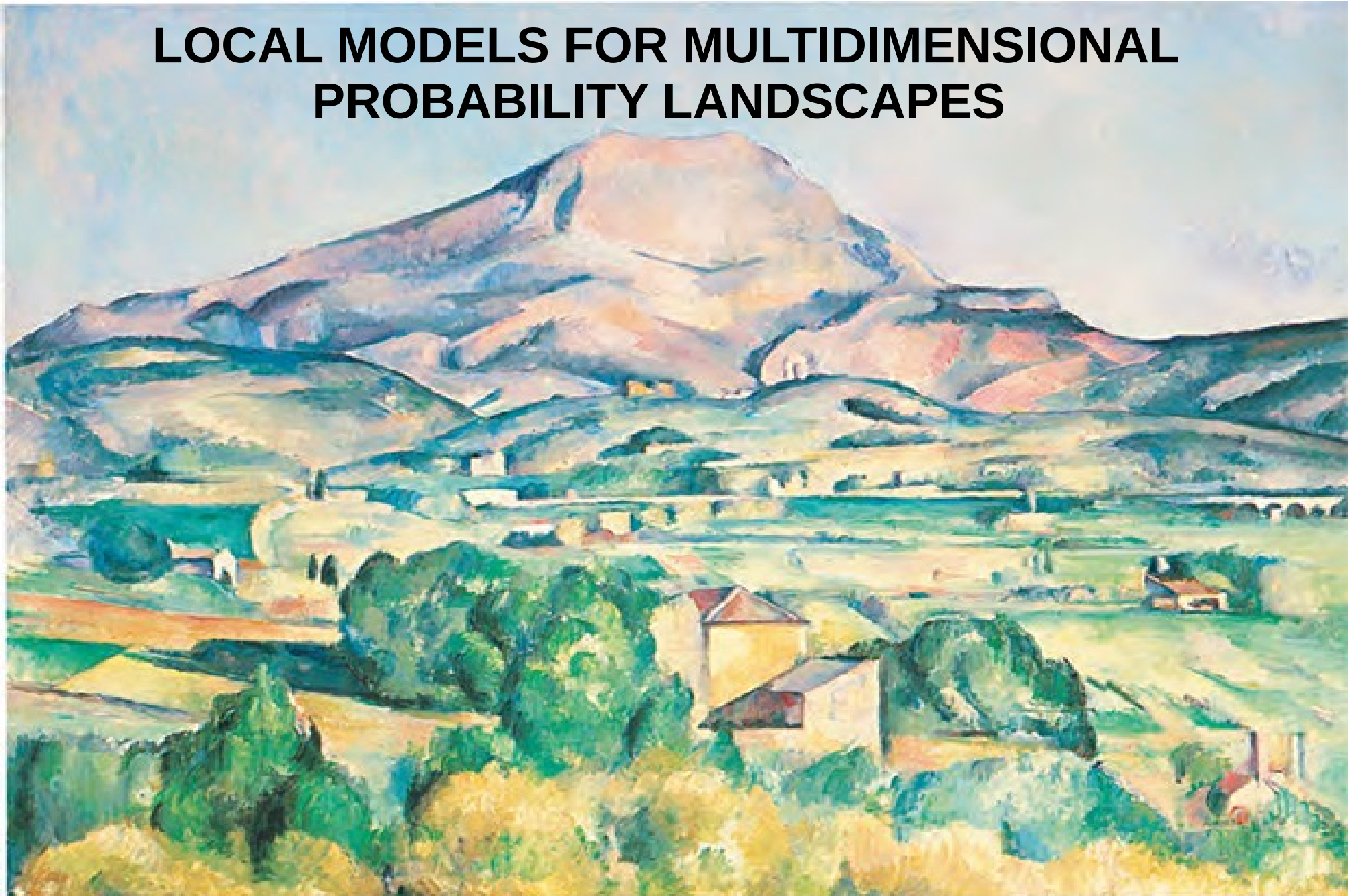


LOCAL MODELS FOR MULTIDIMENSIONAL PROBABILITY LANDSCAPES



Outline



- **Motivation: a rigorous basis for a clustering method**
- **A local model of nearest-neighbor distances under the assumption of locally constant density**
- **Density estimation**
- **Intrinsic dimension estimation**
- **An extension of the model for heterogeneous intrinsic dimension**

