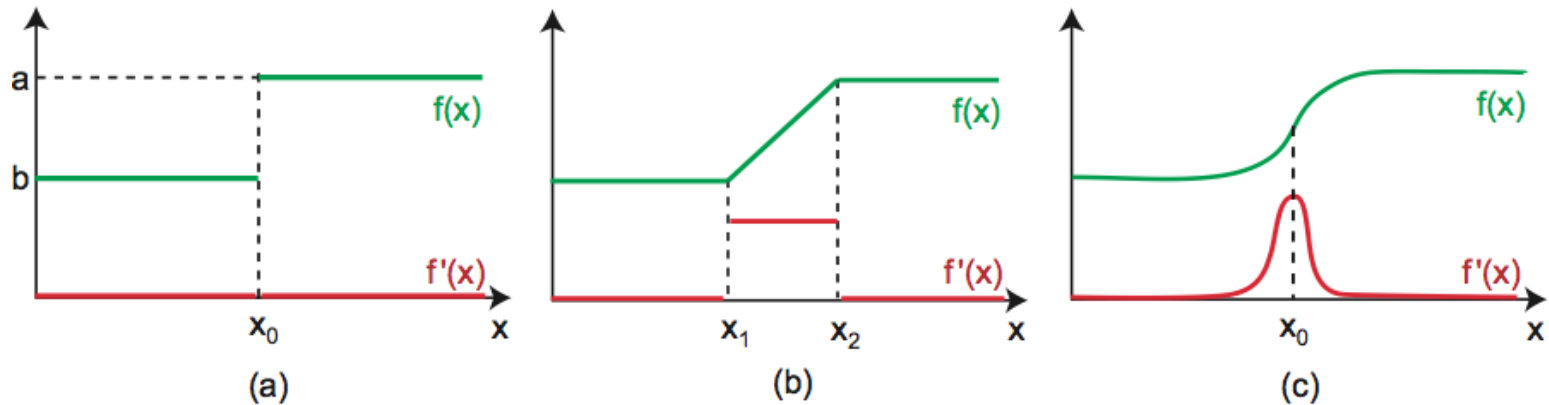
Reconstruction



$$\tilde{f} : D \rightarrow C \quad \tilde{f}(p_i) = f(p_i) = f_i$$

Continuous data, as close as possible to input

# Continuous data

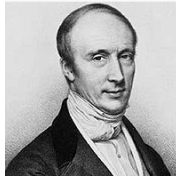


**Figure 3.1.** Function continuity. (a) Discontinuous function. (b) First-order  $C^0$  continuous function. (c) High-order  $C^k$  continuous function.

## Cauchy definition of continuity

A function  $f$  is continuous iff

$\forall \epsilon > 0, \exists \delta > 0$  such that if  $\|x - p\| < \delta, x \in C$  then  $\|f(x) - f(p)\| < \epsilon$ .



- $C^{-1}$  discontinuous (graph of function has “holes”)
- $C^0$  first-order continuous (graph of function has “kinks”)
- $C^k$  first  $k$  derivatives of the function are continuous

# Sampled data

## Functional properties

- **finite**
  - captures continuous signal at a finite set of points (measurements)
- **accurate**
  - can reconstruct a signal close to input accurately
  - reconstruction guarantees continuity properties

## Non-functional properties

- **efficient**
    - reconstruction is fast
  - **compact**
    - store Gbytes of sample points compactly
  - **generic**
    - few data structures cover most dataset types
  - **simple**
    - learn to create & use such data structures quickly
-

# Interpolation

Fundamental tool for signal reconstruction

1. **Reconstruction** formula

$$\tilde{f} = \sum_{i=1}^N f_i \phi_i \quad \phi_i : D \rightarrow \mathbb{C} \text{ are basis (or interpolation) functions}$$

2. **Interpolation**: reconstruction passes through (interpolates) the sampled values

$$\sum_{i=1}^N f_i \phi_i(p_j) = f_j, \forall j. \quad \text{because } \tilde{f}(p_i) = f(p_i) = f_i$$

3. **Orthogonality** of basis functions

$$\phi_i(p_j) = \begin{cases} 1, & i = j, \\ 0, & i \neq j. \end{cases}$$

$$\text{why? Just apply (2) to } f = \begin{cases} 1, & p = p_j \\ 0, & p \neq p_j \end{cases}$$

4. **Normality** of basis functions

$$\sum_{i=1}^N \phi_i(x) = 1, \forall x \in D$$

$$\text{why? } \sum_{i=1}^N \phi_i(p_j) = 1, \forall p_j \text{ (sum (3) over } i = 1..N) \\ \text{and apply above to all } p_i \in D$$



# Practical interpolation: Cells

Recall the interpolation formula


$$\tilde{f} = \sum_{i=1}^N f_i \phi_i$$

This becomes **very inefficient** if

- $N$  is very large and we have to evaluate  $\phi_i$  at all these  $N$  points
- $\phi_i$  have complicated expressions

## Practical basis functions

- are non-zero over small spatial ‘pieces’ of  $D$  only (limited support)
- have the same simple formula at all sample points  $p_i$

 We will discretize our spatial domain  $D$  into **cells**

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# Cells: 1D space

Consider a simple 1D function  $f: \mathbf{R} \rightarrow \mathbf{R}$

1. Sample the 1D axis at some points  $p_i$

2. Define **cells**  $c_i = (p_i, p_{i+1})$

3. Consider the **reference** basis functions for a reference cell  $(0,1)$

$$\phi_{0,1} : [0,1] \rightarrow [0,1], \quad \phi_0(r) = 1-r, \quad \phi_1(r) = r$$

4. Define a linear **transformation**  $T_i$  from the reference to actual cell  $c_i$

$$(x, y, z) = T(r, s, t) = \sum_{i=1}^n p_i \Phi_i^1(r, s, t)$$

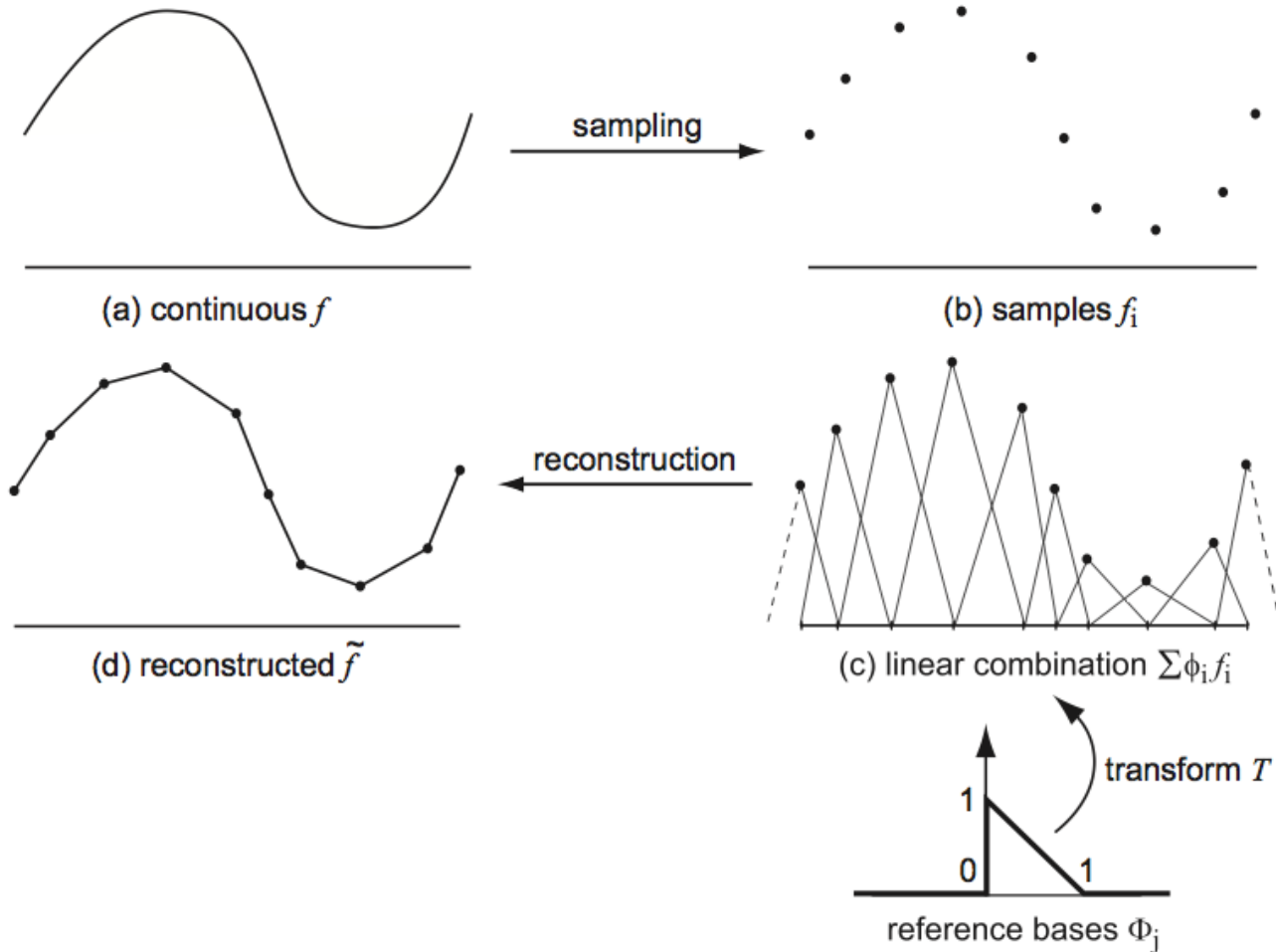
1. For  $c_i$ , define the actual basis functions  $\Phi_{0,1}$  using  $\phi_{0,1}$  and  $T_i^{-1}$  and rewrite the final interpolation

