

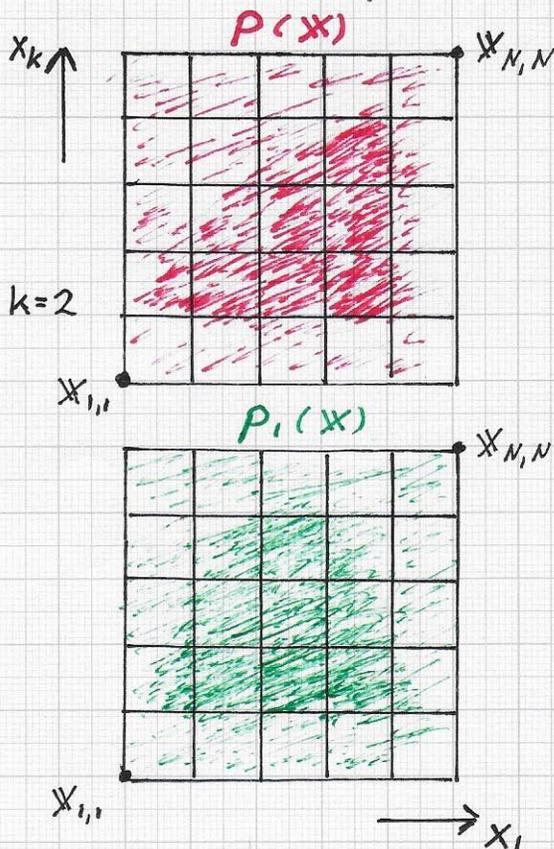
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DISTRIBUTION APPROXIMATION - Cont'd

→ Considerations and Issues (Details):

5) Comparison of discretized NEW

unclassified p-function, $p(x)$, and model p-function, $p_1(x)$, for example:



- $p(x)$ and $p_1(x)$ are represented by samples $p(x_{1,1}, \dots, 1), \dots, p(x_{N,1}, \dots, N)$ and $p_1(x_{1,1}, \dots, 1), \dots, p_1(x_{N,1}, \dots, N)$.

- The number of p-function samples is N^k .

- The number of features should be **SMALL!**

◦ Ex.: $k=5, N=32$
 $\Rightarrow 32^5 = 2^{25}$ p-function samples

- Shaded versions of $p(x)$ and $p_1(x)$ to be compared

◦ Ex.: resolution of 3D image: 512^3 ; (average) number of voxels of an object to be classified: $64^3 = 2^{18}$
 $\Rightarrow n = 2^{18} =$ no. of sites used for DISCRETE SIBSON interpolation

⇒ Root-mean-square (RMS) difference between $p(x), p_1(x)$:

$$d = \sqrt{\sum_{\hat{u}} (p(x_{\hat{u}}) - p_1(x_{\hat{u}}))^2}$$

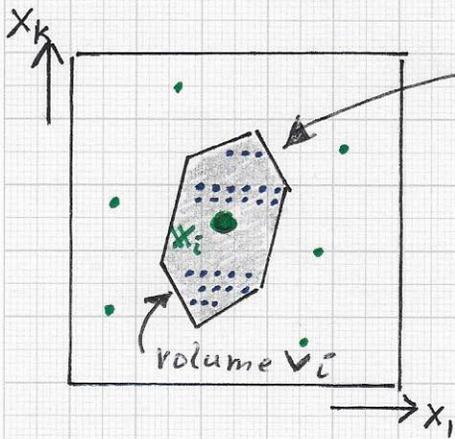
$\hat{u} = (1, \dots, 1), \dots, (N, \dots, N)$

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DISTRIBUTION APPROXIMATION - cont'd.

➔ More Considerations...

6) Computation/estimation of k-dim. tiles
 (= hyper-volumes) for definition of ρ_i
 values associated with original feature points



..... = points
closest to x_i ;
 hypercube volume
 associated with $\square = c$

⇒ $V_i = C \cdot c$

C = cardinality of
 set of all '•'
 inside x_i 's tile

tile of site $x_i (= \bullet)$,
tile hyper-volume = ?

