

Math 189Z

Lecture 3: Time series data, Markov Chains, and HMM

COVID-19: Data Analytics and Machine Learning

PROF. WEIQING GU

SPRING 2020





COVID-19 Dashboard by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU)



Total Confirmed
2,196,109

Confirmed Cases by
Country/Region/Sovereignty

672,303 US

188,068 Spain

172,434 Italy

147,113 France

138,456 Germany

109,769 United Kingdom

83,760 China

79,494 Iran

78,546 Turkey

36,138 Belgium

32,008 Russia

31,161 Canada

30,891 Brazil

30,618 Netherlands

27,078 Switzerland

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Last Updated at (M/D/YYYY)

4/17/2020, 9:38:37 AM



Total Deaths
149,024

22,745 deaths
Italy

19,478 deaths
Spain

17,920 deaths
France

14,576 deaths
United Kingdom

11,477 deaths
New York City New York US

5,163 deaths
Belgium

4,958 deaths
Iran

Deaths Recovered

Total Tested in the US
3,423,034

550,579 tested
New York US

246,400 tested
California US

224,141 tested
Florida US

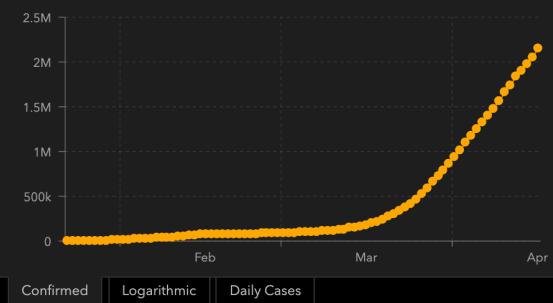
158,547 tested
Texas US

151,830 tested
New Jersey US

141,470 tested
Pennsylvania US

140,773 tested
Massachusetts US

100,000+ tested
US Tested US Hospitalization



185
countries/regions

Lancet Inf Dis Article: [Here](#). Mobile Version: [Here](#).
Lead by JHU CSSE. Automation Support: Esri Living Atlas team and JHU APL. Contact US. FAQ.
Data sources: WHO, CDC, ECDC, NHC, DXY, 1point3acres, Worldometers.info, BNO, the COVID Tracking Project (testing and hospitalizations), state and national government health departments, and local media reports. Read more in this [blog](#).

- <https://coronavirus.jhu.edu/map.html>

Overview



- COVID-19 confirmed cases have been increased but not doubled since our last meeting

In the case of Italy:

27,980 Italy



80,589 Italy



115,242 Italy



147,577 Italy



172,434 Italy

$$R1 = (147-115)/(115-80) = 0.91 < 1$$

$$R2 = (172-147)/(147-115) = \\ .78 < 0.91 < 1$$

USA:

$$US_R0 = (245-86)/(86-0.5) = \\ 1.97$$

$$US_R1 = (475-245)/(245-86) = \\ 1.44$$

$$US_R2 = (672-457)/(475-245) \\ 0.85 < 1$$

5,204 US



86,012 US



245,540 US



475,749 US



672,303 US



Coronavirus COVID-19 Global Cases by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU)



Total Confirmed

1,650,210

Confirmed Cases by Country/Region/Sovereignty

475,749	US
157,053	Spain
147,577	Italy
119,624	Germany
118,790	France
82,941	China
71,078	United Kingdom
68,192	Iran
47,029	Turkey
26,667	Belgium
24,548	Switzerland
23,245	Netherlands
21,243	Canada
18,397	Brazil
15,472	Portugal

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Last Updated at (M/D/YYYY)
4/10/2020, 10:02:32 AM



Cumulative Confirmed Cases

Active Cases

185
countries/regions

Lancet Inf Dis Article: [Here](#). Mobile Version: [Here](#). Visualization: [JHU CSSE](#). Automation Support: [Esri Living Atlas team](#) and [JHU APL](#). Contact US. [FAQ](#).
Data sources: [WHO](#), [CDC](#), [ECDC](#), [NHC](#), [DXY](#), [1point3acres](#), [Worldometers.info](#), [BNO](#), state and national government health departments, and local media reports. Read more in this [blog](#).

