

Clustering-based Gaussian Mixture Reduction

Dennis Schieferdecker - 09.06.09



KIT - The cooperation of Forschungszentrum Karlsruhe GmbH and Universität Karlsruhe (TH)

in der Helmholtz-Gemeinschaft



Gaussian Mixtures

Gaussian Mixture Density

weighted sum of Gaussians

 $f(x;\underline{\eta}) = \sum_{i=1}^{N} \omega_i \cdot \mathcal{N}(x;\mu_i,\sigma_i^2)$

- universal function approximator
- possible applications
 - target tracking,
 - density estimation,
 - ...

Problems in Application

- recursive multiplication of Gaussian mixtures
- number of components grows rapidely (exponential growth)

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Problem Description



Gaussian Mixture Reduction

Goal

- find a mixture η with K < N components (reduced mixture),
- so that a deviation measure $d(\tilde{\eta}, \eta)$ is minimized.

Deviation Measures

- Integrated Squared Distance (ISD): $d(f_1(x), f_2(x)) = \int_{\mathbb{R}} (f_1(x) - f_2(x))^2 dx$
- Kullback-Leibler divergence (KLD): $d(f_1(x), f_2(x)) = \int_{\mathbb{R}} f_1(x) \log \frac{f_1(x)}{f_2(x)} dx$
- normalized variants







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Overview

top-down approaches

- greedy methods
- iteratively replace two Gaussians with one
- chosen according to a deviation measure (local, global, hybrid)

- constructive method
- starts with one Gaussian
- adds components as neccessary
- progressive approximation
 - PGMR





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Clustering Method

Overview

Gaussian Mixture Reduction via Clustering (GMRC)

- modular three-step algorithm
- input:
 - $\tilde{\eta}$ (parameter vector of the original mixture)
 - \overline{K} (number of reduced components)

output:

 η (parameter vector of the reduced mixture)



Conception

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each component <u>n</u> is interpreted as point (site) in a space with an underlying deviation measure







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- compute a preliminary solution $\underline{\eta}$ (i.e. using West, Runnalls, ... \rightarrow initial cluster centers
- associate each original component (site) <u>n</u> with the nearest component of the reduced mixture
- replace each cluster center with a new one, retaining mean and variance of the associated sites





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Clustering Step

- greedy approach
- based on Lloyd's algorihm (k-means algorithm):
 - associate each site $\tilde{\eta}_i$ with the 'nearest' center η
 - recompute centers according to the current association
 - repeat until the deviation no longer changes or is good enough

determine the 'nearest' center

- associate site $\tilde{\eta}_i$ with each center η_i
- temporarily update the affected centers
- compute change in deviation between updated reduced and original mixture (ISD)
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Refinement Step

Parameter Optimization

- optimize parameter vector $\underline{\eta}$ w.r.t. ISD $\min_{\underline{\eta}} \int_{\mathbb{R}} \left(\tilde{f}(x; \underline{\tilde{\eta}}) - f(x; \underline{\eta}) \right)^2 dx$
- non-linear optimization problem \rightarrow Newton approach

finds local optimum

Weight Optimization

- system of linear equations
- finds global optimum

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Simulation Setup

- Office PC (Intel Core2 Duo E8400)
- OpenSUSE 11.0
- Matlab 7.7.0 (R2008b)

- reduction of mixtures with $N \in \{40, 120, 200, 500, 1000\}$ components down to K = 10
- each evaluated with 1 000 simulation runs





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Results Approximation Quality





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algorithm		running time	norm. ISD
GMRC	complete	$\boxed{\textbf{2.793}\pm \textbf{0.052s}}$	$\overline{0.658\pm0.494}$
	w. random init. w/o clustering w/o refinement		$\begin{array}{c} \hline 1.272 \pm 1.561 \\ 0.774 \pm 0.872 \\ 1.697 \pm 0.432 \end{array}$
Runnalls		$1.678\pm0.024s$	$\textbf{3.606} \pm \textbf{0.752}$

(initialization with Runnalls' algorithm; N = 200, K = 10)









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clustering step primarily improves variance

refinement has single-most impact on approximation quality

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Visualization



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- novel top-down, global reduction algorithm
- competitive w.r.t. current state-of-the-art (PGMR)
- combines algorithmic and numerical ideas

Outlook

- extension to multivariate Gaussian mixtures
- impact of different clustering methods and deviation measures
- adaptive reduction of components







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Thank you for your attention!





time for questions

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