⇒ Define a high-resolution
sampling approach to estimate
 hyper-volume V_i ;
determine set of samples in
 x -space closest to x_i ;
 assuming that this sample
 point set $\{ \dots \}$ consists
 of centers of hyper-cubes in
 k-dim. feature space, add the
 hyper-cube volumes to estimate V_i .

⇒ x_i has multiplicity m_i :

$\rho_i = \frac{m_i}{V_i}$

PARALLEL COMPUTATION

i) ρ_i -values for sites resulting from
 "training a high-quality" p -function
 can be pre-generated using high-res. sampling.

ii) ρ_i -values for sites used to define a p -function
 for an object in a NEW 3D scan must be computed
RAPIDLY!

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■ DISTRIBUTION APPROXIMATION - Cont'd.

➔ Considerations...

7) Resolutions are crucially important for storage/memory cost, computational performance cost, and classification quality. These are possible resolution scenarios:

i) 3D image resolution: $512^3 = 2^{27}$

ii) Number of features: $5 = k$

iii) Number of voxels per segmented object: $64^3 = 2^{18}$

iv) Number of same-class objects

(used for "training" the model): $16 = 2^4$

v) Number of sites in feature space

(number of all feature points/vectors of all voxels of all same-class objects):

$$n = 16 \cdot 64^3 = 2^4 \cdot 2^{18} = 2^{22}$$

vi) Resolution of discretized Sibson

interpolation p-function over k-dim. space:

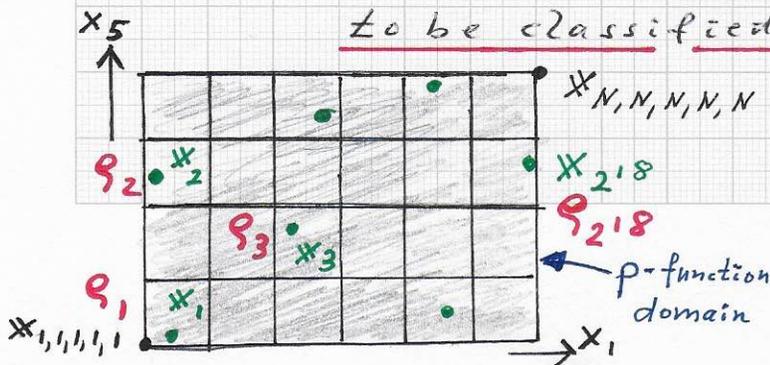
$$2^5 = 32 = N \text{ (each dimension)} \Rightarrow N^k = 2^{25}$$

vii) Number of different materials to

be recognized: $16 = 2^4$

viii) Number of voxels of segmented object

to be classified: $64^3 = 2^{18}$



● Example:

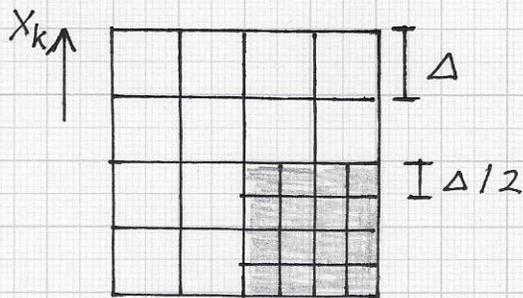
$\{x_1, x_2, \dots, x_{2^{18}}\}$ = single object's feature points in 5-dim. feature space; density values ρ_i used to compute p-values at $x_{i,j,k,l,m}$

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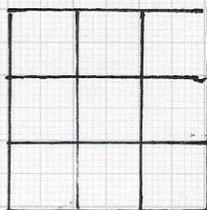
→ Considerations...

8) Adaptive grid spacing (= locally varying resolution) can be used to compute higher-precision p-function values where needed:

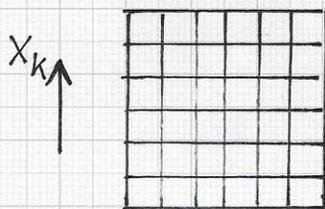


• p-function grids - adaptive

• p-function grids

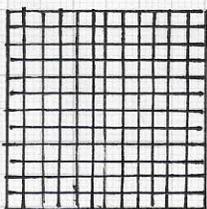


DECREASING



Δ/2

SPACING



Δ/4

→ x_l

• For example, use grid spacing Δ/2 in regions of the p-function's domain where the p-function varies more, i.e., where more feature points / vectors are given.

