



# Hmm, HID HMMs

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# Overview

- The Problem
- HMM Background
- Binomial Field HMMs
- HMMs, a la Kale, et al.
- New Ideas





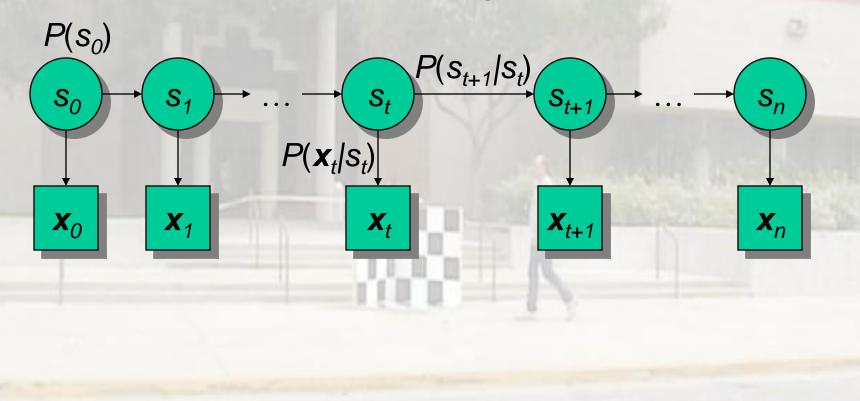
# The Problem

- Identify people
  - Video of profile views
  - Varying surface conditions, shoes, etc.
- Need for building a model robust to
  - Surface conditions, shoes, etc.
  - Local backgrounding deficiencies
    - Missing patches
    - Shadows
    - Noise



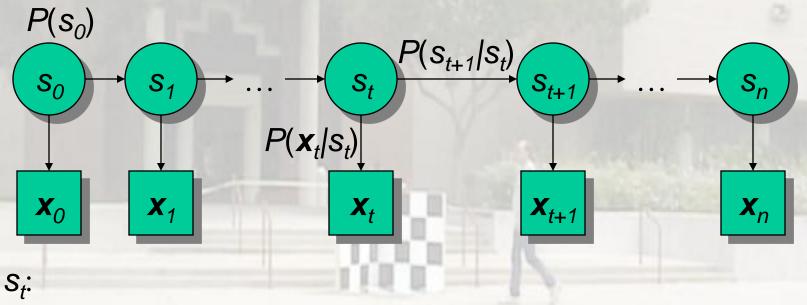








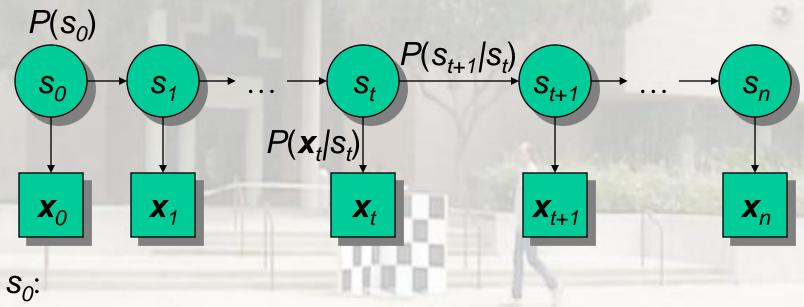




- What: State of the system at time t
- Example 1:  $s_9 = 0 \rightarrow$ The person has their legs together (state/phase 0) in frame 9
- Example 2:  $s_{14} = 4 \rightarrow$ The person in the widest stance (state/phase 4) in frame 14



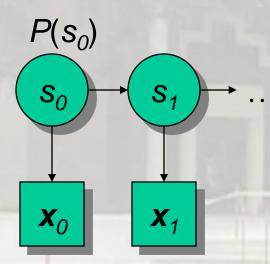




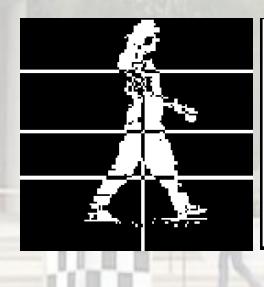
- What: Initial state
- Notes: Useful to model as a non-uniform random variable if you have some idea about how a person starts walking, relative to the first frame.

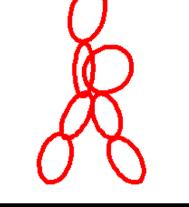






**X**<sub>t</sub>:





#### - What: Observation

- What you measure
- Must be describable in a generative, probabilistic framework
- **Example 1:** The silhouette in frame *t*
- Example 2: Lily's features from the 7 silhouette sections





# **Binomial Field HMMs**

Observation is a binary image

 Assume silhouette pixels produced independently, given the current state.

 $-P(\boldsymbol{x}_t|\boldsymbol{s}_t) = \prod_{u,v} P(\boldsymbol{x}_t(u,v) \mid \boldsymbol{s}_t)$ 

Model may be visualized as a grayscale image







# Some HMM Uses (one HMM per person)

- Make phase assignments
- Help build an appearance model to clean up silhouettes
  - E.g. turn on any pixels in a silhouette when that pixel almost always is on given the most likely state assignment
- Use directly for recognition
  - Determine the likelihood that each person's HMM would generate a test sequence of silhouettes
  - Select the person most likely to generate the sequence





# HMMs, a la Kale, et al.

- Thought process
  - Independence is a bad assumption
  - Not enough data to learn even covariances
  - So, do a dimensionality reduction...