Density peak clustering

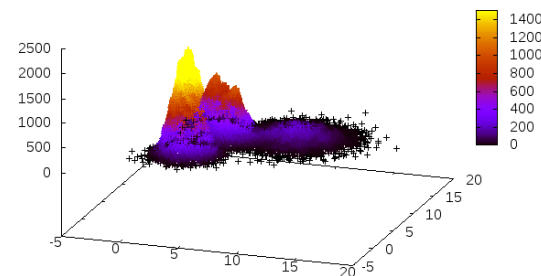
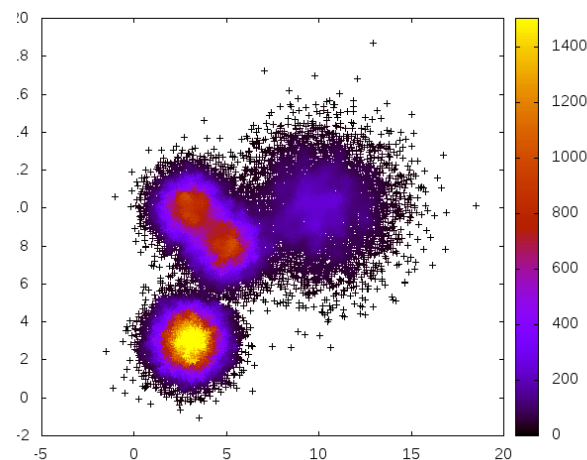
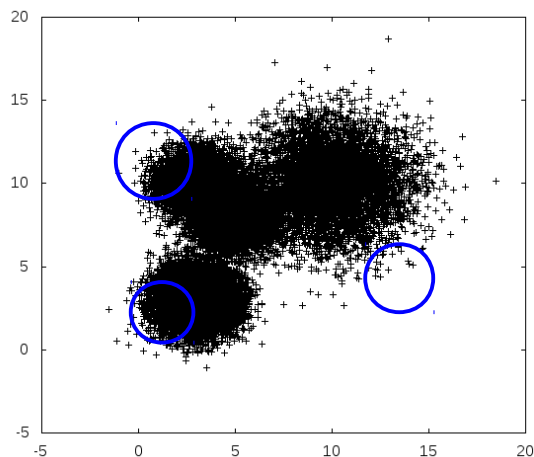
A Rodriguez, A Laio, Science 344, 1492 (2014)



Find modes (peaks) of a density distribution

Reconstruct density around each point with ε -ball counting:

