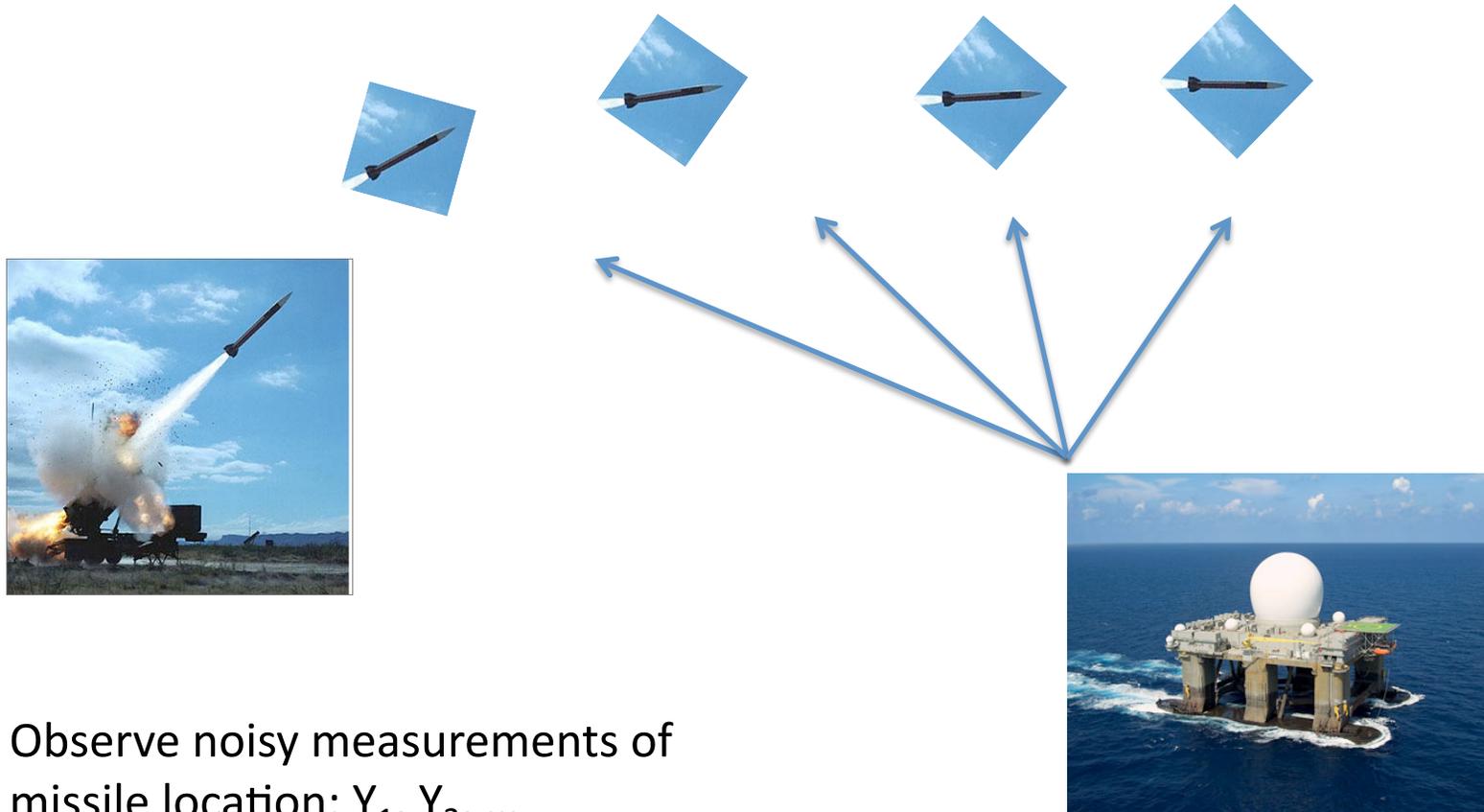


Hidden Markov models

Lecture 22

David Sontag
New York University

Example application: Tracking



Observe noisy measurements of
missile location: Y_1, Y_2, \dots

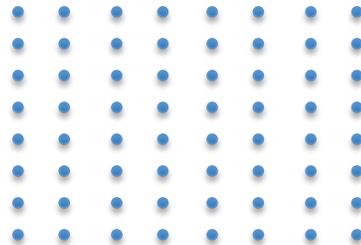
Radar

Where is the missile **now**? Where will it be in 10 seconds?