$$\tilde{f}(x, y) = \sum_{i=1} f_i \Phi_i^1(T^{-1}(x, y))$$

2. Apply (5) to interpolate all points in  $c_i$  using only samples at vertices  $p_i, p_{i+1}$  of  $c_i$

3. Repeat from 4 for next cell  $c_{i+1}$

# Cells: 1D example (cont'd)



## Remarks

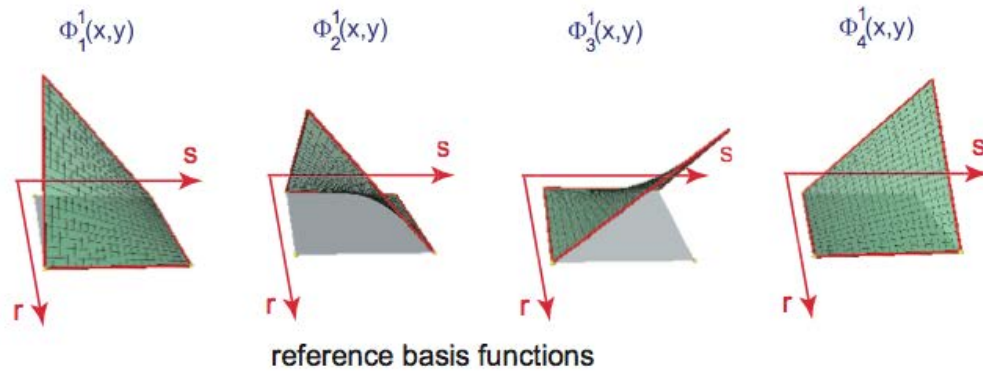
- interpolation & reconstruction goes cell-by-cell
- only need sample points at a cell vertices to interpolate over that cell
- reconstruction is  $C^1$  because  $\phi_i$  are  $C^1$  and interpolation formula is are  $C^\infty$

# 2D cells: Quads

Same as in 1D case, but

- we have to decide on different cells; say we take quads
- quads  $\rightarrow$  4 vertices, 4 basis functions
- particular case: square cells = pixels

## Bilinear basis functions



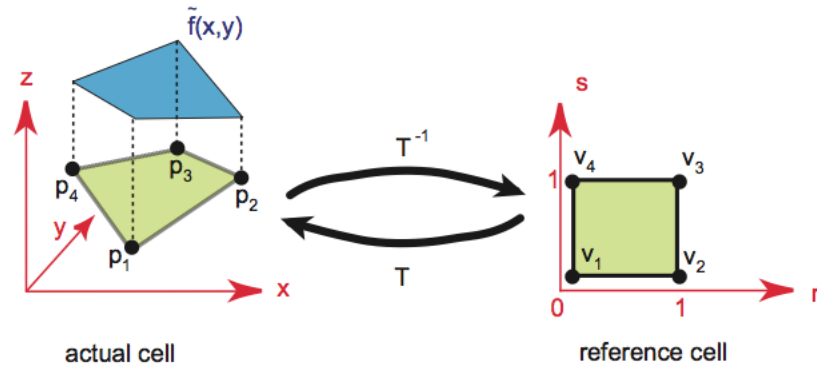
$$\Phi_1^1(r, s) = (1 - r)(1 - s),$$

$$\Phi_2^1(r, s) = r(1 - s),$$

$$\Phi_3^1(r, s) = rs,$$

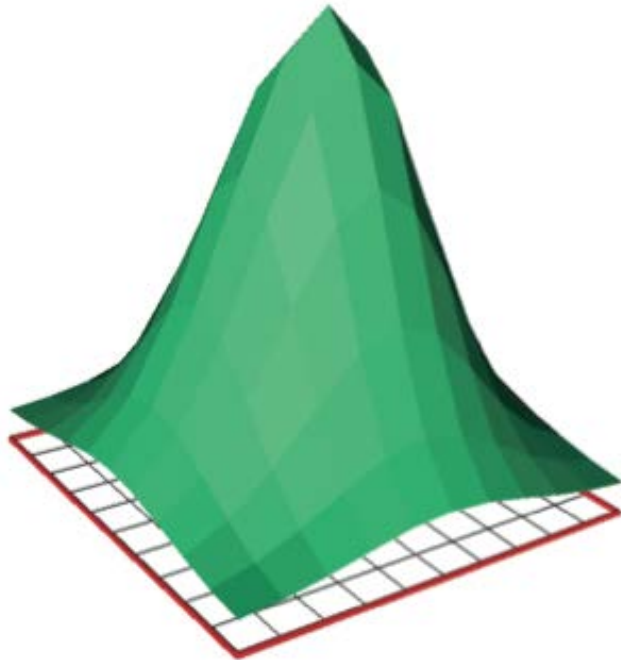
$$\Phi_4^1(r, s) = (1 - r)s;$$

## Bilinear transforms



# 2D cells: Quads

## Bilinear interpolation



$$\Phi_1^1(r, s) = (1 - r)(1 - s),$$

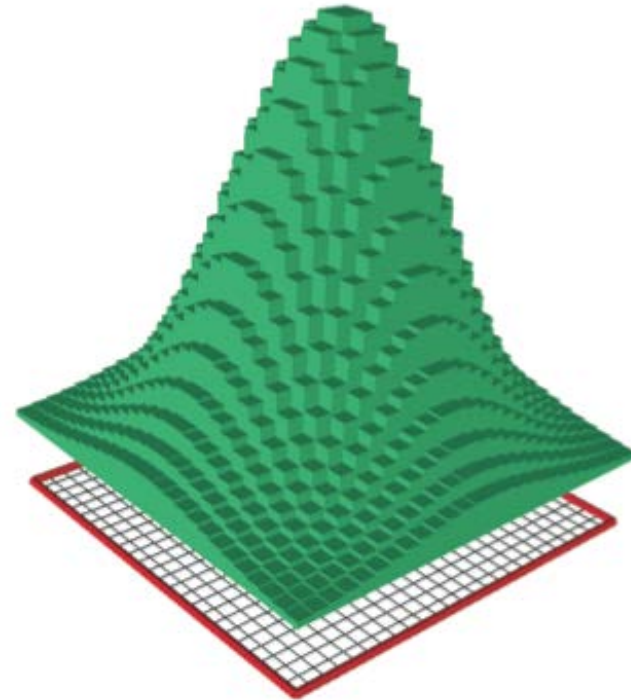
$$\Phi_2^1(r, s) = r(1 - s),$$

$$\Phi_3^1(r, s) = rs,$$

$$\Phi_4^1(r, s) = (1 - r)s;$$

- 4 functions, one **per vertex**
- result:  $C^0$  but never  $C^1$  (why?)
- good for **vertex-based** samples