$$\rho(x_i) = \sum_j \chi(j \in B_\varepsilon(x_i))$$



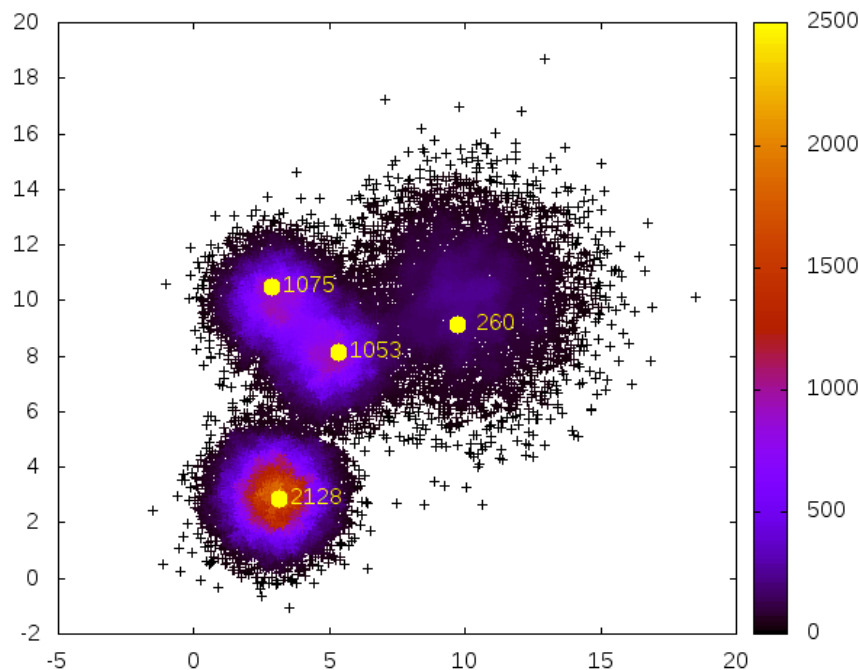
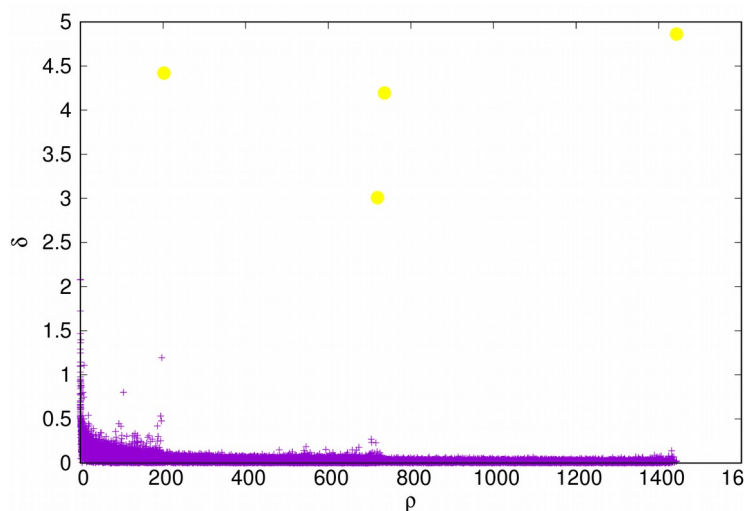
Density Peak Clustering