(The discrete Sibson interpolation step can incorporate adaptive spacing by using spacing-based weighted ρ_i -value additions.)

9) "Accelerated convergence" of discrete Sibson-based p-function via RICHARDSON extrapolation: spacing → 0.

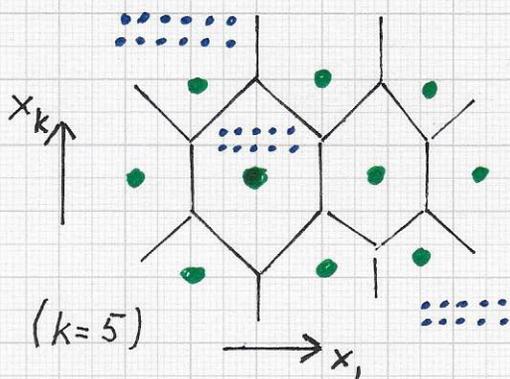
Using a measure for convergence of a gridded p-function, compute a sequence of such gridded functions using spacings $\Delta, \Delta/2, \Delta/4, \dots$ until threshold is met.

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⇒ Considerations...

10) Ratio of number of sites (tiles) of Sibson interpolation function and number of values of the p-function (evaluated on a uniform grid):



$\{ \cdot \}$ = set of sites
= $\{x_i\}$

$\{ \cdot \}$ = p-function values

Example: (see p. 13)

i) no. sites for MODEL: 2^{22}

ii) no. sites for OBJECT to be classified: 2^{18}

iii) no. of p-function values: 2^{25}

• Goal: much larger number of p-function values than number of sites ... !

⇒ Ratio $\frac{iii}{i} = \frac{2^{25}}{2^{22}} = 2^3 = \underline{8}$

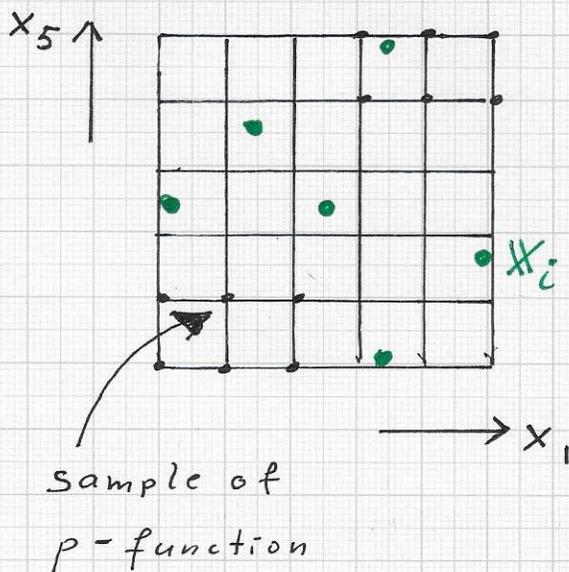
$\frac{iii}{ii} = \frac{2^{25}}{2^{18}} = 2^7 = \underline{128}$

• The feature space between sites (measured) and the feature space sampling density of the p-function (obtained via discrete Sibson interpolation) must be related such that desired classification performance is ensured.

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→ Considerations ...

- III) Normalization and floating-point representation: (i) A "meaningful normalization" of the individual dimensions of feature points/vectors \ast is necessary. (ii) The feature space grid points used to sample the p -function have floating-point coordinates in feature space; the p -values are also represented as floating-point numbers. (iii) All given, measured sites \ast also use floating-point representation.



→ original feature point/vector tuples:

$$\bullet = \ast_i = (x_1^i, \dots, x_5^i)$$

→ "multi-dimensional" (5-dim.) array of uniformly sampled \ast - and associated p -values:

$$\bullet = \ast_{\vec{i}} = (x_1^{\vec{i}}, \dots, x_5^{\vec{i}})$$

with multi-index $\vec{i} = (i_1, i_2, i_3, i_4, i_5)$;

$$p_{\vec{i}} = p_{i_1, i_2, i_3, i_4, i_5}$$

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