## Constant interpolation

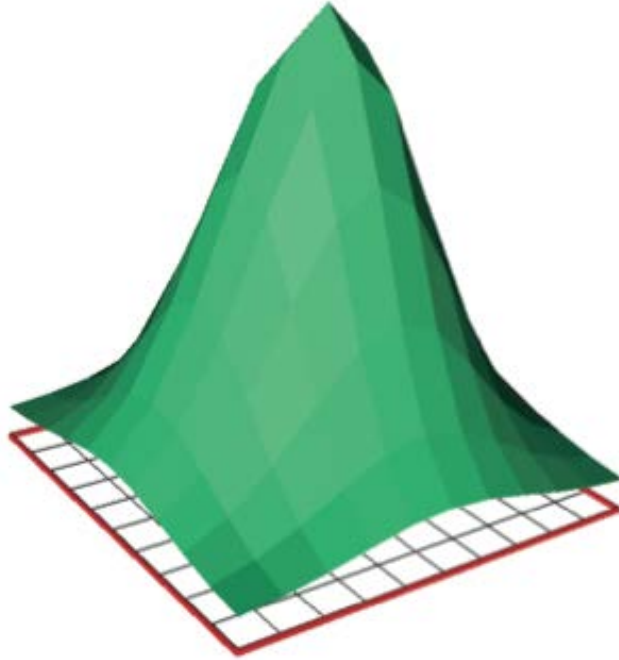


$$\phi_i^0(x) = \begin{cases} 1, & x \in c_i, \\ 0, & x \notin c_i. \end{cases}$$

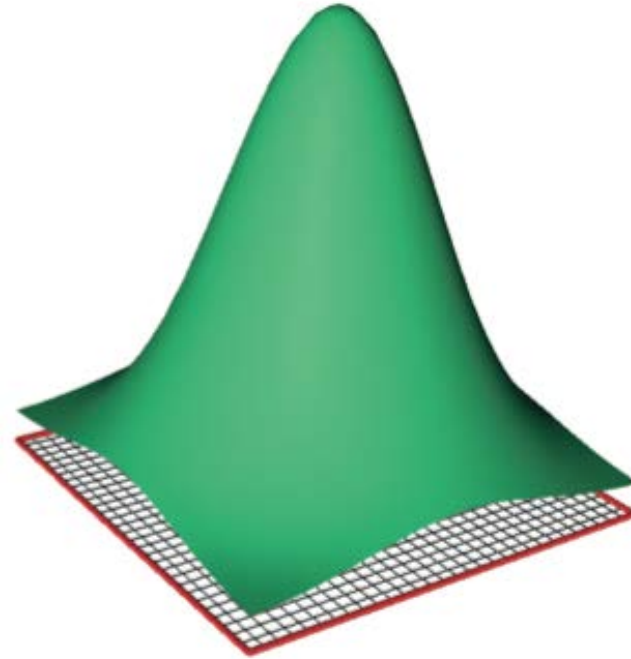
- 1 functions per **whole cell**
- result: not even  $C^0$
- good for **cell-based** samples

# Intermezzo

What is the difference between flat and Gouraud (smooth) shading?



Flat shading



Gouraud shading

- surface: bilinear interpolation
- colors: constant interpolation
- surface: bilinear interpolation
- colors: bilinear interpolation

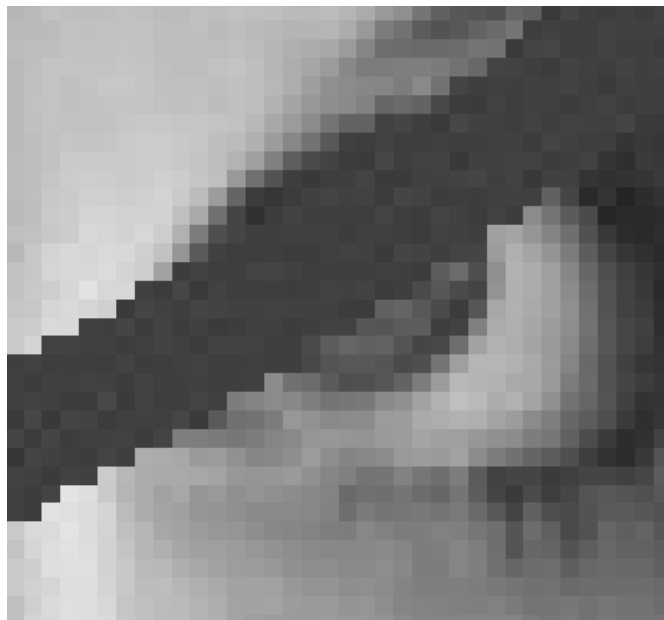
**Note:** do not confuse *Gouraud shading* (color interpolation) with the *Phong lighting model* (color computation from normals)

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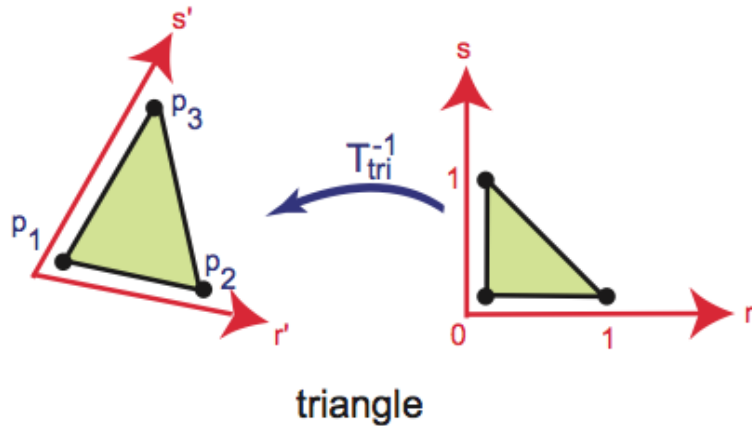
## 2D cells: Quads

### Images (color or grayscale)

- use constant basis functions
- cells = pixels
- data (color) is defined at the **center** of pixels, not corners
- we'll see why this is important in Module 3



# 2D cells: Triangles



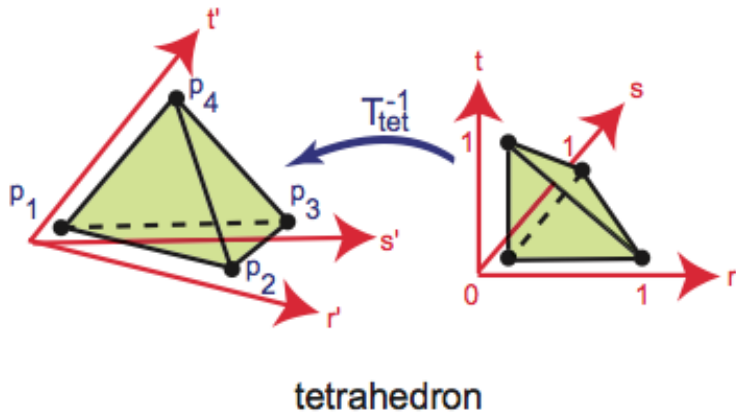
$$\begin{aligned}\Phi_1^1(r, s) &= 1 - r - s, \\ \Phi_2^1(r, s) &= r, \\ \Phi_3^1(r, s) &= s.\end{aligned}$$

## Remarks

- triangles and quads offers largely same pro's and con's
  - quad basis functions are not planes (they are **bi**linear)
  - in graphics/visualization, triangles used more often than quads
    - easier to cover complex shapes with triangles than quads
    - same computational complexity
-



# 3D cells: Tetrahedra

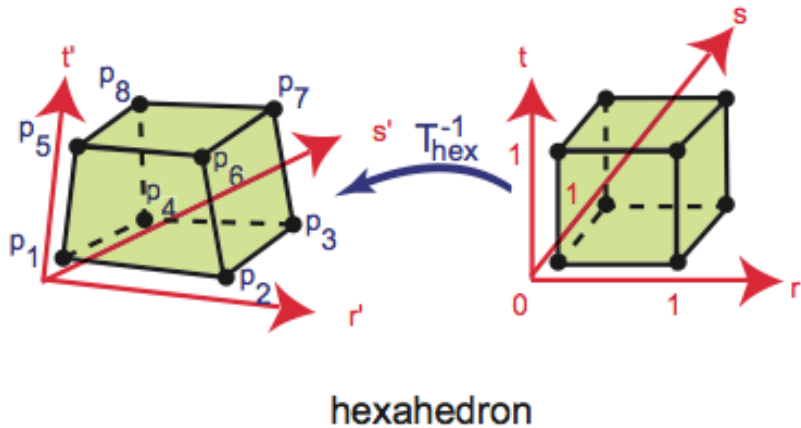


$$\begin{aligned}\Phi_1^1(r, s, t) &= 1 - r - s - t, \\ \Phi_2^1(r, s, t) &= r, \\ \Phi_3^1(r, s, t) &= s, \\ \Phi_4^1(r, s, t) &= t.\end{aligned}$$

## Remarks

- counterparts of triangles in 3D
  - interpolate **volumetric** functions  $f: \mathbf{R}^3 \rightarrow \mathbf{R}$
  - three parametric coordinates  $r, s, t$
  - **trilinear** interpolation
-

# 3D cells: Hexahedra



$$\Phi_1^1(r, s, t) = (1 - r)(1 - s)(1 - t),$$

$$\Phi_2^1(r, s, t) = r(1 - s)(1 - t),$$

$$\Phi_3^1(r, s, t) = rs(1 - t),$$

$$\Phi_4^1(r, s, t) = (1 - r)s(1 - t),$$

$$\Phi_5^1(r, s, t) = (1 - r)(1 - s)t,$$

$$\Phi_6^1(r, s, t) = r(1 - s)t,$$

$$\Phi_7^1(r, s, t) = rst,$$

$$\Phi_8^1(r, s, t) = (1 - r)st.$$

## Remarks

- counterparts of quads in 3D
- interpolate **volumetric** functions  $f: \mathbf{R}^3 \rightarrow \mathbf{R}$
- **trilinear** interpolation
- particular case: cubic cells or voxels (studied later in Module 7)

# Cell types for constant/linear basis functions

## 0D

- point

## 1D

- line

## 2D

- triangle, quad, rectangle

## 3D

- tetrahedron, parallelepiped, box, pyramid, prism, ...

