Spectral Clustering of Time Series Data

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Overview

Project Goals

- Clustering time series data that contain smoothly varying HMM parameters
- Background on HMM & EM
- Current approaches to HMM clustering
- Background on spectral clustering
- The spectral clustering approach to HMM clustering
- Data used
- Current test results
- Conclusions

Project Goals

- Unsupervised learning is a well studied problem
- Typically applied to data in vectorized form
- What about variable length data? Such as time series
- Our goal:
 - Investigate time-series clustering using spectral clustering with probability product kernels
 - □ Compare with existing methods of time-series clustering

Background

HMM clustering using EM

- \Box Train a mixture of *k* HMMs on the data
- Calculate posteriors from likelihoods of each sequence with each HMM in Estep
- Update the parameters of each HMM with the posteriors in M-step
- Each sequence influences the parameters of each HMM by its posterior probability under that HMM

CONTINUE

$$\bar{\pi}_{i}^{(m)} = \frac{\sum_{l=1}^{L} z_{lm} \gamma_{1}^{(lm)}(i)}{\sum_{l=1}^{L} z_{lm}}$$
e
its
at HMM $\bar{a}_{ij}^{(m)} = \frac{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T-1} \xi_{t}^{(lm)}(i,j)}{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T-1} \gamma_{t}^{(lm)}(i)}$
 $\bar{\mu}_{j} = \frac{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T} \gamma_{t}^{(lm)}(i) \cdot O_{t}}{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T} \gamma_{t}^{(lm)}(i)}$
 $\bar{\Sigma}_{j} = \frac{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T} \gamma_{t}^{(lm)}(i) \cdot (O_{t} - \mu_{j})(O_{t} - \mu_{j})'}{\sum_{l=1}^{L} z_{lm} \sum_{t=1}^{T} \gamma_{t}^{(lm)}(i)}$

Background

- Spectral Clustering
 - Calculate the Gram matrix of dataset
 - □ Calculate the normalized Laplacian matrix
 - □ Find the eigenvectors of the Laplacian matrix
 - Run clustering algorithm on the components of the eigenvectors
- Probability Product Kernels
 - Calculate the probability distribution of the HMM over a dataset
 - The kernel affinity is the similarity between two distributions
 - □ Bhattacharya affinity



Spectral Clustering of HMMs

- Train single HMMs over each data sequence
- Find probability of each data sequence over each HMM
- The elements of the gram matrix are the probability product kernel affinities
- Find eigenvectors of the normalized gram matrix
- Run K-means on the components of the eigenvectors

Data

Rotated MOCAP Data – 360 Degrees



Results

- Testing results over MOCAP data Accuracy rates
- Sub-sampled at 5° degree step sizes

	EM-HMMs	Spectral Clustering
Non-rotated MOCAP	100%	100%
Rotated 0 - 45°	100%	100%
Rotated 0 - 90°	~50%	100%
Rotated 0 -120°	~50%	100%
Rotated 0 -150°	~50%	50%
Rotated 0 -180°	~66%	~50%
Rotated 0 -360°	~50%	~50%

- In our testing, spectral clustering was roughly twice as fast as EM-HMM
- Accuracy results were influenced by what we believe to be an Intersecting manifold in HMM parameter space

Advantages

Speed

- □ Like EM-HMM except with only one iteration
- Can control the number of samples used to find the distribution, reduce it to boost speed
- Accuracy
 - On slowly varying data with non-gaussian HMM parameter distributions, the spectral clustering approach outperforms EM-HMM
 - EM-HMM can get stuck on local minima
- Eigengap
 - Can estimate the number of clusters by looking at the eigengap!
- Local Max
 - EM is susceptible to low max, spectral clustering is not



Conclusions

- Spectral clustering works well with slowing varying data. Can handle cases where the HMM parameters define some bendy non-Gaussian manifold.
- Results for time series clustering using spectral clustering is analogous to the results found in clustering non-time series data.

Currently working on

- □ Testing on more real world data (MOCAP, GAIT)
- Testing on multi-class data
- □ Exploring better ways to find K based on the eigengap
- □ Exploring better ways to improve speed