ρ maxima are far from points with lower ρ

Compute minimum distance from point at higher ρ

$$\delta_i = \min_{j: \rho_j > \rho_i} d_{ij}$$

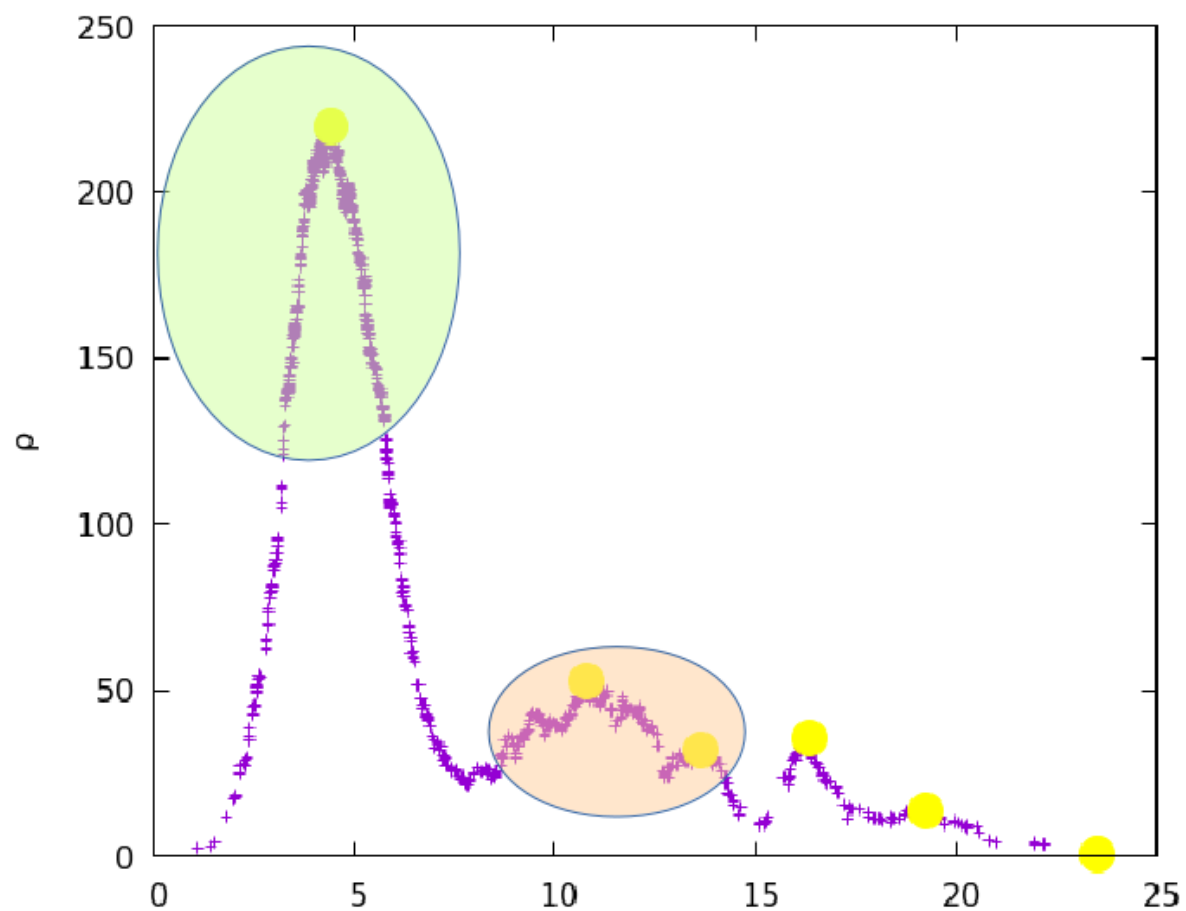
Peak are outliers in *decision graph* ρ_i vs δ_i



Validating DPC

“Significance” of the peaks?

Peaks or density fluctuations?