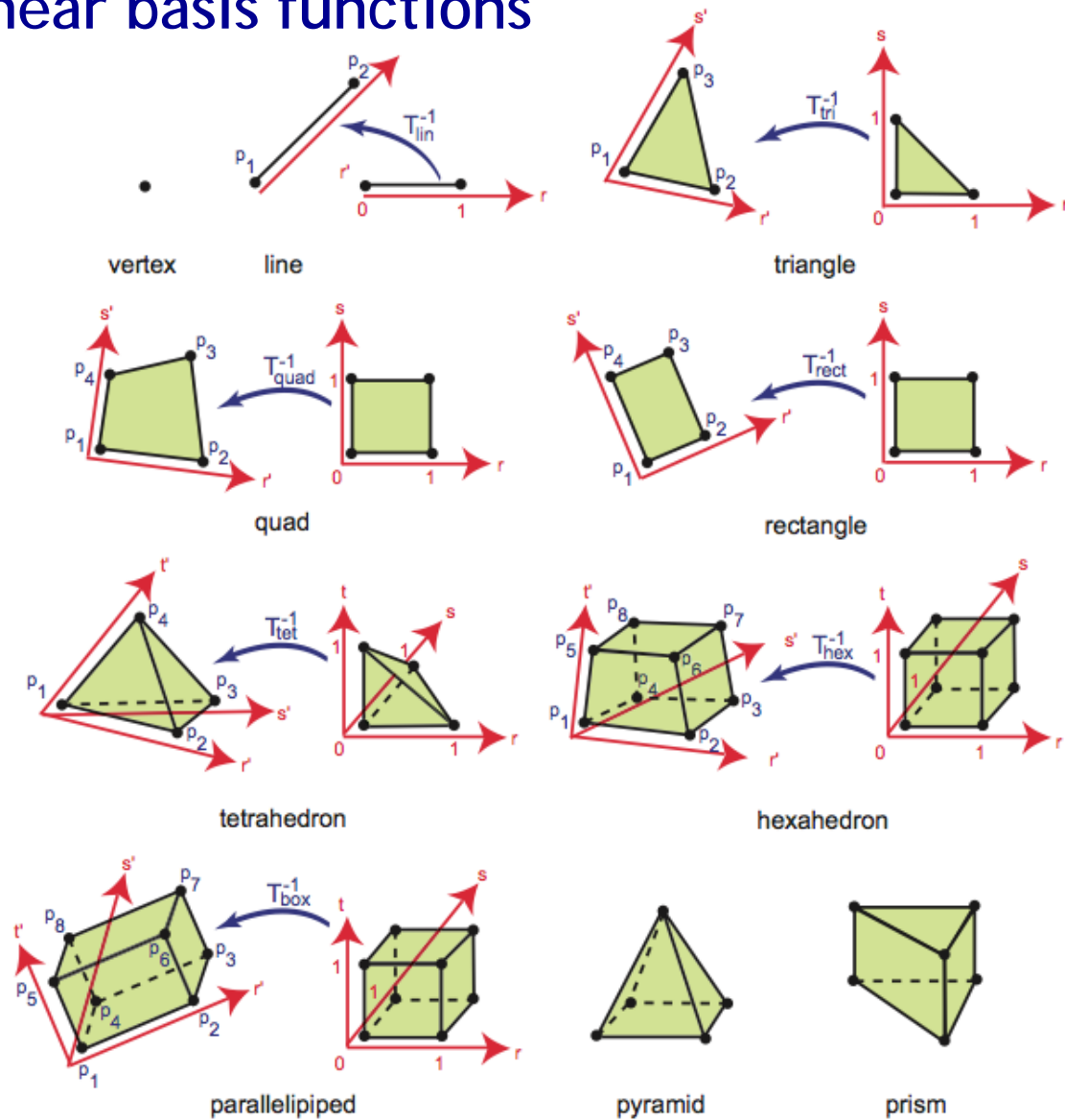
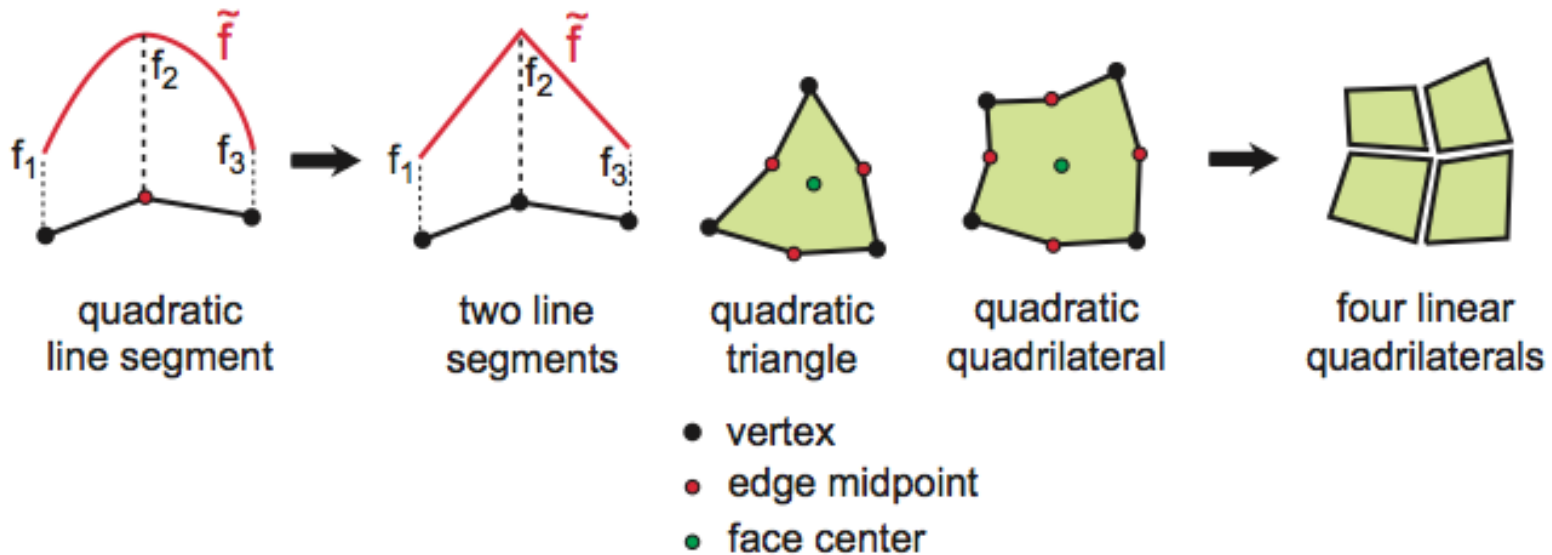


Figure 3.5. Cell types in world and reference coordinate systems.

# Quadratic cells



**Figure 3.6.** Converting quadratic cells to linear cells.

- allow defining **quadratic** basis functions
- higher **precision** for interpolation
- however, we need data samples at extra midpoints, not just vertices
- used in more complex numerical simulations (e.g. finite elements)
- split into linear cells for visualization purposes

# From cells to grids

## Cells

- provide interpolation over a small, simple-shaped spatial region

## Grids

- partition our complex data domain  $D$  into cells
- allow applying per-cell interpolation (as described so far)

Given a domain  $D$ ...

