

Abnormal Event Detection Based on Trajectory Clustering

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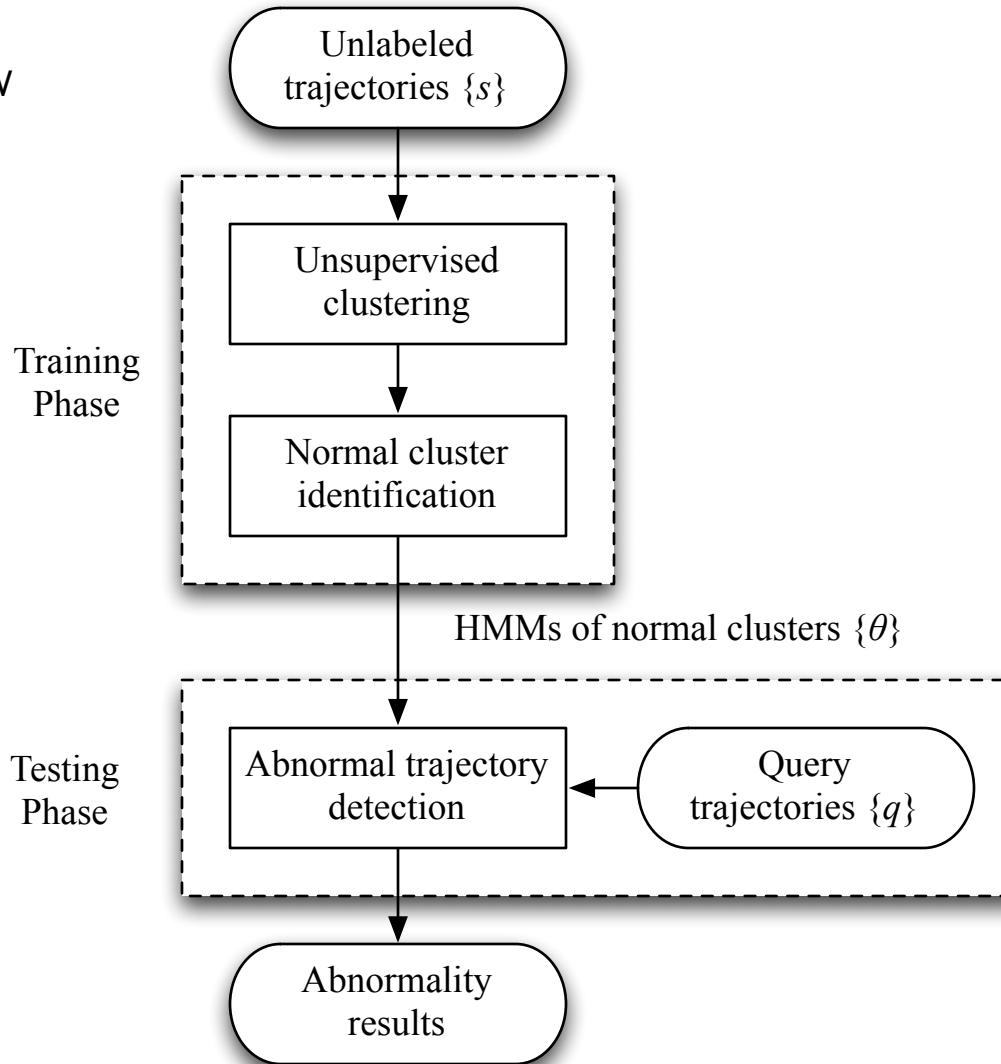
Introduction

- Suspicious / abnormal events in surveillance scene
- Trajectory-based representation



Clustering-Based Approach

System Overview



Dissimilarity Measure

- Bayesian information criterion (BIC)

$$BIC = -\log L + \frac{1}{2}K \log N,$$

- For N trajectories:

$$BIC(i, j, \dots) = -\sum_{n=1}^N \log L_n + \frac{1}{2}NK_0 \log N,$$

- Merge i and j

$$BIC(ij, \dots) = -\sum_{\substack{n=1 \\ n \neq i,j}}^N \log L_n - \log L_{ij} + \frac{1}{2}(N-1)K_0 \log N,$$

