

# Graphical Models and Hidden Markov Models

STAT5241 Section 2

Statistical Machine Learning

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#### Generative models are powerful

Conditional generative model P(zebra images | horse images)



Style Transfer



Input Image



Monet



Van Gogh

Zhou el al., Cycle GAN 2017



#### Generative models are powerful





# Generative models are powerful

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#### **Generative Models**

- An important "unsupervised learning" problem is learning distributions over input variables, also known as "generative models"
  - given such distributions, can do "probabilistic reasoning"
  - does not require "labels"; such unlabeled data is plentiful; big open problem in ML to get "good" generative models!
- Applications of generative models and probabilistic reasoning, especially for time series analysis:
  - Computer vision and graphics
  - Natural language processing
  - Information retrieval
  - Robotic control
  - Computational biology
  - Medical diagnosis
  - Finance and economics





#### However, generative models are...

- very specific (e.g. multivariate Gaussian distribution, or mixture of multivariate Gaussians) so that they might not be applicable to your given dataset ("data does not look Gaussian!")
- have a very large number of parameters (e.g. kernel density estimation)



#### **Graphical Models**

- Graphical models thread this needle: flexible, compact, and also interpretable (uses graphs, that are intuitive even to lay users)
- A marriage of probability theory (distributions among random variables) and graph theory (uses graphs to represent distributions)



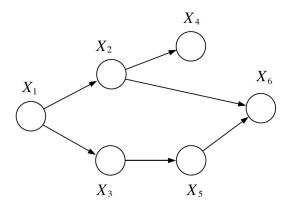
# **Graphical Models**

- Directed Graphical Models
- Undirected Graphical Models



# **Directed Graphical Models**

- Directed Graph is a pair G = (V, E), where V is a set of nodes, E is a set of directed (also called oriented) edges. We will assume G is acyclic.
- Each node  $i \in V$  is associated with a random variable  $X_i$ .
- Letting  $V = \{1, 2, ..., n\}$ , the set of random variables is  $\{X_1, X_2, ..., X_n\}$ .
- We will use node i and the associated random variable  $X_i$  interchangeably (though graph nodes and random variables are different formal objects!)

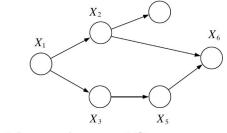




# **Directed Graphical Models**

• Each node  $i \in V$  has a set of parents  $\pi_i$ 

$$- \pi_6 = \{X_2, X_5\}.$$



• Let  $V = \{1, 2, \dots, n\}$ . Given a set of functions  $\{f_i(x_i, x_{\pi_i}) : i \in V\}$ , we define a joint probability distribution:

$$p(x_1,\ldots,x_n):=\prod_{i=1}^n f_i(x_i,x_{\pi_i}).$$
 Only assumption \* Non-negative \* Sum to one as

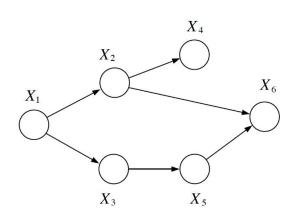
Only assumptions:

- \* Sum to one as a function of x i

- Is it a proper distribution?
  - Is it non-negative?
  - Does it sum to one? (Yes, provided each function  $f_i(x_i, x_{\pi_i})$  sums to one as a function of  $x_i$ .



# **Directed Graphical Models**





#### **Chain Rule**

$$p(x_1, x_2, x_3, x_4, x_5, x_6)$$

$$= p(x_1)p(x_2 \mid x_1)p(x_3 \mid x_1, x_2)p(x_4 \mid x_1, x_2, x_3)p(x_5 \mid x_1, x_2, x_3, x_4)p(x_6 \mid x_1, x_2, x_3, x_4, x_5)$$

In General:

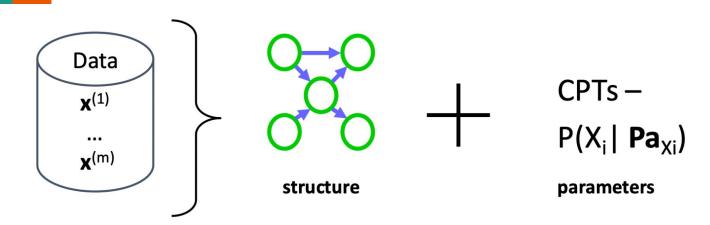
$$p(x_1, x_2, \dots, x_n) = \prod_{i=1}^n p(x_i \mid x_1, \dots, x_{i-1}).$$
 Holds for all distributions

Compare to:

$$p(x_1, x_2, \ldots, x_n) = \prod_{i=1}^n p(x_i \, | \, x_{\pi_i})$$
. Holds for directed graphical model distributions



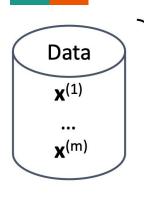
#### **Learning Directed Graphical Models**



Given set of m independent samples (assignments of random variables), find the best (most likely?) Bayes Net (graph Structure + CPTs)



# **Learning CPTs with Given Structure**



For each discrete variable X<sub>k</sub>

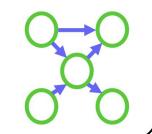
Compute MLE or MAP estimates for

$$p(x_k|pa_k)$$



MLE: 
$$P(X_i = x_i \mid X_j = x_j) = \frac{\text{Count}(X_i = x_i, X_j = x_j)}{\text{Count}(X_i = x_j)}$$

MAP: Add psuedocounts





# **Learning CPTs with Given Structure**

Given structure, log likelihood of data

onlyon

In structure, log likelihood of data 
$$\log P(\mathcal{D} \mid \theta_{\mathcal{G}}, \mathcal{G})$$

$$= \log \prod_{j=1}^{m} P(f^{(j)}) P(a^{(j)}) P(s^{(j)} \mid f^{(j)}, a^{(j)}) P(h^{(j)} \mid s^{(j)}) P(h^{(j)} \mid s^{(j)})$$

$$= \sum_{j=1}^{m} [\log P(f^{(j)}) + \log P(a^{(j)}) + \log P(a^{(j)} \mid f^{(j)}, a^{(j)}) + \log P(h^{(j)} \mid s^{(j)}) + \log P(h^{(j)} \mid s^{(j)})$$

$$= \sum_{j=1}^{m} \log P(f^{(j)}) + \sum_{j=1}^{m} \log P(a^{(j)}) + \sum_{j=1}^{m} \log P(a^{(j)} \mid f^{(j)}, a^{(j)}) + \sum_{j=1}^{m} \log P(h^{(j)} \mid s^{(j)})$$
Depends 
$$\theta_{\mathsf{F}} \qquad \theta_{\mathsf{A}} \qquad \theta_{\mathsf{F,A}} \sum_{j=1}^{m} \log P(h^{(j)} \mid s^{(j)}) + \sum_{j=1}^{m} \log P(h^{(j)} \mid s^{(j)})$$
Only on



# Information Theoretic Interpretation

$$\begin{split} \log P(\mathcal{D} \mid \theta_{\mathcal{G}}, \mathcal{G}) &= \sum_{j=1}^m \sum_{i=1}^n \log P\left(X_i = x_i^{(j)} \mid \mathbf{Pa}_{X_i} = \mathbf{x}_{\mathbf{Pa}_{X_i}}^{(j)}\right) \\ &= \sum_{i=1}^n \sum_{x_i} \sum_{\mathbf{x}_{\mathbf{Pa}_{X_i}}} \mathrm{count}(X_i = x_i, \mathbf{Pa}_{X_i} = \mathbf{x}_{\mathbf{Pa}_{X_i}}) \log P\left(X_i = x_i \mid \mathbf{Pa}_{X_i} = \mathbf{x}_{\mathbf{Pa}_{X_i}}\right) \end{split}$$