Total Deaths

100,376

18,849 deaths
Italy

15,970 deaths
Spain

12,210 deaths
France

8,958 deaths
United Kingdom

5,150 deaths
New York City **New York US**

4,232 deaths
Iran

3,216 deaths
Hubei China

3,019 deaths
Belgium

Total Recovered

368,669

77,791 recovered
China

55,668 recovered
Spain

52,407 recovered
Germany

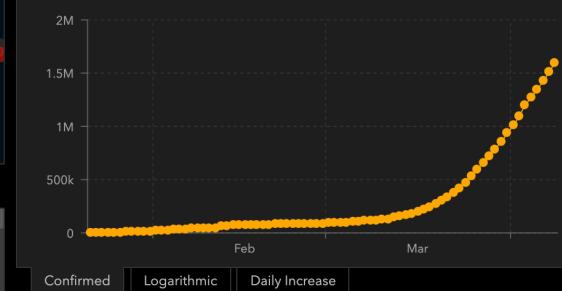
35,465 recovered
Iran

30,455 recovered
Italy

26,645 recovered
US

23,469 recovered
France

10,600 recovered
Switzerland



Confirmed Logarithmic Daily Increase



Coronavirus COVID-19 Global Cases by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU)



Total Confirmed

1,016,128

Confirmed Cases by Country/Region/Sovereignty

245,540 US

115,242 Italy

112,065 Spain

84,794 Germany

82,456 China

59,929 France

50,468 Iran

34,173 United Kingdom

18,827 Switzerland

18,135 Turkey

15,348 Belgium

14,788 Netherlands

11,284 Canada

11,129 Austria

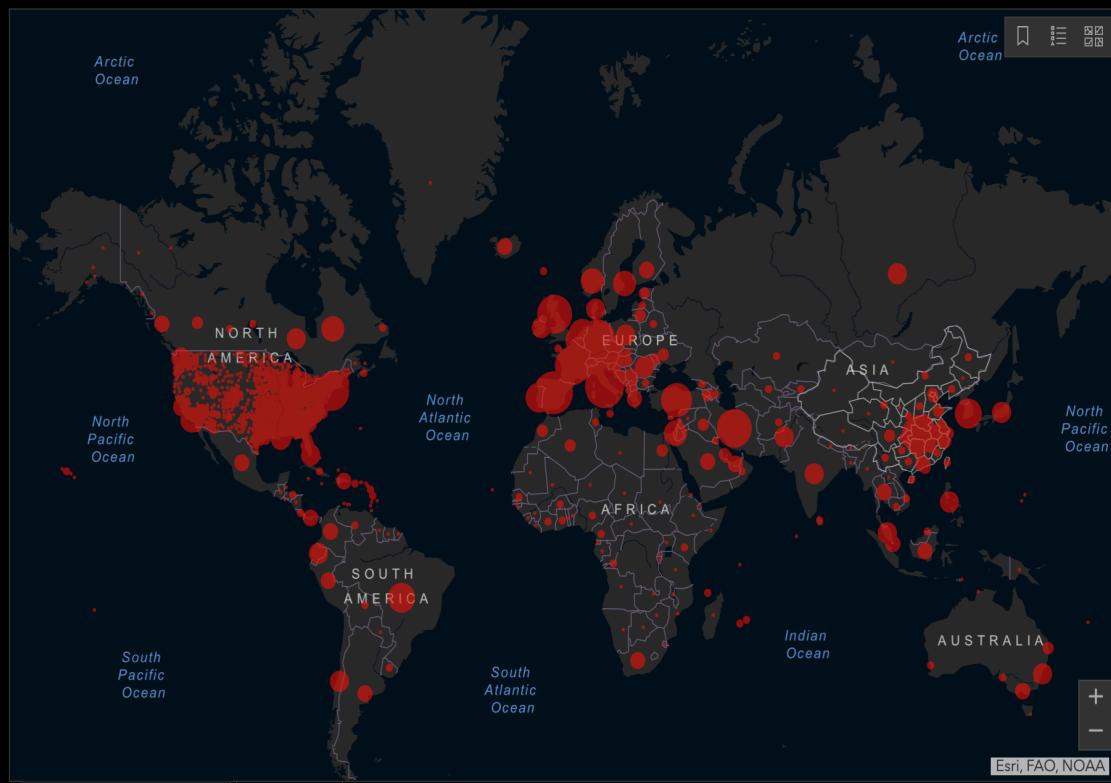
10,062 Korea, South

8,024 Other

Admin0 Admin1 Admin2

Last Updated at (M/D/YYYY)

4/2/2020, 9:12:43 PM



181
countries/regions

Lancet Inf Dis Article: [Here](#). Mobile Version: [Here](#). Visualization: JHU CSSE. Automation Support: Esri Living Atlas team and JHU APL. Contact US. FAQ.