Probabilistic approach

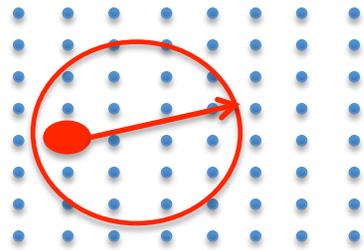
- Our measurements of the missile location were Y_1, Y_2, \dots, Y_n
- Let X_t be the *true* <missile location, velocity> at time t
- To keep this simple, suppose that everything is discrete, i.e. X_t takes the values $1, \dots, k$

Grid the space:



Probabilistic approach

- First, we specify the *conditional* distribution $\Pr(X_t | X_{t-1})$:



From basic physics, we can bound the distance that the missile can have traveled

- Then, we specify $\Pr(Y_t | X_t = \langle (10, 20), 200 \text{ mph toward the northeast} \rangle)$:

With probability $\frac{1}{2}$, $Y_t = X_t$ (ignoring the velocity). Otherwise, Y_t is a uniformly chosen grid location

Hidden Markov models

1960's

- Assume that the **joint** distribution on X_1, X_2, \dots, X_n and Y_1, Y_2, \dots, Y_n factors as follows:

$$\Pr(x_1, \dots, x_n, y_1, \dots, y_n) = \Pr(x_1) \Pr(y_1 | x_1) \prod_{t=2}^n \Pr(x_t | x_{t-1}) \Pr(y_t | x_t)$$

- To find out where the missile is *now*, we do **marginal inference**:

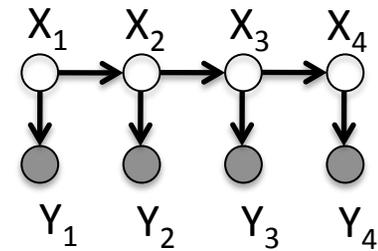
$$\Pr(x_n | y_1, \dots, y_n)$$

- To find the most likely *trajectory*, we do **MAP (maximum a posteriori) inference**:

$$\arg \max_{\mathbf{x}} \Pr(x_1, \dots, x_n | y_1, \dots, y_n)$$

Inference

- Recall, to find out where the missile is now, we do marginal inference: $\Pr(x_n \mid y_1, \dots, y_n)$



- How does one **compute** this?
- Applying rule of conditional probability, we have:

$$\Pr(x_n \mid y_1, \dots, y_n) = \frac{\Pr(x_n, y_1, \dots, y_n)}{\Pr(y_1, \dots, y_n)}$$

- Naively, would seem to require k^{n-1} summations,

$$\Pr(x_n, y_1, \dots, y_n) = \sum_{x_1, \dots, x_{n-1}} \Pr(x_1, \dots, x_n, y_1, \dots, y_n)$$

Is there a more efficient algorithm?

Marginal inference in HMMs

- Use **dynamic programming**

$$\begin{aligned}
 \Pr(x_n, y_1, \dots, y_n) &= \sum_{x_{n-1}} \Pr(x_{n-1}, x_n, y_1, \dots, y_n) && \Pr(A = a) = \sum_b \Pr(B = b, A = a) \\
 &= \sum_{x_{n-1}} \Pr(x_{n-1}, y_1, \dots, y_{n-1}) \Pr(x_n, y_n \mid x_{n-1}, y_1, \dots, y_{n-1}) && \Pr(\vec{A} = \vec{a}, \vec{B} = \vec{b}) = \Pr(\vec{A} = \vec{a}) \Pr(\vec{B} = \vec{b} \mid \vec{A} = \vec{a}) \\
 &= \sum_{x_{n-1}} \Pr(x_{n-1}, y_1, \dots, y_{n-1}) \Pr(x_n, y_n \mid x_{n-1}) && \text{Conditional independence in HMMs} \\
 &= \sum_{x_{n-1}} \Pr(x_{n-1}, y_1, \dots, y_{n-1}) \Pr(x_n \mid x_{n-1}) \Pr(y_n \mid x_n, x_{n-1}) && \Pr(A = a, B = b) = \Pr(A = a) \Pr(B = b \mid A = a) \\
 &= \sum_{x_{n-1}} \Pr(x_{n-1}, y_1, \dots, y_{n-1}) \Pr(x_n \mid x_{n-1}) \Pr(y_n \mid x_n) && \text{Conditional independence in HMMs}
 \end{aligned}$$

