



**Aalto University**  
School of Science

# Mixture Models from Multiresolution data

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# Management Summary

- ▶ Motivation for the Work
- ▶ Multiresolution Data
- ▶ Algorithms and Experiments
- ▶ Summary and Conclusions

# Modelling : the general perspectives

“ *Modeling is like vintage wine; it matures with time.* ”

— DECISIONCRAFT INC.  
*[www.decisioncraft.com](http://www.decisioncraft.com)*

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“ *People may mature with time but models mature only with increasing data.* ”

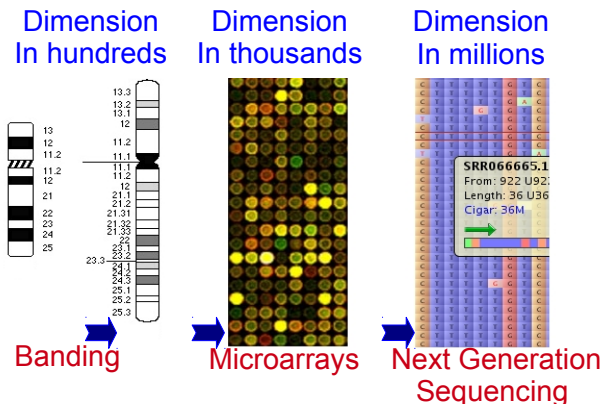
— PREM RAJ ADHIKARI  
*PhD Student*

# Importance of Using More Samples

## The Square Root Law

Accuracy of Information =  $\sqrt{\text{Volume of Information}}$

# The Multiresolution data



- Multiresolution data is everywhere: biology, computer vision, telecoms ...
- Older Generation Technology  $\Rightarrow$  Data in Coarse Resolution
- Newer Generation Technology  $\Rightarrow$  Data in Fine Resolution

# Mixture Modeling of Multiresolution 0–1 Data

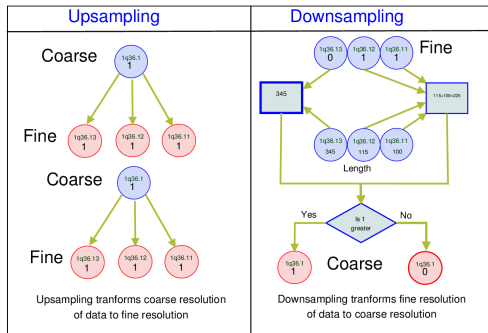
- ▶ Why Mixture Models?
- ▶ Cancer is a heterogeneous collection of several diseases and mixture models are well known for their ability to model heterogeneity
- ▶ Mixture models generally cannot model multiresolution data

$$P(x) = \sum_{j=1}^J \pi_j P(x|\theta_j) = \sum_{j=1}^J \pi_j \prod_{i=1}^d \theta_{ji}^{x_i} (1 - \theta_{ji})^{1-x_i}$$

↓  
i is different for each resolution and requires different models for each resolution

- ▶ Only mixture modeling solution to multiresolution data is to model each resolution separately

# Data transformation for multiresolution modelling



We upsample and downsample the data and integrate the data in same resolution before mixture modelling

P. R. Adhikari, J. Hollmén, UP'2010  
Patterns from Multiresolution 0-1 data



# Results of Mixture Models

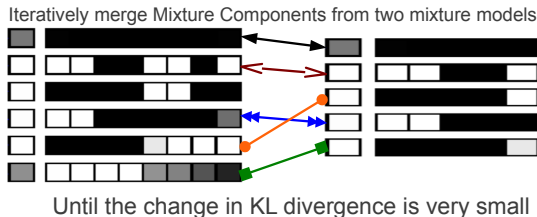
<b>Data Resolution</b>	<b>J</b>	<b>Likelihood</b>
Original in Coarse	8	-3.39
Original in Fine	8	-4.75
Downsampled to Coarse	6	-3.41
Upsampled to Fine	6	-5.23
Combined in Coarse	7	-3.36
Combined in Fine	7	-5.11

**Table :** Results of experiments on chromosome-17. J denotes the selected number of component distributions.

P. R. Adhikari, J. Hollmén, UP'2010  
**Patterns from Multiresolution 0-1 data**

# Multiresolution Mixture Modelling by Merging of Mixture Components

- What is done?