Data sources: WHO, CDC, ECDC, NHC, DXY, 1point3acres, Worldometers.info, BNO, state and national government health departments, and local media reports. Read more in this [blog](#).

Total Deaths

53,146

13,915 deaths
Italy

10,348 deaths
Spain

5,387 deaths
France

3,203 deaths
Hubei China

3,160 deaths
Iran

2,921 deaths
United Kingdom

1,562 deaths
New York City New York US

1,339 deaths
Netherlands

1,107 deaths
Korea, South

1,012 deaths
Other

Total Recovered

211,615

76,724 recovered
China

26,743 recovered
Spain

22,440 recovered
Germany

18,278 recovered
Italy

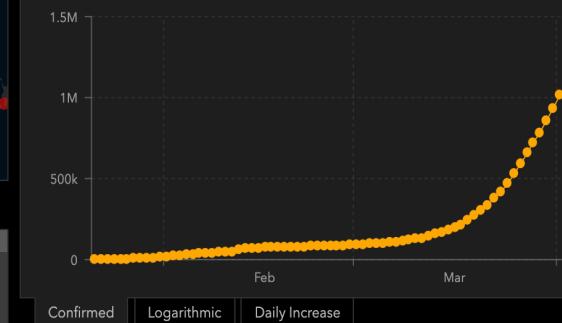
16,711 recovered
Iran

12,548 recovered
France

9,148 recovered
US

6,021 recovered
Korea, South

4,012 recovered
Other





Coronavirus COVID-19 Global Cases by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins ...