A **grid**  $G = \{c_i\}$  is a set of cells such that

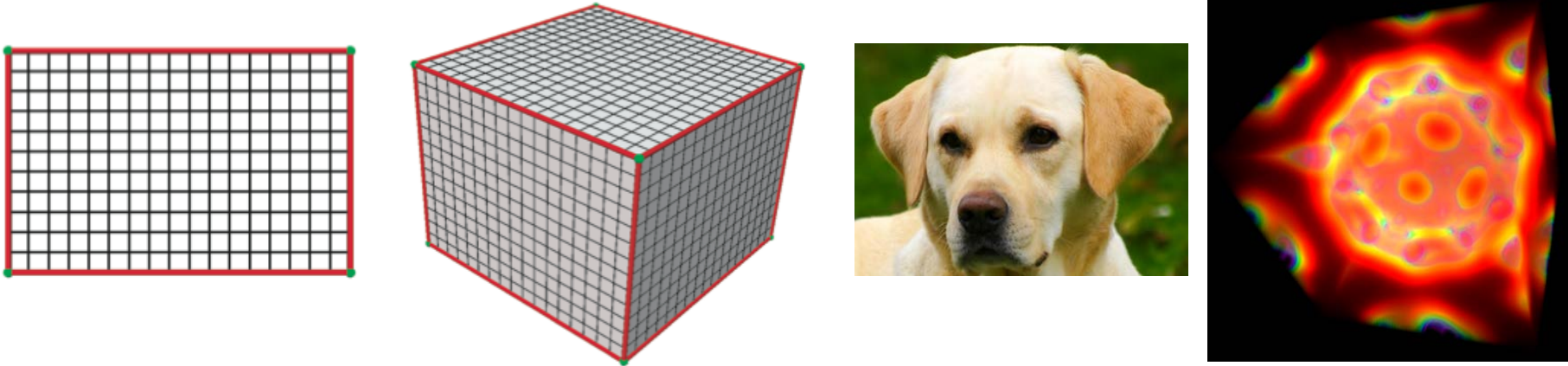
$c_i \cap c_j = \emptyset, \forall i \neq j$       no two cells overlap (why? Think about interpolation)

$\bigcup_i c_i = D$       the cells cover all our domain (why? Think about our end goal)

The dimension of the domain  $D$  constrains which cell types we can use: **see next**

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# Uniform grids



**Figure 3.7.** Uniform grids. 2D rectangular domain (left) and 3D box domain (right).

image

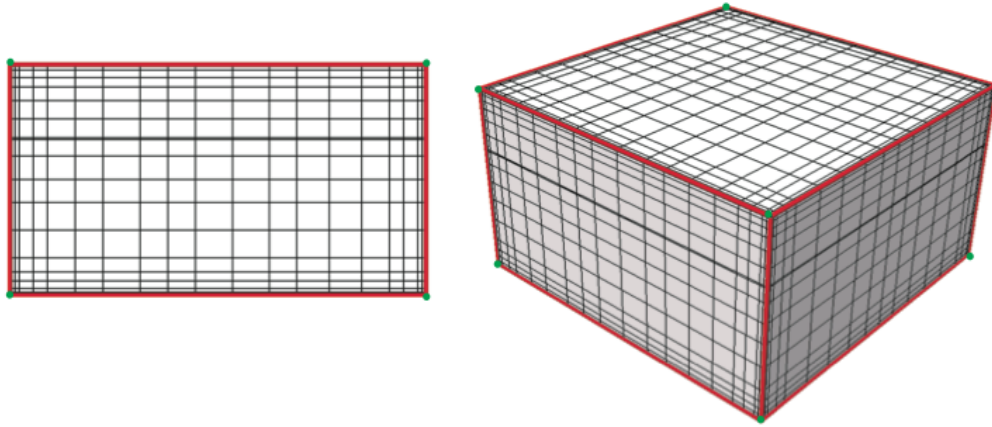
volume

- all cells have identical size and type (typically, square or cubic)
- cannot model non-axis-aligned domains

## Storage requirements

- $m$  integers for the #cells along each of the  $m$  dimensions of  $D$  (e.g.  $m=2$  or  $3$ )
-

# Rectilinear grids



**Figure 3.8.** Rectilinear grids. 2D rectangular domain (left) and 3D box domain (right).

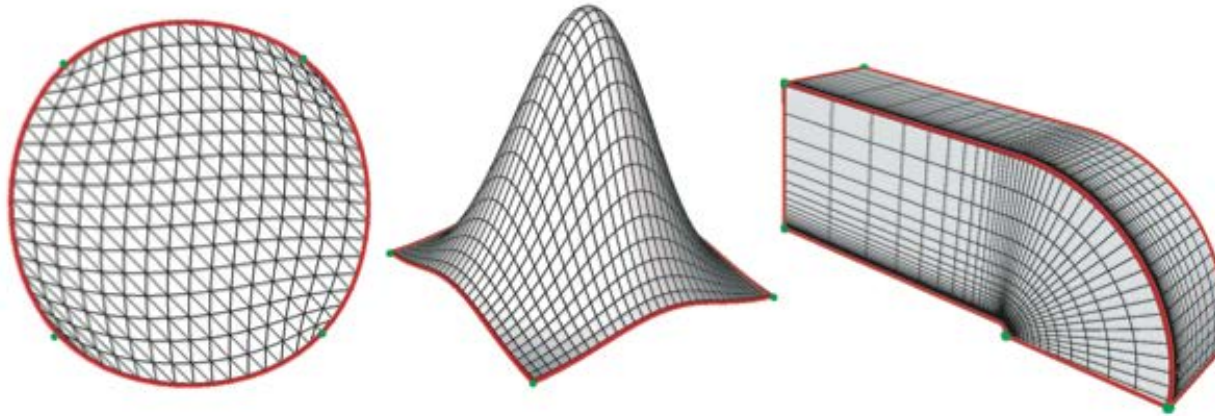
- all cells have same type
- cells can have different dimensions but share them along axes
- cannot model non-axis-aligned domains

## Storage requirements

$\sum_{i=1}^m d_i$  floats (coordinates of vertices along each of the  $m$  axes of  $D$ )

---

# Structured grids



**Figure 3.9.** Structured grids. Circular domain (left), curved surface (middle), and 3D volume (right). Structured grid edges and corners are drawn in red and green, respectively.

- all cells have same type
- cell vertex coordinates are freely (explicitly) specifiable...
- ...as long as cells assemble in a matrix-like structure
- can approximate more complex shapes than rectilinear/uniform grids

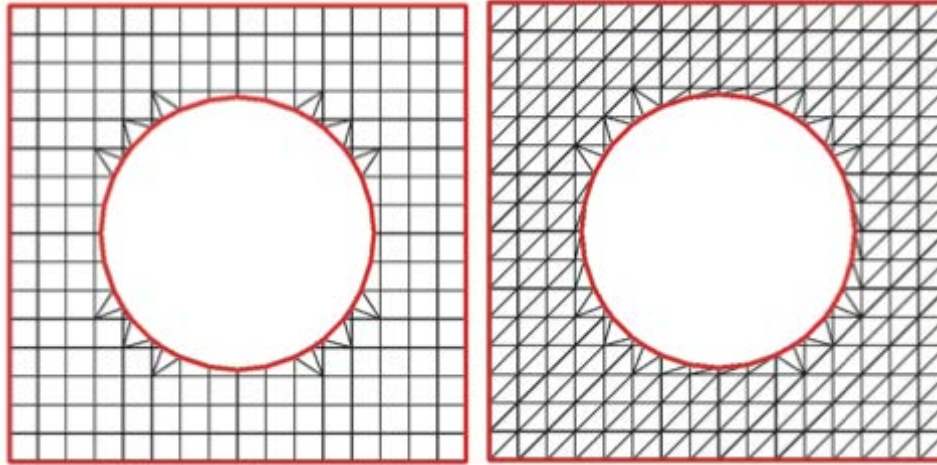
## Storage requirements

$\prod_{i=1}^m d_i$  floats (coordinates of all vertices)



# Unstructured grids

Consider the domain  $D$ : a square with a hole in the middle



We cannot cover such a domain with a structured grid (why?)