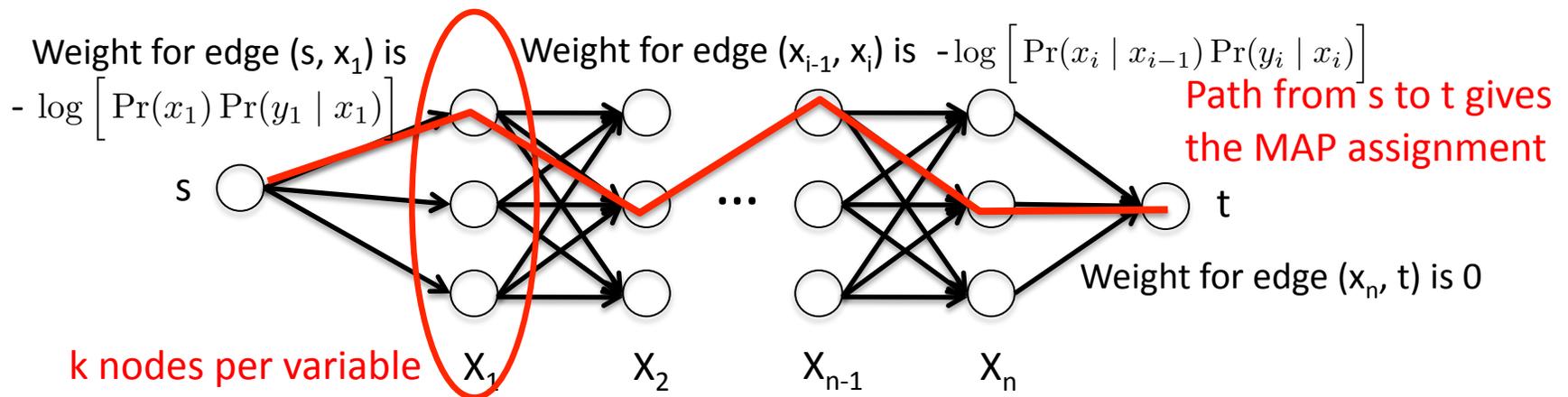
- For $n=1$, initialize $\Pr(x_1, y_1) = \Pr(x_1) \Pr(y_1 \mid x_1)$
- Total running time is $O(nk)$ – linear time! **Easy to do filtering**

MAP inference in HMMs

- MAP inference in HMMs can *also* be solved in linear time!

$$\begin{aligned} \arg \max_{\mathbf{x}} \Pr(x_1, \dots, x_n \mid y_1, \dots, y_n) &= \arg \max_{\mathbf{x}} \Pr(x_1, \dots, x_n, y_1, \dots, y_n) \\ &= \arg \max_{\mathbf{x}} \log \Pr(x_1, \dots, x_n, y_1, \dots, y_n) \\ &= \arg \max_{\mathbf{x}} \log \left[\Pr(x_1) \Pr(y_1 \mid x_1) \right] + \sum_{i=2}^n \log \left[\Pr(x_i \mid x_{i-1}) \Pr(y_i \mid x_i) \right] \end{aligned}$$

- Formulate as a shortest paths problem



Called the Viterbi algorithm

Applications of HMMs

- Speech recognition
 - Predict phonemes from the sounds forming words (i.e., the actual signals)
- Natural language processing
 - Predict parts of speech (verb, noun, determiner, etc.) from the words in a sentence
- Computational biology
 - Predict intron/exon regions from DNA
 - Predict protein structure from DNA (locally)
- And many many more!