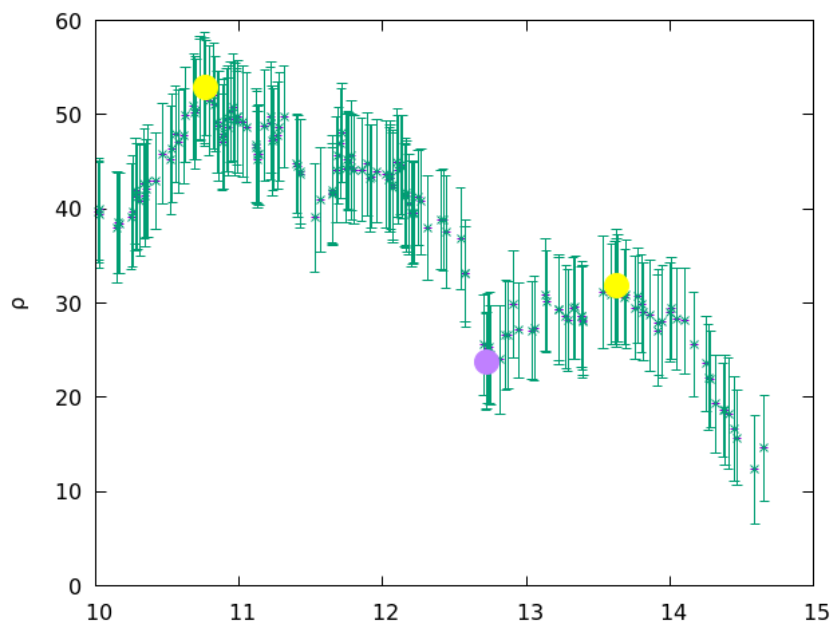
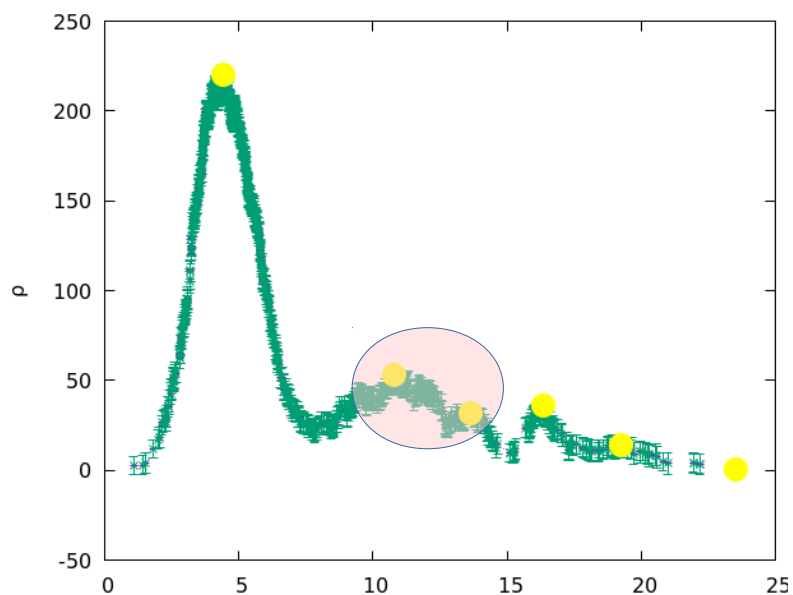


Validating DPC

- Find **peaks and saddle points**

- Compute **error $\Delta\rho$**

- Significance: $\rho_{peak} - \rho_{saddle} \geq z(\sqrt{(\Delta\rho_{peak})^2 + (\Delta\rho_{saddle})^2})$





A local model for the data

No global model for the data

Only two **broad assumptions**:

H1) the data points x_i are **independent samples** from a density $\rho(x)$.

H2) local uniformity: for all x_i , there exists (small) k such that $\rho(x) \sim \text{const.}$ in the region containing the first k neighbors of x_i



local model for the distribution of neighbor distances around each point

points within a small region around each point follows **Poisson process**
with **parametric dependence on ρ**

$$\text{prob}(\mathcal{N}(\mathcal{A}) = n) = \frac{\rho(x_i)^n \text{vol}(\mathcal{A})^n}{n!} e^{-\rho(x_i) \text{vol}(\mathcal{A})}$$

Estimating the density

k nearest neighbors of i at distances $r_{i1}, r_{i2}, r_{i3}, \dots$

hyperspherical shells S_j enclosed between neighbors

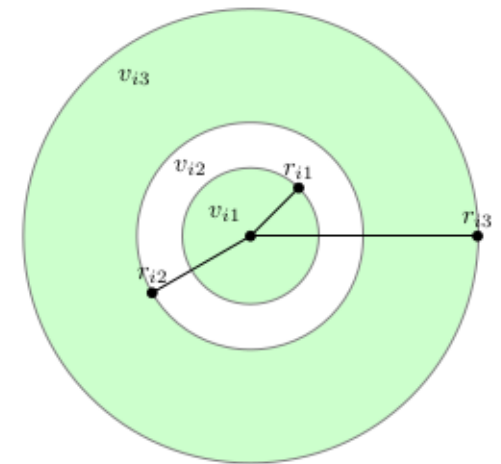
volumes $v_j = \omega_d r_{ij}^d - \omega_d r_{i(j-1)}^d$

distribution of shell volumes V_j follows from Poisson process

$$\text{prob}(\mathcal{N}(s_j) = 0) = e^{-\rho(x_i)v_j} = \text{prob}(V_j > v_j)$$

$$\mathcal{L}(V_j = v_j) = \rho(x_i)e^{-\rho(x_i)v_j}$$

- Considering all volumes $\mathcal{L}(\{V_j = v_j\}) = \rho(x_i)^k e^{-\rho(x_i)\text{vol}(B_{r_{ik}}(x_i))}$
- **Local model of NN distances depending on ρ**