- it's not of genus 0, so cannot be covered with a matrix-like distribution of cells

For this, we need [unstructured grids](#) (see next)

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# Unstructured grids



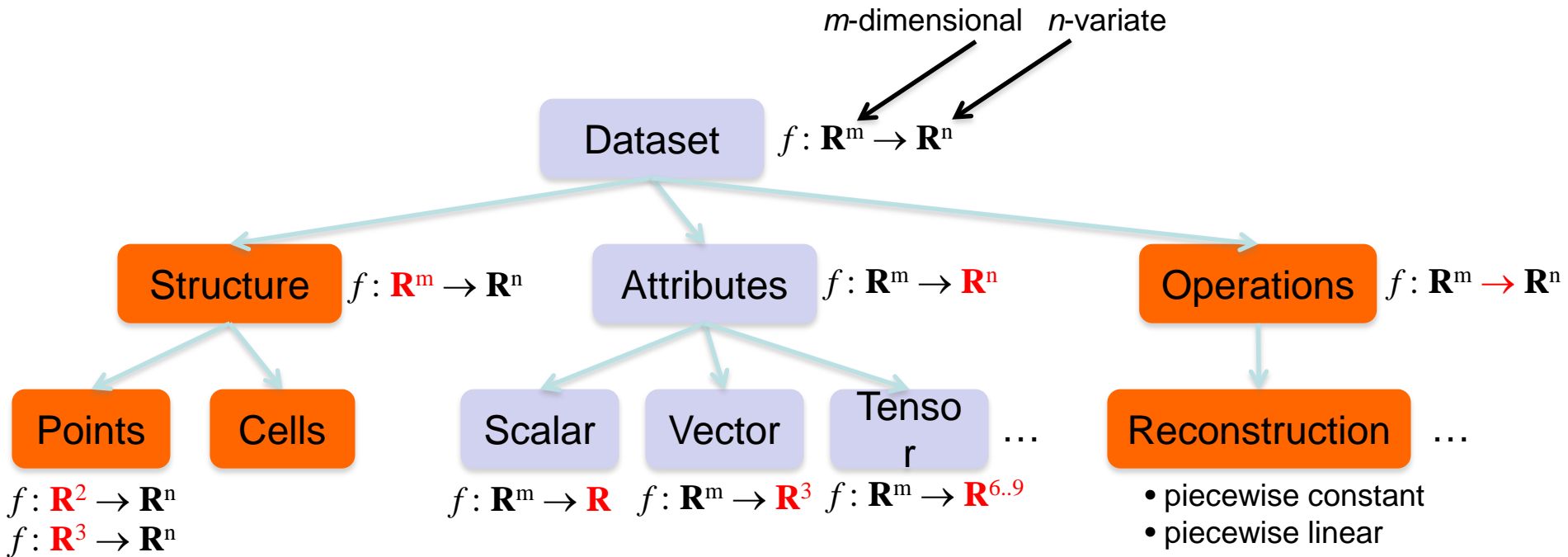
- different cell types can be mixed (though it's not usual)
- both vertex coordinates and cell themselves are freely (explicitly) specifiable
- implementation
  - vertex set  $V = \{v_i\}$
  - cell set  $C = \{c_i = (\textit{indices of vertices in } V)\}$
- most flexible, but most complex/expensive grid type

## Storage requirements

$m\|V\| + s\|C\|$  for a  $m$ -dimensional grid with cells having  $s$  vertices each

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# Recapitulation: Dataset



- We discussed about **these** (grids, interpolation, reconstruction)
  - We discuss next about **attributes**
-

# Data attributes

$$f: \mathbf{R}^m \rightarrow \mathbf{R}^n$$

- $n=0$  no attributes (we model a shape only e.g. a surface)
- $n=1$  scalars (e.g. temperature, pressure, curvature, density)
- $n=2$  2D vectors
- $n=3$  3D vectors (e.g. velocity, gradients, normals, colors)
- $n=6$  symmetric tensors (e.g. diffusion, stress/strain – Modules 5..6)
- $n=9$  assymetric general tensors (not very common)

## Remarks

- an attribute is usually specified for **all** sample points in a dataset (why?)
  - different measurements will generate different attributes
  - each attribute is interpolated separately
  - different visualization methods for each  $n$  (see Module 3 next)
-

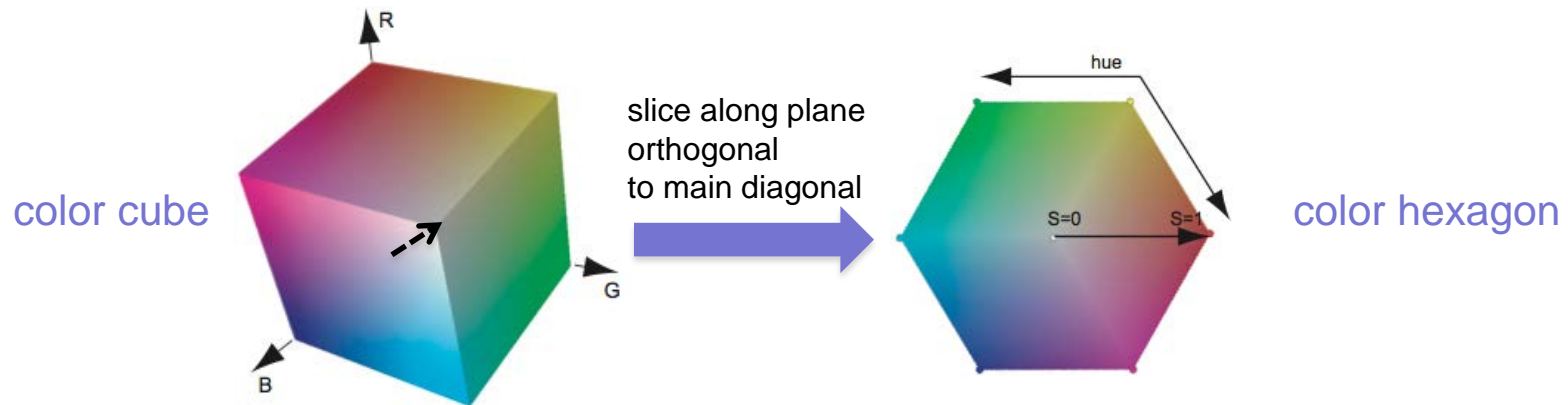
# Data attributes: Color

- complex topic (measurement, perception, representation)
- we'll mainly focus on **representation** and a bit on perception

## RGB color system

$$c = (c_R, c_G, c_B) \in [0, 1]^3$$

- three floating-point components in  $[0, 1]$
- additive system (add, or mix, components to obtain result)



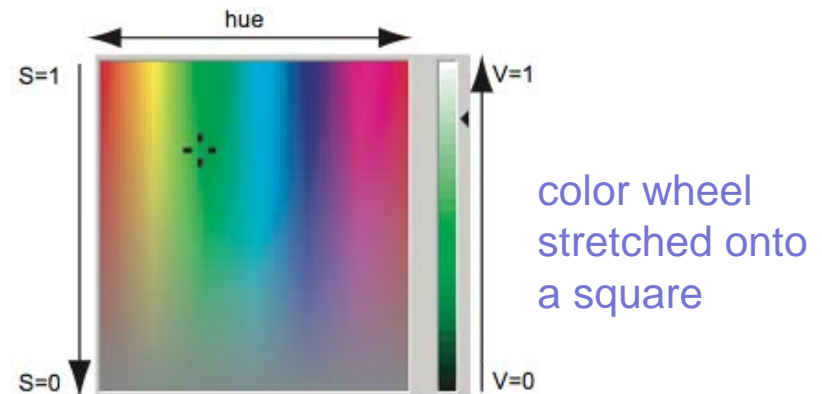
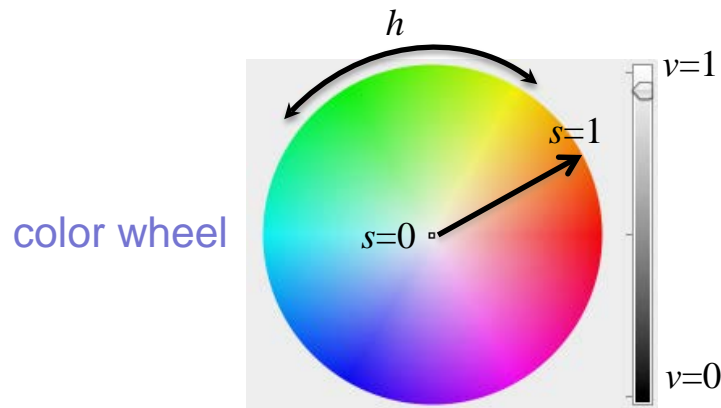
- perfect for synthesis (e.g. in the graphics card)
- unintuitive for humans, who think easier in **hues**