#### HMMs, *a la* Kale, *et al.*: Dimensionality Reduction #1

• Calculate "width vectors"



 $66 \times 48 =$  3,168 dimensions

#### 66 dimensions







## HMMs, *a la* Kale, *et al.*: Dimensionality Reduction #1 (cont.)

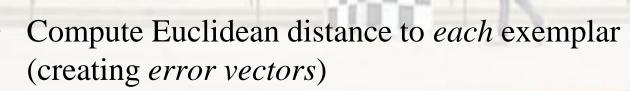
- General covariance estimation requirements
  - Assume our data has k dimensions
  - Covariance estimation involves  $\frac{k(k+1)}{2}$  unknowns
  - Each data point supplies k equations
  - Need  $\frac{k+1}{2}$  data points to avoid degeneracy
- In our case...
  - $k = 66 \rightarrow \left\lceil \frac{66+1}{2} \right\rceil = 34$  data points (frames) per phase
  - But, we only have  $\frac{200 \text{ frames}}{8 \text{ phases}} = 25 \text{ data points per phase}$





#### HMMs, a la Kale, et al.: Dimensionality Reduction #2

Choose phase exemplars
 (5 phases used in this case)





#### → [36.84 25.29 **20.99** 39.44 54.26]





### HMMs, a la Kale, et al.: Dimensionality Reduction #2 (cont.)

• Model error vectors as a joint Gaussian

 For 8 phases, have ~5 equations for each unknown in the covariance





## HMMs, a la Kale, et al.: Training

- Estimate the walking period
- Pick a set of equally-spaced frames from one period

- "...we use the 5 stances which lead to minimum error in the 5-d vector sequences. (in the sense of minimizing the norm)."

- Train the HMM
  - Update the mean and covariances of the error vectors
  - No updating of the exemplars...





## HMMs, a la Kale, et al.: Training (cont.)

• M-Step: Find  $\hat{\theta}_i$  where

$$\widehat{\theta}_i = \arg \max_{\theta_i} \sum_{l=1}^{L} \sum_{t=0}^{n_l} \gamma_t^{(l)}(i) \log P(\mathbf{x}_t^{(l)} | \theta_i)$$

 Unfortunately, this expands to something really ugly, where the nasty part includes

$$-\frac{1}{2} \left( \left( \begin{array}{c} \left\| \mathbf{x}_t - S_1 \right\| \\ \left\| \mathbf{x}_t - S_2 \right\| \\ \dots \\ \left\| \mathbf{x}_t - S_N \right\| \end{array} \right) - \mu_i \right)^T \mathbf{\Sigma}^{-1} \left( \left( \begin{array}{c} \left\| \left\| \mathbf{x}_t - S_1 \right\| \\ \left\| \left\| \mathbf{x}_t - S_2 \right\| \right\| \\ \dots \\ \left\| \left\| \mathbf{x}_t - S_N \right\| \right\| \end{array} \right) - \mu_i \right)$$





# New Ideas

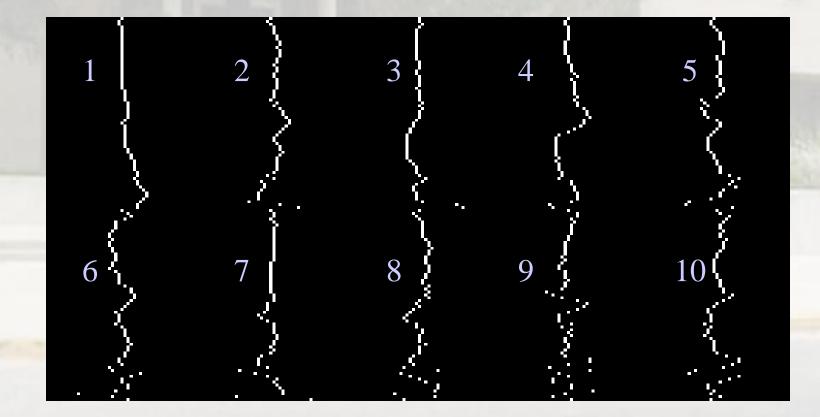
- Cannot easily update exemplars in Kale's approach, so...
- Use projections onto "exemplars" instead of distances from them
- Optimal "exemplars" are the PCA vectors





#### New Ideas:

## Top 10 Eigenvectors (Sequence 1)







#### New Ideas: How Many Eigenvectors?

1 dim, rms:21.1



4 dim, rms:16.1



7 dim, rms:9.6



2 dim, rms:20.1



5 dim, rms:13.7



8 dim, rms:9.3



3 dim, rms:16.2



Left side: Original width vector

**Right side:** Reconstruction after projection onto the *n*dimensional eigenspace

Note: Max possible RMS error is 48

6 dim, rms:13.0

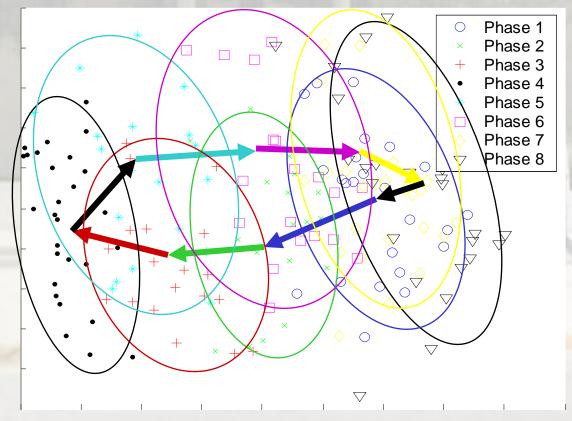


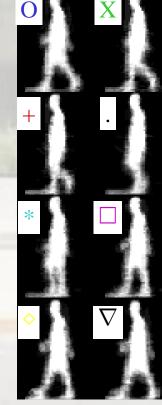
9 dim, rms:9.2





## New Ideas: Projection Onto 2D Subspace





Projection of each frame (phase determined by Binomial Field HMMs) BFHMM State Models