Total Confirmed

553,244



Confirmed Cases by Country/Region/Sovereignty

86,012 US

81,897 China

80,589 Italy

64,059 Spain

47,373 Germany

32,332 Iran

29,581 France

12,311 Switzerland

11,830 United Kingdom

9,332 Korea, South

8,641 Netherlands

7,393 Austria

7,284 Belgium

4,268 Portugal

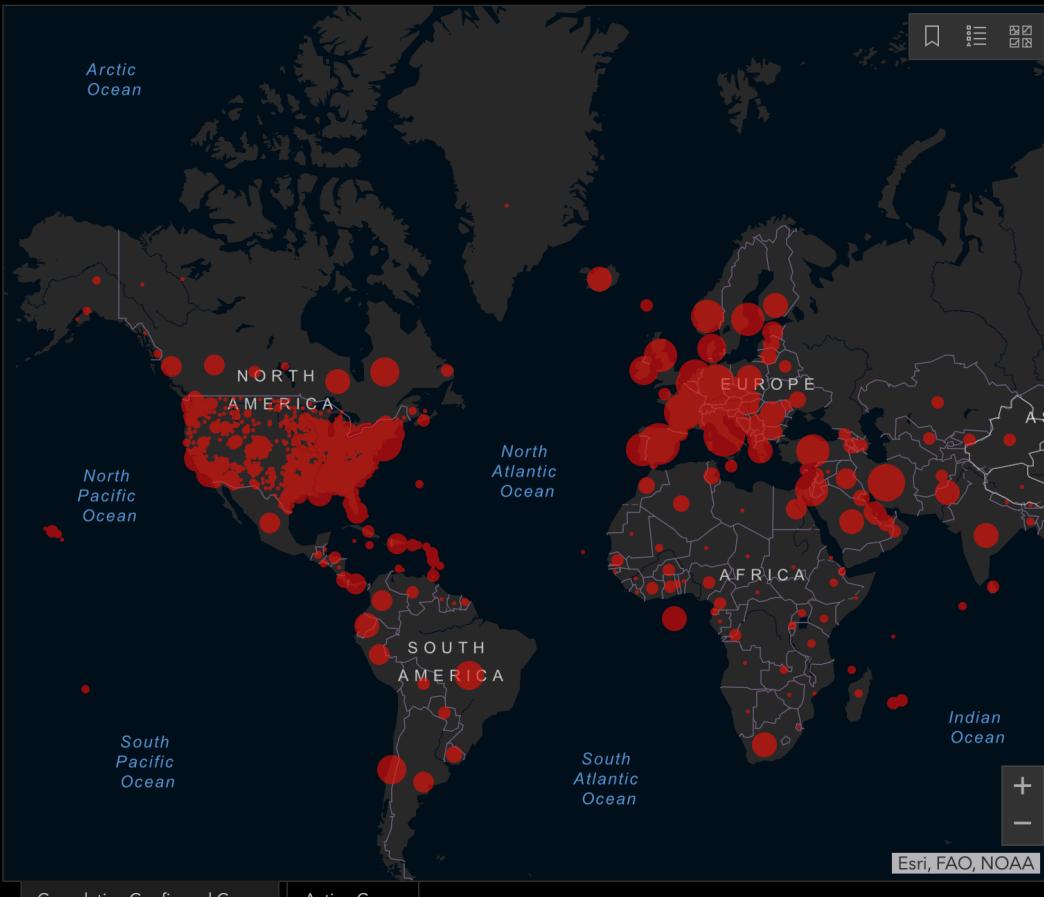
4,046 Canada

3,207 Norway

Admin1

Last Updated at (M/D/YYYY)

3/27/2020, 8:13:47 AM



Cumulative Confirmed Cases

Active Cases

176
countries/regions

Lancet Inf Dis Article: [Here](#). Mobile Version: [Here](#). Visualization: JHU CSSE. Automation Support: Esri Living Atlas team and JHU APL. Contact US. FAQ.

Data sources: WHO, CDC, ECDC, NHC, DXY, 1point3acres, Worldometers.info, BNO, state and national government health departments, and local media reports. Read more in this [blog](#).

Downloadable database: GitHub: [Here](#) Feature layer: [Here](#)