# HSV color system

- three floating-point components in  $[0,1]$

$$c = (h, s, v) \in [0, 1]^3$$

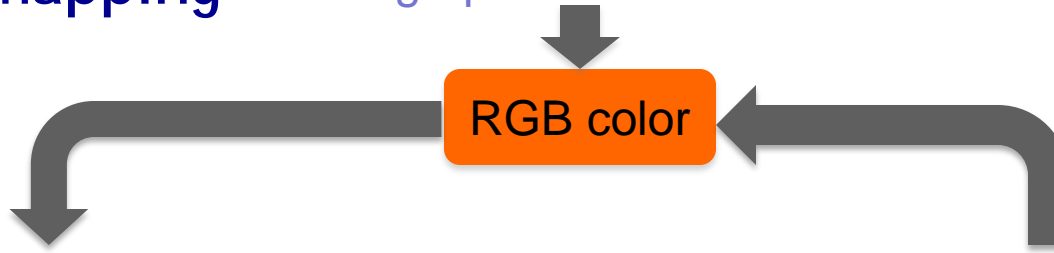
- hue: tint of the color (red, green, blue, yellow, cyan, magenta, yellow, ...)
- saturation: strong color ( $s=1$ ), grayish color ( $0 < s < 1$ ) or gray ( $s=0$ )
- value: luminance; white ( $v=1$ ), dark ( $0 < v < 1$ ), or black ( $v=0$ )



- HSV widgets: typically specify  $h$  and  $s$  in a 2D canvas and  $v$  separately (slider)
- show a 'surface slice' in the RGB cube

# RGB-HSV mapping

graphics software

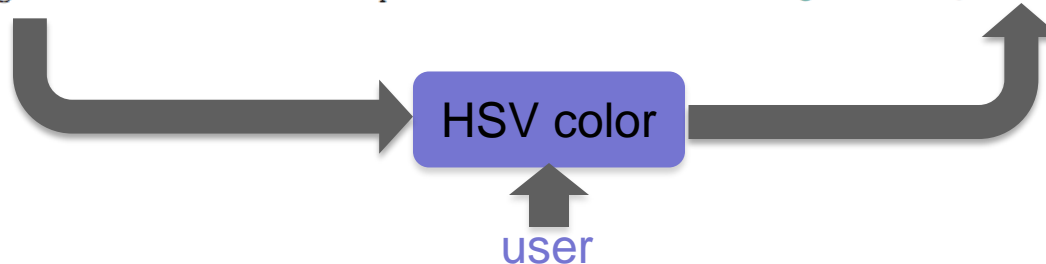


```
void rgb2hsv(float r, float g, float b,
            float& h, float& s, float& v)
{
    float M = max(r, max(g, b));
    float m = min(r, min(g, b));
    float d = M-m;
    v = M; //value = max(r,g,b)
    s = (M>0.00001)? d/M:0; //saturation
    if (s==0) h = 0; //achromatic case, hue=0 by convention
    else //chromatic case
    {
        if (r==M) h = (g-b)/d;
        else if (g==M) h = 2 + (b-r)/d;
        else h = 4 + (r-g)/d;
        h /= 6;
        if (h<0) h += 1;
    }
}
```

```
void hsv2rgb(float r, float g, float b,
            float& h, float& s, float& v)
{
    int hueCase = (int)(h*6);
    float frac = 6*h-hueCase;
    float lx = v*(1 - s);
    float ly = v*(1 - s*frac);
    float lz = v*(1 - s*(1 - frac));
    switch (hueCase)
    {
        case 0:
        case 6: r=v; g=lz; b=lx; break; // 0<hue<1/6
        case 1: r=ly; g=v; b=lx; break; // 1/6<hue<2/6
        case 2: r=lx; g=v; b=lz; break; // 2/6<hue<3/6
        case 3: r=lx; g=ly; b=v; break; // 3/6<hue<4/6
        case 4: r=lz; g=lx; b=v; break; // 4/6<hue<5/6
        case 5: r=v; g=lx; b=ly; break; // 5/6<hue<1
    }
}
```

Listing 3.2. Mapping colors from RGB to the HSV space.

Listing 3.3. Mapping colors from HSV to the RGB space.



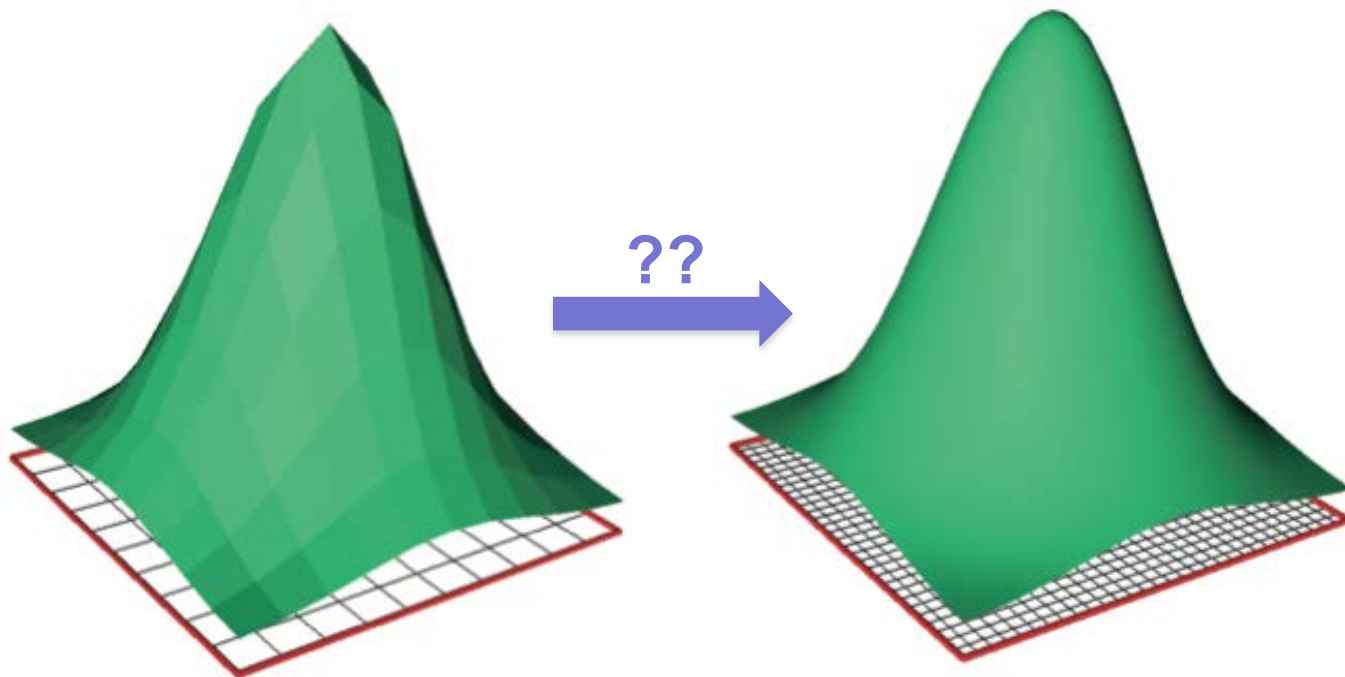
- simple conversions
- for details, see Chapter 3, pages 72-74



# Advanced data representation issues

## Data resampling

- consider building a Gouraud-shaded surface plot



Flat shading

- normals computed per **cell**

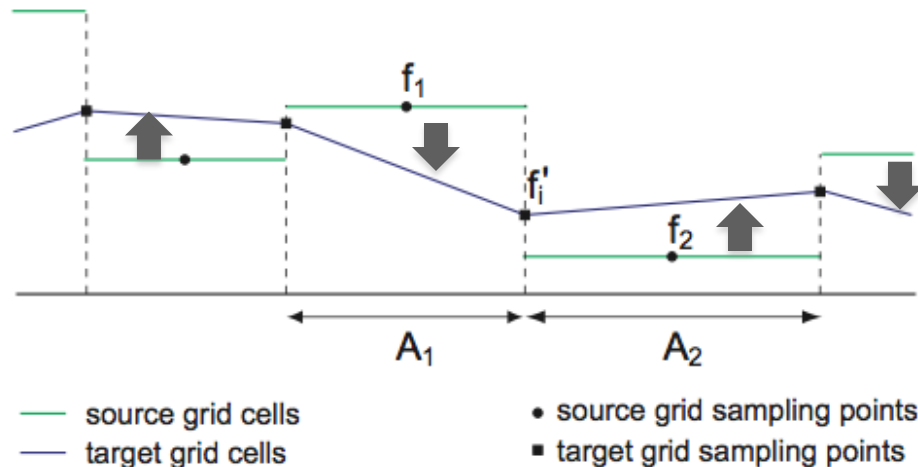
Gouraud shading

- normals required per **vertex**

- how to compute vertex attributes (normals) when we have cell attributes?
-



# Data resampling: cell data to vertex data and back



**Figure 3.16.** Converting cell to vertex attributes. The vertex value  $f'_i$  equals  $\frac{A_1 f_1 + A_2 f_2}{A_1 + A_2}$ , the area-weighted average of the cell values using vertex  $i$ .