Estimating the density

By max likelihood estimate ρ and error $\Delta\rho$

$$\rho(x_i) = \frac{k}{\omega_d r_{ik}^d}, \quad \Delta\rho(x_i) = \frac{\sqrt{k}}{\omega_d r_{ik}^d}$$

Two problems:

- 1) what is right k ?
- 2) what is right d ?

What is right k ?

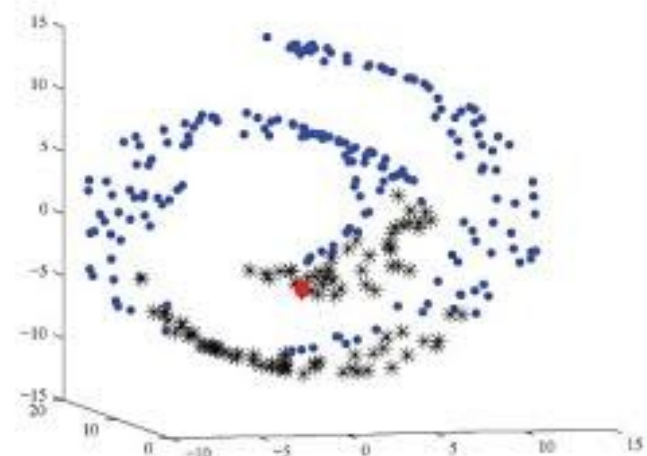
increase k until local model fails

[Rodriguez et al., JCTC 2017]

What is right d ?

Intrinsic dimension

- The data lie on d -dimensional hypersurface
- ρ should be evaluated on this hypersurface





ID estimation: TWO-NN

E Facco, M D'Errico, A Rodriguez, A Laio, Scientific Reports 7, 12140 (2017)

local model for $k=2$

$$\nu = v_2/v_1$$

distribution of ν is independent of ρ

$$\mathcal{L}(\nu) = \frac{1}{1 + \nu^2}$$

$$\mu = r_{i2}/r_{i1}$$

$$\nu = \mu^d - 1$$

under local model, distribution of μ depends only on d

$$\mathcal{L}(\mu) = de^{-(d+1)\mu}$$

ID can be inferred from the **μ of all points** collectively

This is independent of the estimates of ρ
(assuming ρ is constant over scale of first 2 neighbors)



ID estimation: heterogeneous case

M Allegra, E Facco, A Laio and A Mira, in prep. (2018)

ID may not be uniform in the dataset!

H3) $\rho(x)$ has support on the union of a finite number K of manifolds

with different intrinsic dimensions $d_1 \dots d_K$

Mixture model
$$\rho(x) = \sum_{k=1}^K p_k \rho(x|k)$$

Under H1), H2) one can still predict the expected distribution of the μ

mixture of Pareto distributions
$$P(\mu_i) = \sum_{k=1}^K p_k d_k \mu_i^{-d_k-1}$$

The likelihood of the data is
$$\mathcal{L}(\boldsymbol{\mu}|\mathbf{d}, \mathbf{p}) = \prod_{i=1}^N \sum_{k=1}^K p_k d_k \mu_i^{-d_k-1}$$

Then we can again estimate
$$\mathbf{d} = d_1 \dots d_K, \quad \mathbf{p} = p_1 \dots p_K$$