- Information-based dissimilarity

$$\begin{aligned} d(i, j) &= BIC(ij, \dots) - BIC(i, j, \dots) \\ &= \log L_i + \log L_j - \log L_{ij} - \frac{1}{2}K_0 \log N. \quad (5) \end{aligned}$$

Trajectory Clustering

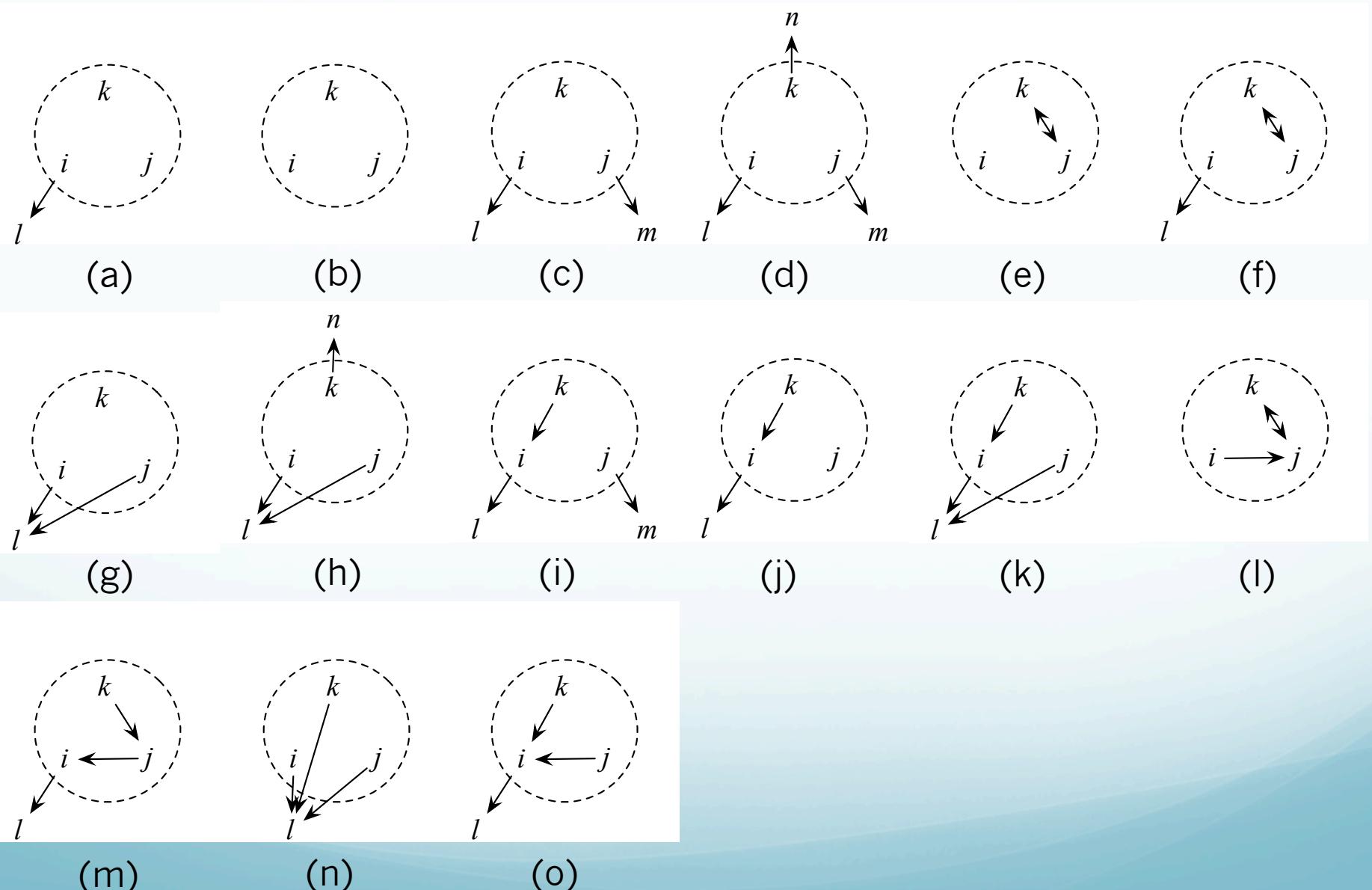
- Dynamic Hierarchical Clustering (DHC)