Resampling a signal  $\tilde{f}$  over some target domain  $D'$  should yield a 'similar' signal  $\tilde{f}'$

$$\int_{c'_i} \tilde{f}' ds \approx \int_{c'_i} \tilde{f} ds, \quad \forall \text{ cells } c'_i \in D' \quad \xrightarrow{\text{see Sec. 3.9.1}} \quad f'_i = \frac{\sum_{c_j \in \text{cells}(p_i)} A(c_j) f_j}{\sum_{c_j \in \text{cells}(p_i)} A(c_j)}$$

•this is the classical area-weighted normal averaging used in Gouraud shading

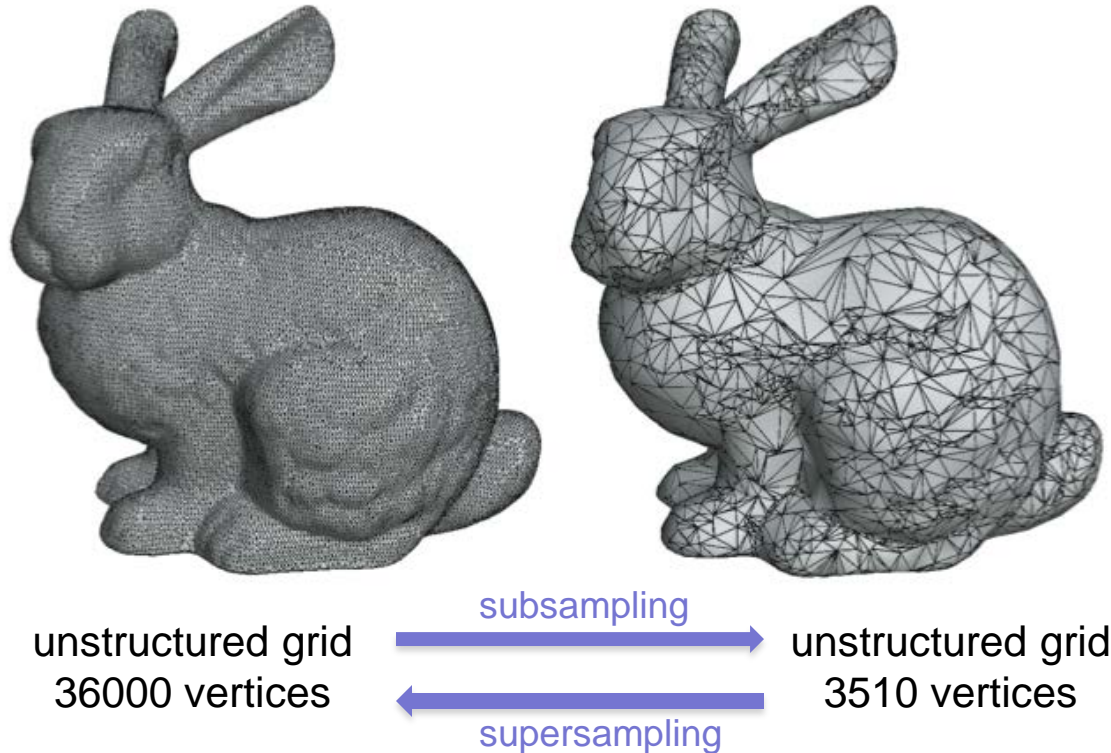
Resampling vertex data to cell data (same reasoning as above)

$$f_i = \frac{\sum_{p_j \in \text{points}(c_i)} f'_j}{C}$$

•this is the classical averaging of vertex values to compute cell values

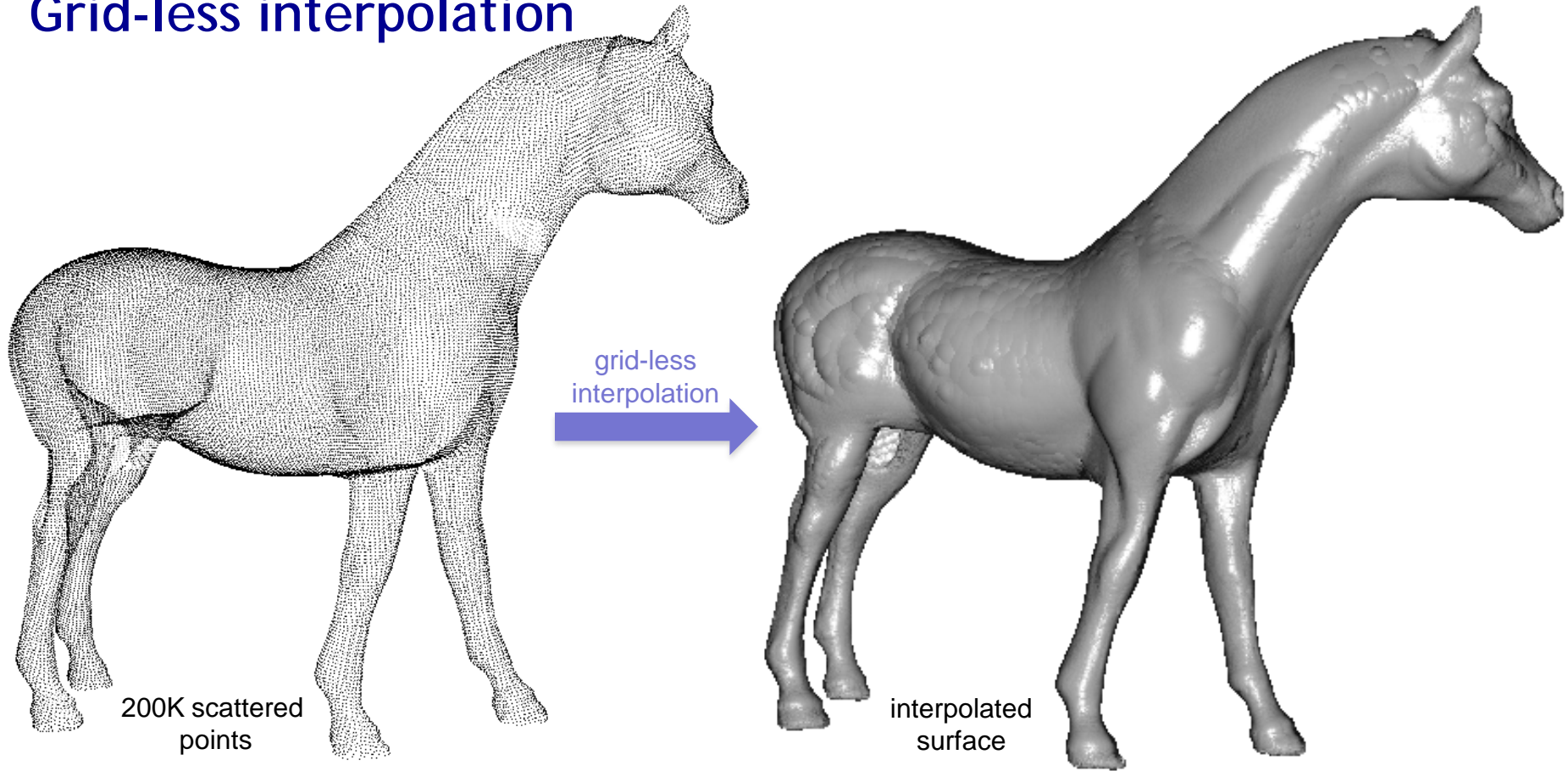
# Data super/subsampling

- we have data on some grid
- we want data on a 'similar' grid having more or less cells
- the interpolation functions stay the **same** (unlike in resampling)



- this is an advanced topic – treated separately in Module 6
-

# Grid-less interpolation



- we have data at some 'scattered' point locations in  $D$
  - we have no grid (cells connecting points)
  - we can
    - construct such a grid ([triangulation](#))
    - interpolate without a grid ([radial basis functions](#))
  - These advanced topics are discussed separately in Module 6
-

# Summary

## Data Representation (book Chapter 2)

- reconstruct continuous representations of sampled signals
  - efficiently
  - accurately
- interpolation, grids, and cells
- data attributes (scalars, vectors, tensors)
- advanced issues (resampling, grid-less interpolation)
- read Ch. 2 *in detail* to understand all the math!

## Next module

- visualization algorithms

Happy so far?

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