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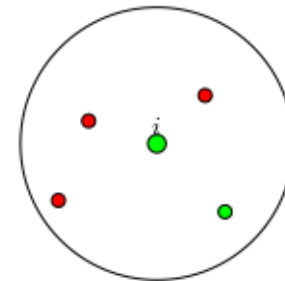
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ID estimation: heterogeneous case

- Introduce **latent variables** (manifold membership of each point) $\mathbf{Z} = Z_1, \dots, Z_N$
- Likelihood with latent variables $\mathcal{L}(\mu|\mathbf{d}, \mathbf{p}, \mathbf{Z}) = \prod_{i=1}^N p_{Z_i} d_{Z_i} \mu_i^{-d_{Z_i}-1}$
- Estimate jointly $\mathbf{d}, \mathbf{p}, \mathbf{Z}$
- K fixed by trying increasing values in $[1, K_{\max}]$ and performing a model selection test e.g. likelihood ratio test

Problem:

- Pareto distributions with different d are highly overlapping
- **estimation of manifold membership fails**
- diagnostic: **neighboring points have different Z**



Hidalgo

- **H4)** the first q neighbors of a point mostly belong to the same manifold

Probabilistic requirement on the Z of neighboring points

Z : be probability that a neighbour of i belong to same manifold as i

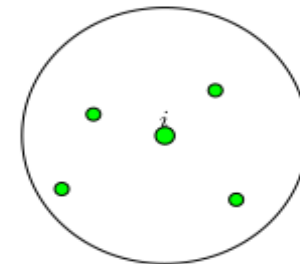
n_i^{in} # neighbors with same Z as i

n_i^{out} # neighbors with different Z

additional term in the likelihood

$$\mathcal{L}(n_i^{in} | \mathbf{Z}) = \frac{\zeta^{n_i^{in}} (1 - \zeta)^{n_i^{out}}}{Z}$$

$\zeta > \frac{1}{2}$ Controls the degree of uniformity



Two types of data analysis



Confirmatory analysis

- Starts from assumed *model* for the data, given a priori
- Uses *statistics* to verify whether the data fit the assumed model
- Can be rigid: fail to exploit richness of the data



Exploratory analysis

- Procedures (algorithms) to find structure in the data
- Often, *no formal evaluation* of the results
- Danger of falling into magical thinking (seeing structures that are not there)



A possible compromise



- We started with E.D.A. method (density peak clustering) with no statistical validation of results
- for statistical validation, some assumption on the data was needed
- we introduced *minimal assumptions* on the data, allowing to maintain high flexibility
- as a result, we developed complex procedure to reconstruct probability density, its intrinsic dimension and peaks in high dimensional space

Acknowledgments



Alessandro Laio



Maria d'Errico



Elena Facco



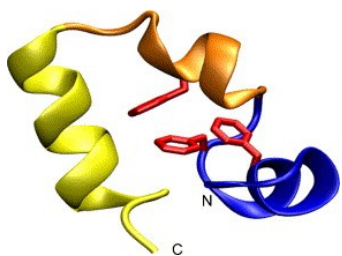
Alex Rodriguez



Antonietta Mira

Hidalgo

Example: molecular dynamics



- consider a MD of unfolding/refolding villing headpiece
- for each of the $N \sim 32000$ configurations, $D=32$ dihedral angles.

We find four manifolds

$d=12$

$d=13$

$d=13$

$d=23$

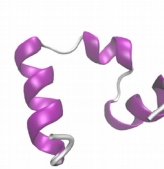
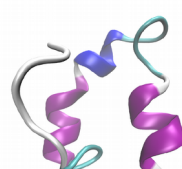
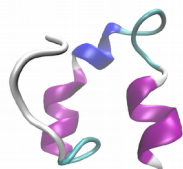
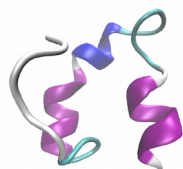
$Q=0.53$

$Q=0.58$

$Q=0.64$

$Q=0.89$

Fraction of native contacts



The folded state is recognized from its higher ID!