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- 0) *Initialization*: each trajectory in the dataset forms a group and is fitted with an HMM. There are N groups with N HMMs;
 - 1) *Dissimilarity Measurement*: calculate $d(i, j)$ of any two groups i and j in the dataset by (5);
 - 2) *Merging*: the two groups \hat{i} and \hat{j} with smallest $d(\hat{i}, \hat{j})$ ($d(\hat{i}, \hat{j}) < 0$) are merged; if no $d(i, j) < 0$, the clustering terminates;
 - 3) *Reclassification*: a new HMM $\theta_{\hat{i}\hat{j}}$ is trained, replacing $\theta_{\hat{i}}$ and $\theta_{\hat{j}}$; then based on the $(N - 1)$ HMMs, all trajectories are reclassified into $(N - 1)$ groups by the maximum likelihood (ML) criterion;
 - 4) *Retraining*: $(N - 1)$ HMMs are retrained based on the updated $(N - 1)$ data groups respectively;
 - 5) *Update*: $N = N - 1$; go back to step 1).
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2-Depth Search Strategy

- At each merging step, consider all pairs and all triplets:
 - Make two merges that decrease BIC the most
- However, need to calculate all trajectory pairs ($d(i,j)$), all trajectory triplets ($d(i,j,k)$)
 - Huge computation!

15 Categories of trajectory groups according to nearest neighbors



Branch Pruning

- Given pairwise measurement, the triple measurement can be possibly avoided
- Example: for category (f), merging of i,j,k together can be rejected if
 - $BIC(ij, k, l, \dots) > BIC(il, jk, \dots)$
 - Sufficient condition:

$$d(i, j) - d(j, k) + d(i, k) - d(i, l) > \frac{1}{2}K_0 \log N.$$

Proof of Sufficient Condition for (f)

$$\begin{aligned} d(i, j) - d(j, k) + d(i, k) - 2d(i, l) &> \frac{1}{2}K_0 \log N \\ \implies (\log L_i + \log L_j - \log L_{ij} - \frac{1}{2}K_0 \log N) \\ &\quad - (\log L_j + \log L_k - \log L_{jk} - \frac{1}{2}K_0 \log N) \\ &\quad + (\log L_i + \log L_k - \log L_{ik} - \frac{1}{2}K_0 \log N) \\ &\quad - 2(\log L_i + \log L_l - \log L_{il} - \frac{1}{2}K_0 \log N) \\ &\qquad\qquad\qquad > \frac{1}{2}K_0 \log N \\ \implies -\log L_{ij} - \log L_{jk} - \log L_{ik} \\ &\quad + 2\log L_{jk} + 2\log L_{il} - 2\log L_l > 0 \\ \implies -\log L_{ijk}^{ij} - \log L_{ijk}^{jk} - \log L_{ijk}^{ik} \\ &\quad + 2\log L_{jk} + 2\log L_{il} - 2\log L_l > 0 \\ \implies -\log(L_{ijk})^2 + 2\log L_{jk} + 2\log L_{il} - 2\log L_l &> 0 \\ \implies -\log L_{ijk} - \log L_l + \log L_{il} + \log L_{jk} &> 0 \\ \implies BIC(ijk, l, \dots) - BIC(il, jk, \dots) &> 0 \\ \implies BIC(ijk, l, \dots) &> BIC(il, jk, \dots) \end{aligned}$$