#### Plugging in MLE estimates: ML score



# Information Theoretic Interpretation

$$\begin{split} \log \widehat{P}(\mathcal{D} \mid \widehat{\theta}_{\mathcal{G}}, \mathcal{G}) &= m \sum_{i=1}^{n} \sum_{x_{i}} \sum_{\mathbf{x}_{\mathbf{Pa}_{X_{i}}}} \widehat{P}(x_{i}, \mathbf{x}_{\mathbf{Pa}_{X_{i}}}) \log \widehat{P}\left(x_{i} \mid \mathbf{x}_{\mathbf{Pa}_{X_{i}}}\right) \\ &= -m \sum_{i=1}^{n} \widehat{H}(X_{i} \mid \mathbf{Pa}_{X_{i}}) \\ &= m \sum_{i=1}^{n} \left[\widehat{I}(X_{i}, \mathbf{Pa}_{X_{i}}) - \widehat{H}(X_{i})\right] \\ &= \text{Doesn't depend on graph structure} \mathcal{G} \end{split}$$

ML score for graph structure  $\mathcal{G}$ 

$$\arg\max_{\mathcal{G}}\log\widehat{P}(\mathcal{D}\mid\widehat{\theta}_{\mathcal{G}},\mathcal{G}) \ = \arg\max_{\mathcal{G}}\sum_{i=1}^{n}\widehat{I}(X_{i},\mathbf{Pa}_{X_{i}})$$

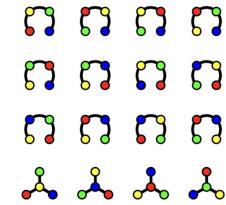


# What if graphical structure is not given?

# How many trees are there?

- Trees every node has at most one parent
- n<sup>n-2</sup> possible trees (Cayley's Theorem)



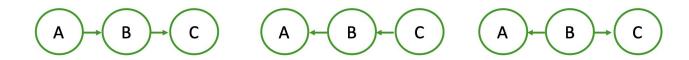




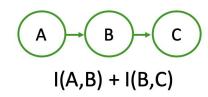
# **Focusing on Tree Structure**

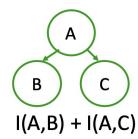
$$\arg\max_{\mathcal{G}}\log\widehat{P}(\mathcal{D}\mid\widehat{\theta}_{\mathcal{G}},\mathcal{G}) = \arg\max_{\mathcal{G}}\sum_{i=1}^{n}\widehat{I}(X_{i},\mathbf{Pa}_{X_{i}})$$

Equivalent Trees (same score): I(A,B) + I(B,C)



Score provides indication of structure:







#### **Chow-Liu Algorithm**

- For each pair of variables X<sub>i</sub>, X<sub>i</sub>
  - Compute empirical distribution:  $\widehat{P}(x_i, x_j) = \frac{\mathsf{Count}(x_i, x_j)}{m}$
  - Compute mutual information:

$$\widehat{I}(X_i, X_j) = \sum_{x_i, x_j} \widehat{P}(x_i, x_j) \log \frac{\widehat{P}(x_i, x_j)}{\widehat{P}(x_i) \widehat{P}(x_j)}$$

- Define a graph
  - Nodes  $X_1,...,X_n$
  - Edge (i,j) gets weight  $\widehat{I}(X_i, X_j)$
- Optimal tree BN
  - Compute maximum weight spanning tree (e.g. Prim's, Kruskal's algorithm O(nlog n))
  - Directions in BN: pick any node as root, breadth-first-search defines directions



# Learning Bayesian Networks for General Graphs

**Theorem**: The problem of learning a BN structure with at most d parents is NP-hard for any (fixed) d>1 (Note: tree d=1)

- Mostly heuristic (exploit score decomposition).
- Chow-Liu: provides best tree approximation to any distribution.
- Graph that maximizes ML score: complete graph!
  - Adding a parent always increase ML score!
  - The more edges, the fewer independence assumptions, the higher the likelihood of the data, but will overfit...
- Start with Chow-Liu tree.
  - Add, delete, invert edges.
  - Evaluate ML score + penalty