- How is it done?
- Fast approximation of KL divergence (P. R. Adhikari, J. Hollmén, DS 2012)

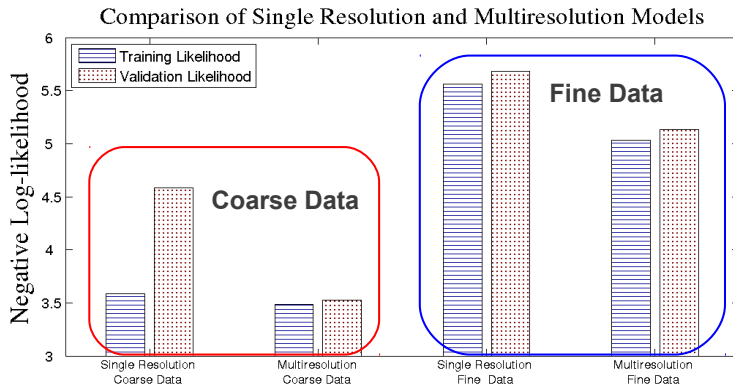
$$KL = \sum_{i \in X^*} \pi_{\alpha} \prod_{m=1}^d \left( \alpha_m^{X_{im}^*} (1 - \alpha_m)^{(1 - X_{im}^*)} \right) - \sum_{i \in Y^*} \pi_{\beta} \prod_{n=1}^{d'} \left( \beta_n^{Y_{in}^*} (1 - \beta_n)^{(1 - Y_{in}^*)} \right)$$

- Retrain the mixture models in different resolutions

P. R. Adhikari, J. Hollmén, ACML'2012

**Multiresolution Mixture Modeling using Merging of Mixture Components**

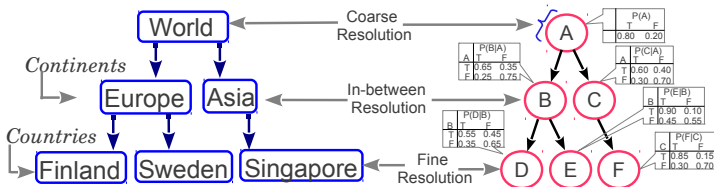
# Performance of Multiresolution Mixture Model



Better generalization through merging of mixture components

P. R. Adhikari, J. Holmén, ACML'2012  
Multiresolution Mixture Modeling using Merging of Mixture Components

# Multiresolution Mixture Components

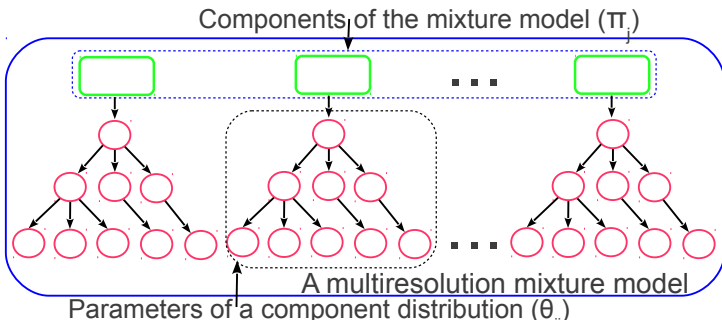


- ▶ Domain ontology provides information about relationships between features in different resolutions
- ▶ We can create a tree structure where features in the coarse resolution form the root and features in the fine resolution leaves of the tree

P. R. Adhikari, J. Hollmén, DS'2013

**Mixture Models from Multiresolution 0-1 Data.**

# Structure of Mixture Model

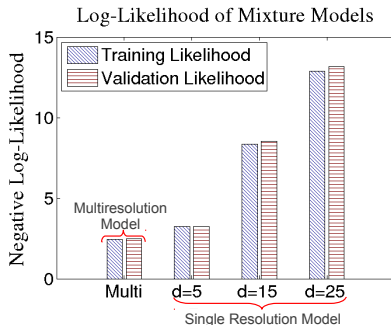


- ▶ The components of the mixture model are Bayesian networks themselves
- ▶ Now, the problem is to learn the parameters  $\Theta = \{J, \{\pi_j, \theta_j\}_{j=1}^J\}$

P. R. Adhikari, J. Hollmén, DS'2013

**Mixture Models from Multiresolution 0-1 Data.**

# Multiresolution Mixture Model Results



- ▶ The Y-axis shows the negative log likelihood, therefore, the shorter the bar, better the result
- ▶ The multiresolution model outperforms single resolution models

P. R. Adhikari, J. Hollmén, DS'2013  
**Mixture Models from Multiresolution 0-1 Data.**

# Summary and Conclusions

- ▶ Mixture Modeling of Multiresolution 0–1 Data in three ways:
  - ▶ Data Transformation
  - ▶ Merging of mixture components
  - ▶ Bayesian network as component distributions
- ▶ Experiments on multiresolution chromosomal datasets
- ▶ Experiments show that multiresolution models outperform single resolution models

# Questions, Comments, Feedback and Acknowledgment



HELSINKI  
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ALCODAN  
Algorithmic Data Analysis

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