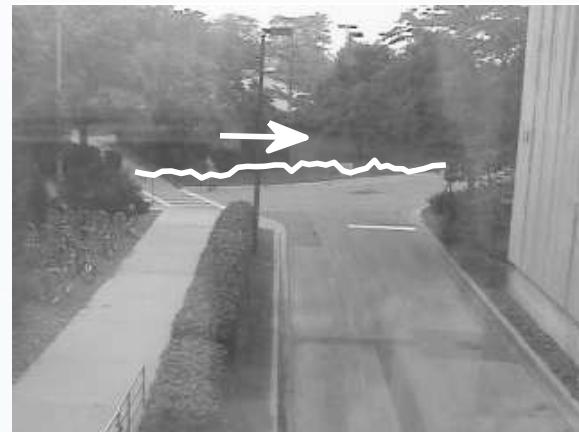
Category	Exclusions	Sufficient conditions
(a)	$BIC(ijk, \dots) > BIC(i, j, k, \dots)$	$d(i, j) + d(j, k) + d(i, k) > \frac{1}{2}K_0 \log N$
(b)	$BIC(ijk, l, \dots) > BIC(il, j, k, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$
(c)	$BIC(ijk, l, m, \dots) > BIC(il, jm, k, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) - 2d(j, m) > \frac{1}{2}K_0 \log N$
(d)	$BIC(ijk, l, m, n, \dots) > BIC(il, jm, kn, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) - 2d(j, m) - 2d(k, n) > \frac{1}{2}K_0 \log N$
(e)	$BIC(ijk, \dots) > BIC(i, jk, \dots)$	$d(i, j) - d(j, k) + d(i, k) > \frac{1}{2}K_0 \log N$
(f)	$BIC(ijk, l, \dots) > BIC(il, jk, \dots)$	$d(i, j) - d(j, k) + d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$
(g)	$BIC(ijk, l, \dots) > BIC(il, j, k, \dots)$ or $BIC(ijk, l, \dots) > BIC(i, jl, k, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$ or $d(i, j) + d(j, k) + d(i, k) - 2d(j, l) > \frac{1}{2}K_0 \log N$
(h)	$BIC(ijk, l, n, \dots) > BIC(il, j, kn, \dots)$ or $BIC(ijk, l, n, \dots) > BIC(i, jl, kn, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) - 2d(k, n) > \frac{1}{2}K_0 \log N$ or $d(i, j) + d(j, k) + d(i, k) - 2d(j, l) - 2d(k, n) > \frac{1}{2}K_0 \log N$
(i)	$BIC(ijk, l, \dots) > BIC(il, j, k, \dots)$ or $BIC(ijk, \dots) > BIC(ik, j, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$ or $d(i, j) + d(j, k) - d(i, k) > \frac{1}{2}K_0 \log N$
(j)	$BIC(ijk, l, m, \dots) > BIC(il, jm, \dots)$ or $BIC(ijk, m, \dots) > BIC(ik, jm, \dots)$	$d(i, j) + d(j, k) + d(i, k) - 2d(i, l) - 2d(j, m) > \frac{1}{2}K_0 \log N$ or $d(i, j) + d(j, k) - d(i, k) - 2d(j, m) > \frac{1}{2}K_0 \log N$
(k)	$BIC(ijk, l, \dots) > BIC(ik, jl, \dots)$	$d(i, j) + d(j, k) - d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$
(l)	$BIC(ijk, \dots) > BIC(i, jk, \dots)$	$d(i, j) - d(j, k) + d(i, k) > \frac{1}{2}K_0 \log N$
(m)	$BIC(ijk, l, \dots) > BIC(il, jk, \dots)$	$d(i, j) - d(j, k) + d(i, k) - 2d(i, l) > \frac{1}{2}K_0 \log N$

2-Depth DHC

- 0) *Initialization*: each trajectory in the dataset forms a group and is fitted with an HMM. There are N groups with N HMMs;
 - 1) *Pairwise Group Measurement*: calculate $d(i, j)$ of any two groups i and j in the dataset by (5);
 - 2) *Triple Group Measurement*: calculate only $d(i, j, k)$ of those three groups i, j, k that do not satisfy the exclusions in Table I;
 - 3) *Merging*: merge two group pairs or three groups together, based on which results in the greatest decrease of BIC. The clustering process terminates if no merge can decrease BIC;
 - 4) *Reclassification*: new HMMs are trained for the merged groups; then based on the $(N - 2)$ HMMs, all trajectories are reclassified into $(N - 2)$ groups by the maximum likelihood (ML) criterion;
 - 5) *Retraining*: $(N - 2)$ HMMs are retrained based on the updated $(N - 2)$ data groups respectively;
 - 6) *Update*: $N = N - 2$; go back to step 1).
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Normal Trajectories

- Examples:



Abnormal Trajectories

- Examples:



Comparison Results

- Compare 4 methods:
 - SC: spectral clustering (using CLR-based dissimilarity measure)
 - HC1: classic hierarchical clustering (1-depth, no dynamics)
 - DHC1: dynamic hierarchical clustering (1-depth)
 - DHC2: dynamic hierarchical clustering (2-depth)

Detection Error Rates