# **Learning Graphical Models**

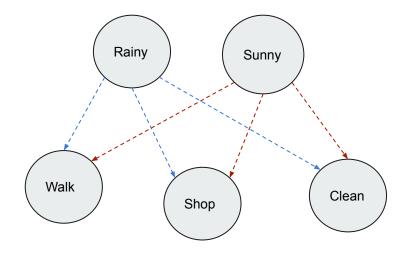
- Two Step Procedures:
  - ▶ 1. **Model Selection**; estimate graph structure
  - ▶ 2. Parameter Inference given graph structure
- Score Based Approaches: search over space of graphs, with a score for graph based on parameter inference
- Constraint-based Approaches: estimate individual edges by hypothesis tests for conditional independences
- Caveats: (a) difficult to provide guarantees for estimators; (b) estimators are NP-Hard



# **Recall Bayesian networks**

#### Consists of 2 parts:

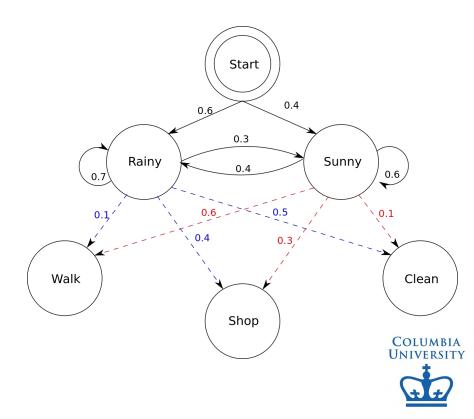
- Probability belongs to each class
- Class conditional probability





# **Limitation of Bayesian networks**

- Doing full Bayesian networks are computationally expensive;
- Performance is suboptimal on high-dimensional data;
- Bayesian networks are directed acyclic graphical models, where the directed edges are used to capture causal relationship between random variables, but we need undirected graphical model with potential loops in between nodes to model correlations.
- Bayesian networks cannot be used to model temporal/sequence data.



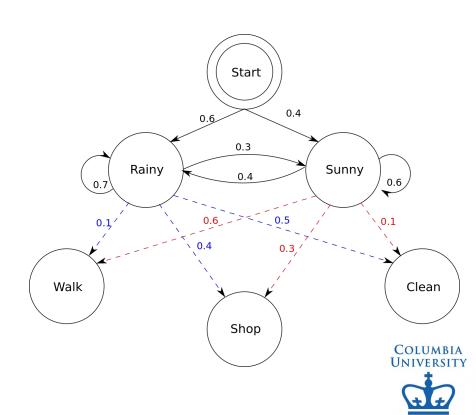
#### **Hidden Markov models**

Model a set of observations using a set of hidden states:

(observations, hidden states) such as:

- (pixel inputs, features)
- (range/visual sensor, location)
- (sound/visual signal, language/situation/words)
- (facial expression, results from the driving test)

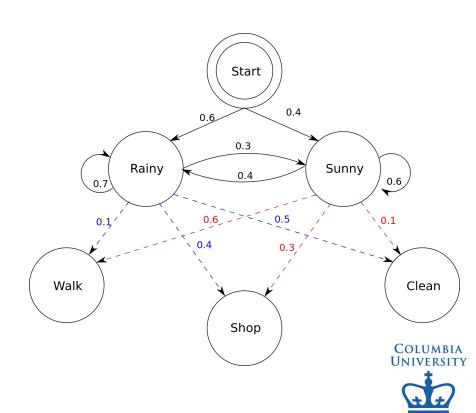
hidden state generates observation; hidden state transitions to other hidden states



# **Example: Daisy's diary**

Daisy kept a diary on what she was doing but she forgot to keep track of the weather. Let's treat the weather conditions as hidden states:

- With 60%/40%, today is rainy/sunny
- If today is rainy, it would either remain rainy (70%), or could be sunny tomorrow (30%).

