

# Chapter 7 Manifold forests

- Learning the structure of high-dimensional data & Mapping it onto a lower dimensional space
- Real data - large number of dimensions
- more direct visualization, easier data interpretation
- Properties of MF:
  - ① computational efficiency
  - ② automatic selection of discriminative features via information-based optimization
  - ③ other (i.e. code reusability)

## 7.1 Manifold Learning & Dimensionality Reduction in the lit

- ① Principal Component Analysis (PCA) - linear
- ② Isometric feature mapping

Laplacian eigenmaps - DR

→ try to preserve local pairwise point distances only

# Medical Image Analysis

## 7.2 Specializing the Forest Model for Manifold L

Given a set of  $k$  unlabeled observations

$$\{v_1, v_2, \dots, v_i, \dots, v_k\} \text{ with } v_i \in \mathbb{R}^d$$

We wish to find a smooth mapping

$$f: \mathbb{R}^d \rightarrow \mathbb{R}^{d'} \text{ with } f(v_i) = v'_i \text{ that approx.}$$

preserves the observations relative

geodesic distances, with  $d' \ll d$ .

- collection of clustering trees

- extra: a model of affinity bt. data points

### 7.2.1 The training objective function

$$\text{RNO} \quad \theta_j = \arg \max_{\theta \in \mathcal{T}_j} I(S_j, \theta)$$

$$I(S_j, \theta) = \log(|\Delta(S_j)|) - \sum \frac{|S_j^i|}{|S_j|} \log(|\Delta(S_j^i)|)$$

### 7.2.2 The Prediction Model

$$(7.3) \quad P_t(v) = \frac{\pi_{t(v)}}{\sum_i} N(v; \mu_t(v), \Delta_t(v))$$

## 7.2.3 The Affinity Model

→ important to estimate similarity / affinity / distance between data points

Partition of input points:

$$l(v) : \mathbb{R}^d \rightarrow \mathcal{C} \subset \mathbb{N}$$

Affinity matrix:  $W_{ij}^T = e^{-Q^T(v_i, v_j)}$

Distance  $Q$ :

① Mahalanobis affinity

② Gaussian affinity

③ Binary Affinity ← simplest & interesting

$$Q^T(v_i, v_j) = \begin{cases} 0 \\ \infty \end{cases}$$

## 7.2.4 The Ensemble Model

→ smoother affinity matrix

$$W = \frac{1}{T} \sum_{t=1}^T W^t \quad (7.9)$$

→ add robustness

## 7.2.5 Estimating the Embedding Function

→ Laplacian eigenmaps

$$\text{matrix: } L = I - \gamma^{-\frac{1}{2}} W \gamma^{-\frac{1}{2}} \quad (7.10)$$

$$v_i^l = f(v_i) = (\tilde{e}_{i1}, \dots, \tilde{e}_{ij}, \dots, \tilde{e}_{id})^T$$

## 7.2.6 Mapping Previously unseen points

→ interpolate the point position given the already available embedding

$$v^l = \frac{1}{T} \sum_t \frac{1}{\gamma t} \sum_i (e^{-Qt(v, v_i)} f(v_i))$$

## 7.2.7 Properties and Advantages

(1) Ensemble Clustering for Distance Estimation

complex data → use pairwise affi. defined by tree structure

(2) Choosing the Feature Space

→ automatically select discriminative features and corresponding parameters.

### (3) Computational Efficiency

①  $d' \ll k$  bottom eigenvectors

②  $L$  is very sparse

### (4) Estimating the Target Intrinsic Dimensionality

Choose value of  $d'$ :

① profile of eigenvalues  $\lambda_j$

② minimum number of eigenvalues

## 7.3 Experiments & Effect of Model Parameters

### (1) Forest size

Large  $T \rightarrow$  positive effects (robustness  
to noise)

better approximate pairwise graph affinity

### (2) Higher Dimensions

### (3) Discovering the Manifold Intrinsic Dimensionality

small value of  $d'$

Sharper elbow:

→ higher values at  $T$

→ Gaussian affinities