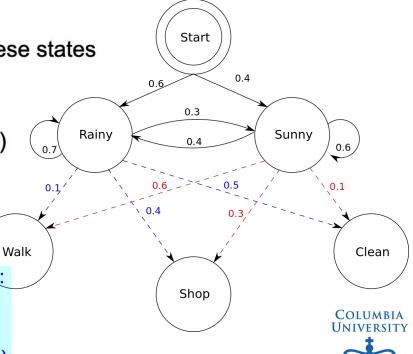


#### Definition of a hidden Markov model

- A set of states {s<sub>1</sub> ... s<sub>n</sub>}
  - In each time point we are in exactly one of these states denoted by  $\mathbf{q}_{\scriptscriptstyle{t}}$
- Π<sub>i</sub>, the probability that we start at state s<sub>i</sub>
- A transition probability model, P(q<sub>t</sub> = s<sub>i</sub> | q<sub>t-1</sub> = s<sub>i</sub>)
- A set of possible outputs Σ
  - At time t we emit a symbol  $\sigma \in \Sigma$
- An emission probability model, p(o<sub>t</sub> = σ | s<sub>i</sub>)

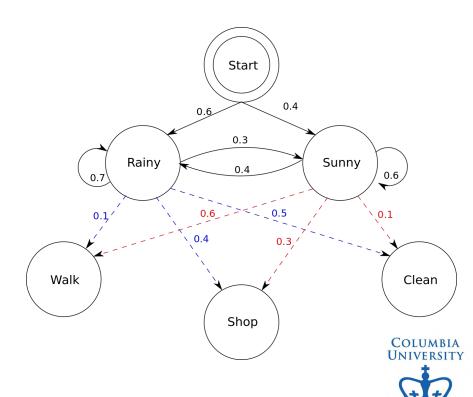
An important aspect of this definition is the Markov property:  $q_{t+1}$  is conditionally independent of  $q_{t-1}$  (and any earlier time points) given  $q_t$ 

More formally  $P(q_{t+1} = s_i | q_t = s_j) = P(q_{t+1} = s_i | q_t = s_j, q_{t-1} = s_j)$ 



Several questions people can ask:

- What is the current weather?
- What is the probability for shopping tomorrow?
- What is the probability for shopping next week?



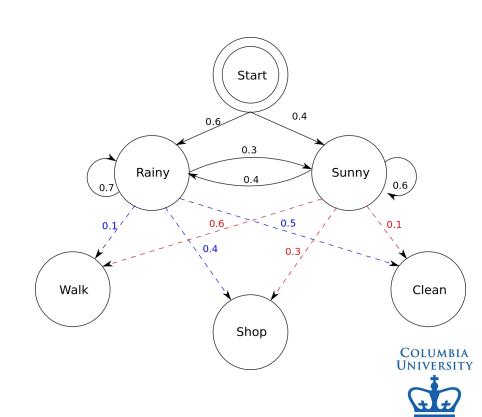
#### When no observation:

- What is the current weather?
  - P[rainy] = 0.6 P[sunny] = 0.4
- What is the weather tomorrow?

$$P[rainy] = 0.6 \times 0.7 + 0.4 \times 0.4$$

$$P[sunny] = 0.6 \times 0.3 + 0.4 \times 0.6$$

• What is the weather on the t-th day?



#### When no observation:

- What is the current weather?
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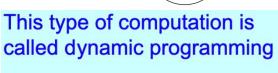
What is the weather on the t-th day?

Lets define  $p_t(i)$  = probability state i at time  $t = p(q_t = s_i)$ 

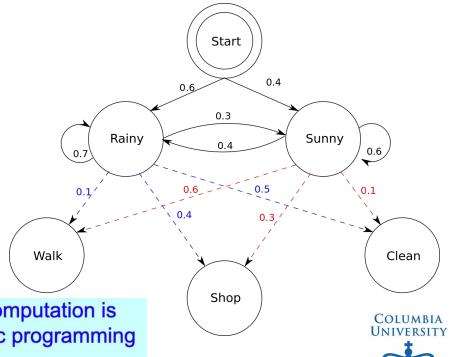
We can determine p<sub>t</sub>(i) by induction

1. 
$$p_1(i) = \Pi_i$$

2. 
$$p_t(i) = \sum_j p(q_t = s_i | q_{t-1} = s_j)p_{t-1}(j)$$



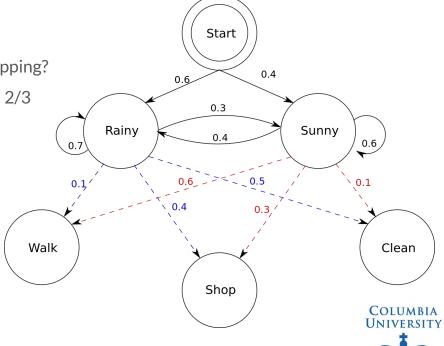
called dynamic programming Complexity: O(n<sup>2</sup>\*t)



#### When with observations:

What is the current weather given that Daisy is shopping?

 $P[rainy|shopping] = 0.6 \times 0.4/(0.6 \times 0.4 + 0.4 \times 0.3) = 2/3$ 



We want to compute P(q<sub>t</sub> = A | O<sub>1</sub> ... O<sub>t</sub>)

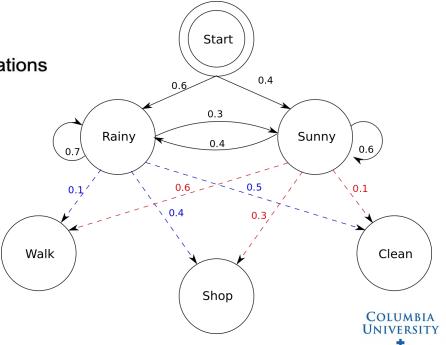
 For ease of writing we will use the following notations (commonly used in the literature)

•  $a_{j,i} = P(q_t = s_i | q_{t-1} = s_j)$ 

•  $b_i(o_t) = P(o_t | s_i)$ 

Emission probability

Transition probability



- We want to compute P(q<sub>t</sub> = A | O<sub>1</sub> ... O<sub>t</sub>)
- Lets start with a simpler question. Given a sequence of states Q, what is P(Q | O<sub>1</sub> ... O<sub>t</sub>) = P(Q | O)?
  - It is pretty simple to move from P(Q|O) to  $P(q_t = A|O)$
  - In some cases P(Q | O) is the more important question
    - Speech processing
    - NLP



$$P(Q|O) = \frac{P(O|Q)P(Q)}{P(O)}$$
Easy, P(O | Q) = P(o<sub>1</sub> | q<sub>1</sub>) P(o<sub>2</sub> | q<sub>2</sub>) ... P(o<sub>t</sub> | q<sub>t</sub>)
$$P(Q) = P(q_1) P(q_2 | q_1) ... P(q_t | q_{t-1})$$

But it is comparatively hard to compute P(O),

i.e. the probability of seeing a set of observations.



- What is the probability of seeing a set of observations:
  - An important question in it own rights, for example classification using two HMMs
- Define  $\alpha_t(i) = P(o_1, o_2 ..., o_t \land q_t = s_i)$
- α<sub>t</sub>(i) is the probability that we:
  - 1. Observe o<sub>1</sub>, o<sub>2</sub> ..., o<sub>t</sub>
  - 2. End up at state i



- We want to compute P(Q | O)
- For this, we only need to compute P(O)
- We know how to compute α<sub>t</sub>(i)

```
From now its easy  \alpha_t(i) = P(o_1, o_2 ..., o_t \land q_t = s_i)  so  P(O) = P(o_1, o_2 ..., o_t) = \Sigma_i P(o_1, o_2 ..., o_t \land q_t = s_i) = \Sigma_i \alpha_t(i)  note that  p(q_t = s_i \mid o_1, o_2 ..., o_t) = \frac{\alpha_t(i)}{\sum_j \alpha_t(j)}
```



Q represents a set of state Q =  $\{s_1, s_2 \dots s_t\}$ 

O represents a set of emitted values  $O = \{o_1, o_2 \dots o_t\}$ 

- Computing P(Q) and  $P(q_t = s_i)$ 
  - If we cannot look at observations
- Computing  $P(Q \mid O)$  and  $P(q_t = s_i \mid O)$ 
  - When we have observation and care about the last state only
- Computing argmax<sub>Q</sub>P(Q | O)
  - When we care about the entire path



# **Computational complexity**

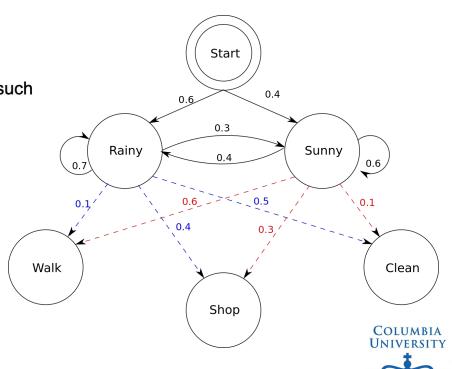
- How long does it take to compute P(Q | O)?
- P(Q): O(t)
- P(O|Q): O(t)
- P(O): O(n<sup>2</sup>t)



- We are almost done ...
- One final question remains
   How do we find the most probable path, that is Q\* such that

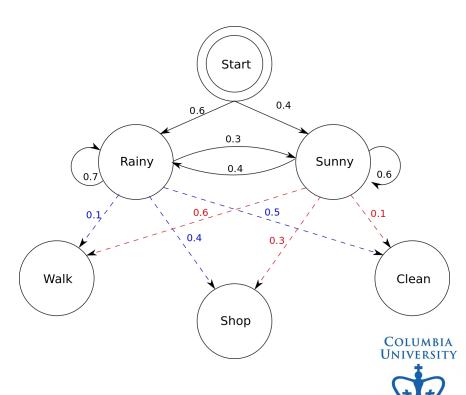
$$P(Q^* \mid O) = argmax_Q P(Q \mid O)$$
?

- This is an important path
  - The words in speech processing
  - The set of genes in the genome
  - etc.



What's the most likely sequence for you to see
 Daisy goes out for walking for 7 days in a week?

$$\arg \max_{Q} P(Q \mid O) = \arg \max_{Q} \frac{P(O \mid Q)P(Q)}{P(O)}$$
$$= \arg \max_{Q} P(O \mid Q)P(Q)$$



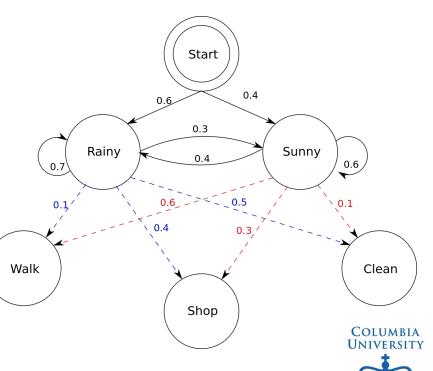
$$\operatorname{arg\,max}_{\mathcal{Q}} P(Q \mid O) = \operatorname{arg\,max}_{\mathcal{Q}} \frac{P(O \mid Q)P(Q)}{P(O)}$$

We will use the following definition:

$$\delta_t(i) = \max_{q_1...q_{t-1}} p(q_1...q_{t-1} \wedge q_t = s_i \wedge O_1...O_t)$$

In other words we are interested in the most likely path from 1 to t that:

- 1. Ends in S<sub>i</sub>
- 2. Produces outputs O<sub>1</sub> ... O<sub>t</sub>



$$\operatorname{arg\,max}_{\mathcal{Q}} P(Q \mid O) = \operatorname{arg\,max}_{\mathcal{Q}} \frac{P(O \mid Q)P(Q)}{P(O)}$$

We will use the following definition:

$$\delta_t(i) = \max_{q_1...q_{t-1}} p(q_1...q_{t-1} \land q_t = s_i \land O_1...O_t)$$

In other words we are interested in the most likely path from 1 to t that:

- 1. Ends in S<sub>i</sub>
- 2. Produces outputs O<sub>1</sub> ... O<sub>t</sub>

Q: Given  $\delta_t(i)$ , how can we compute  $\delta_{t+1}(i)$ ?

A: To get from  $\delta_t(i)$  to  $\delta_{t+1}(i)$  we need to

- 1. Add an emission for time t+1  $(O_{t+1})$
- 2. Transition to state s<sub>i</sub>

$$\begin{split} \delta_{t+1}(i) &= \max_{q_1 \dots q_t} p(q_1 \dots q_t \wedge q_{t+1} = s_i \wedge O_1 \dots O_{t+1}) \\ &= \max_{j} \delta_t(j) p(q_{t+1} = s_i \mid q_t = s_j) p(O_{t+1} \mid q_{t+1} = s_i) \\ &= \max_{j} \delta_t(j) a_{j,i} b_i(O_{t+1}) \end{split}$$



# Inference in HMM: the Viterbi algorithm

$$\delta_{t+1}(i) = \max_{q_1 \dots q_t} p(q_1 \dots q_t \land q_{t+1} = s_i \land O_1 \dots O_{t+1})$$

$$= \max_{j} \delta_t(j) p(q_{t+1} = s_i \mid q_t = s_j) p(O_{t+1} \mid q_{t+1} = s_i)$$

$$= \max_{j} \delta_t(j) a_{j,i} b_i(O_{t+1})$$

- Once again we use dynamic programming for solving  $\delta_{t}\!\!(i)$
- Once we have  $\delta_t(i)$ , we can solve for our  $P(Q^*|O)$

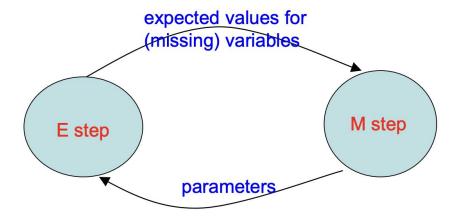
By:

$$P(Q^* \mid O) = argmax_Q P(Q \mid O) = Path defined by max_i \delta_t(i)$$



# Learning HMMs: EM algorithm revisit

- E step: Fill in the expected values for the missing variables
- M step: Regular maximum likelihood estimation (MLE) using the values computed in the E step and the values of the other variables
- Guaranteed to converge (though only to a local minima).





# EM algorithm for HMMs (Baum-Welch)

#### E step:

• Compute  $S_t(i)$  and  $S_t(i,j)$  for all t, i, and j  $P(q_t = s_i \mid O_1, \cdots, O_T) = S_t(i)$   $P(q_t = s_i, q_{t+1} = s_i \mid O_1, \cdots, O_T) = S_t(i,j)$ 

#### M step:

Compute transition probabilities:  $a_{i,j} = \frac{\hat{n}(i,j)}{\sum_{k} \hat{n}(i,k)}$   $\hat{n}(i,j) = \sum_{t} S_{t}(i,j)$ 

Compute emission probabilities (here we assume a multinomial distribution):

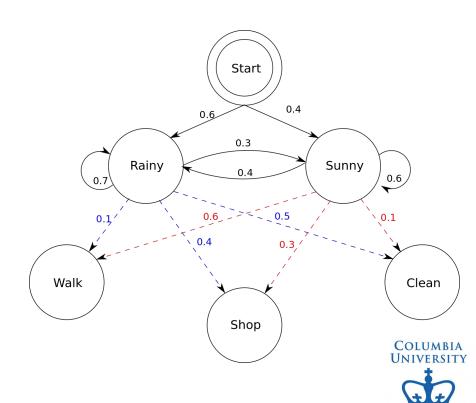
$$b_k(j) = \frac{B_k(j)}{\sum_{i} B_k(i)}$$

$$B_k(j) = \sum_{t|o_t=j} S_t(k)$$



# **Takeaways**

- Why HMMs? Which applications are suitable?
- Inference in HMMs
  - No observations
  - Probability of next state with observations
  - Maximum scoring path (Viterbi)
- Learning HMMs:
  - EM algorithm



#### References

- Christopher Bishop: Pattern Recognition and Machine Learning, Chapter 5
- Ziv Bar-Joseph, Tom Mitchell, Pradeep Ravikumar and Aarti Singh: CMU 10-701
- Ryan Tibshirani: CMU 10-725
- Ruslan Salakhutdinov: CMU 10-703
- https://machinelearningmastery.com/what-are-generative-adversarial-networks-gans/