Total Deaths

25,035

8,215 deaths
Italy

4,858 deaths
Spain

3,174 deaths
Hubei China

2,378 deaths
Iran

1,696 deaths
France

578 deaths
United Kingdom

546 deaths
Netherlands

365 deaths
New York City New York US

Total Recovered

127,567

61,732 recovered
Hubei China

11,133 recovered
Iran

10,361 recovered
Italy

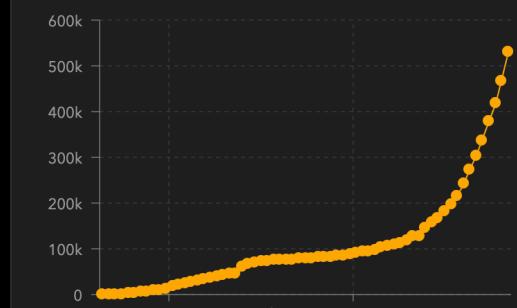
9,357 recovered
Spain

5,673 recovered
Germany

4,948 recovered
France

4,528 recovered
Korea, South

1,337 recovered
Guangdong China



Confirmed

Daily Increase



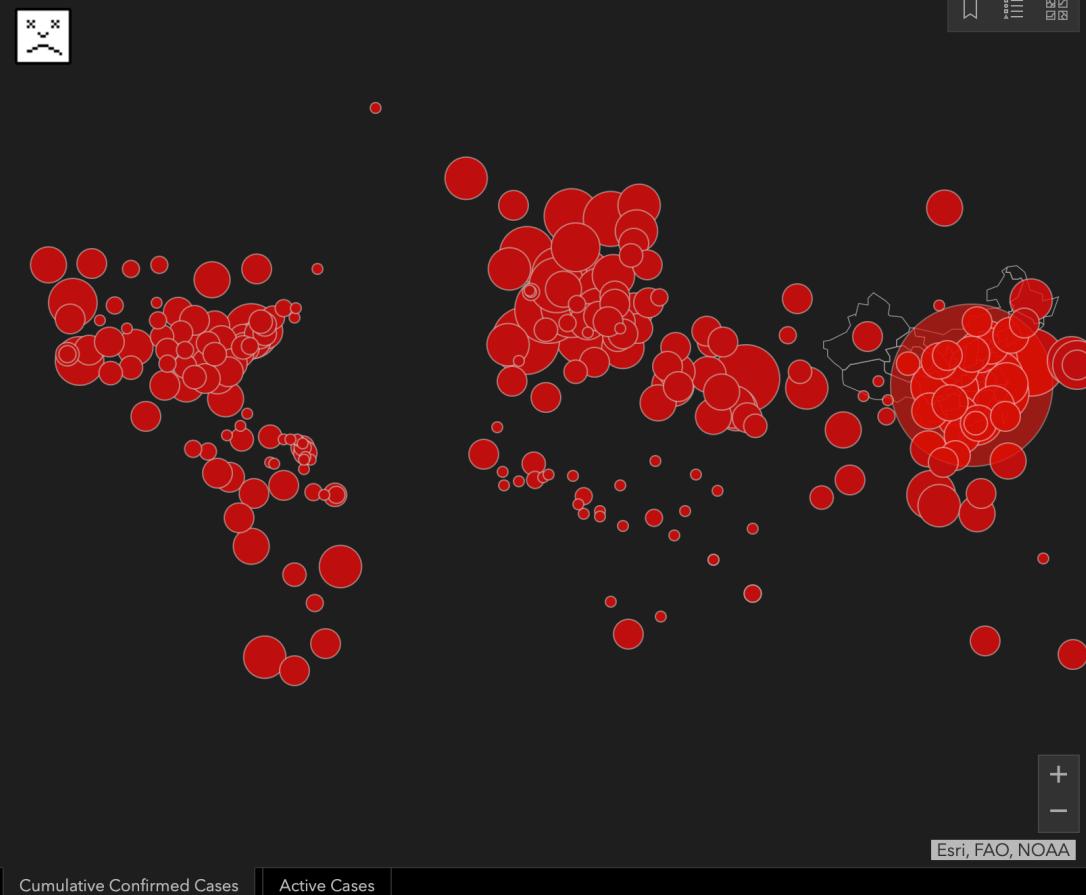
Total Confirmed
190,124

Confirmed Cases by
Country/Region/Sovereignty

81,058 China
27,980 Italy
16,169 Iran
11,309 Spain
8,604 Germany
8,320 Korea, South
6,664 France
5,204 US
2,700 Switzerland
1,960 United Kingdom
1,708 Netherlands
1,443 Norway
1,332 Austria
1,243 Belgium
1,190 Sweden
1,024 Denmark

Country/Region/Sovereign...

Last Updated at (M/D/YYYY)
3/17/2020, 9:33:04 AM



155
countries/regions

Lancet Inf Dis Article: [Here](#). Mobile Version: [Here](#). Visualization: [JHU CSSE](#). Automation Support: [Esri Living Atlas team](#) and [JHU APL](#). Data sources: WHO, CDC, ECDC, NHC and DXY and local media reports. Read more in this [blog](#). [Contact Us](#). [FAQ](#). Downloadable database: GitHub: [Here](#). Feature layer: [Here](#).

Total Deaths
7,516

3,111 deaths
Hubei China

2,158 deaths
Italy

988 deaths
Iran

509 deaths
Spain

148 deaths
France France

81 deaths
Korea, South

55 deaths
United Kingdom United Kingdom

48 deaths
Washington US

Total Recovered
80,643

56,003 recovered
Hubei China

5,389 recovered
Iran

2,749 recovered
Italy

1,407 recovered
Korea, South

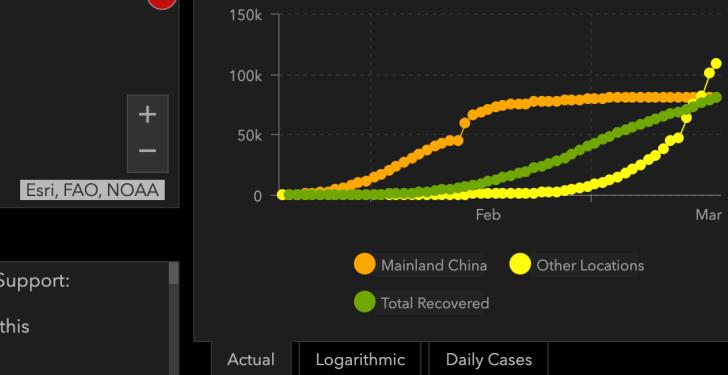
1,307 recovered
Guangdong China

1,250 recovered
Henan China

1,216 recovered
Zhejiang China

1,028 recovered
Spain

1,014 recovered



Today: Overview

Today:

1. Time series data
2. Markov Chains
3. HMM

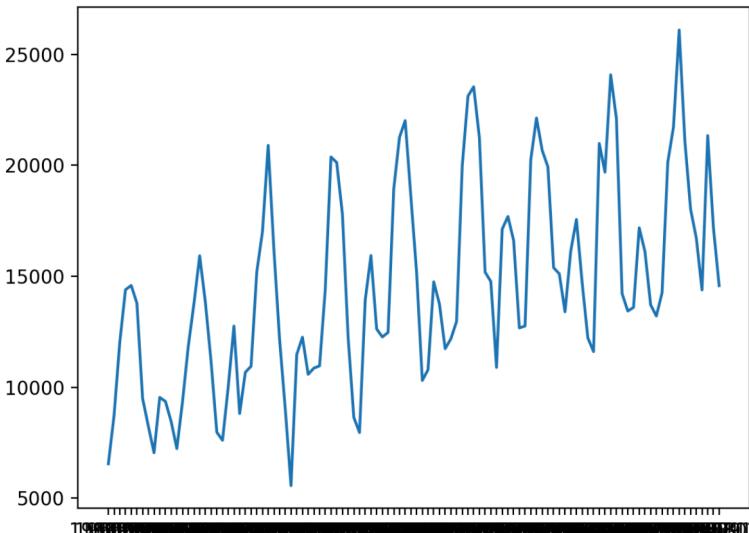
Recall: Last time, we covered

1. NMF (Non-negative Matrix Factorization)
2. LSA (Latent Semantic Analysis)

- <https://math189covid19.github.io/>

What is a Time series (data)?

- A **time series** is a **series** of data points indexed (or listed or graphed) in **time** order.
- Most commonly, a **time series** is a sequence taken at successive equally spaced points in **time**.
- Thus it is a sequence of discrete-**time** data.



This dataset is monthly and has nine years, or 108 observations. In our testing, will use the last year, or 12 observations, as the test set.

```
"Month", "Sales"  
"1960-01", 6550  
"1960-02", 8728  
"1960-03", 12026  
"1960-04", 14395  
"1960-05", 14587  
"1960-06", 13791  
"1960-07", 9498  
"1960-08", 8251  
"1960-09", 7049  
"1960-10", 9545  
"1960-11", 9364  
"1960-12", 8456  
"1961-01", 7237  
"1961-02", 9374  
"1961-03", 11837  
"1961-04", 13784  
"1961-05", 15926  
"1961-06", 13821  
"1961-07", 11143  
"1961-08", 7975  
"1961-09", 7610  
"1961-10", 10015  
"1961-11", 12759  
"1961-12", 11143
```

https://raw.githubusercontent.com/jbrownlee/Datasets/master/monthly_car-sales.csv

```
1 # load
2 series = read_csv('monthly-car-sales.csv', header=0, index_col=0)
```

Once loaded, we can summarize the shape of the dataset in order to determine the number of observations.

```
1 # summarize shape
2 print(series.shape)
```

We can then create a line plot of the series to get an idea of the structure of the series.

```
1 # plot
2 pyplot.plot(series)
3 pyplot.show()
```

We can tie all of this together; the complete example is listed below.

```
1 # load and plot dataset
2 from pandas import read_csv
3 from matplotlib import pyplot
4 # load
5 series = read_csv('monthly-car-sales.csv', header=0, index_col=0)
6 # summarize shape
7 print(series.shape)
8 # plot
9 pyplot.plot(series)
10 pyplot.show()
```

Stock data is time series data



Task example: Find patterns in stock time series data



Markov Chains

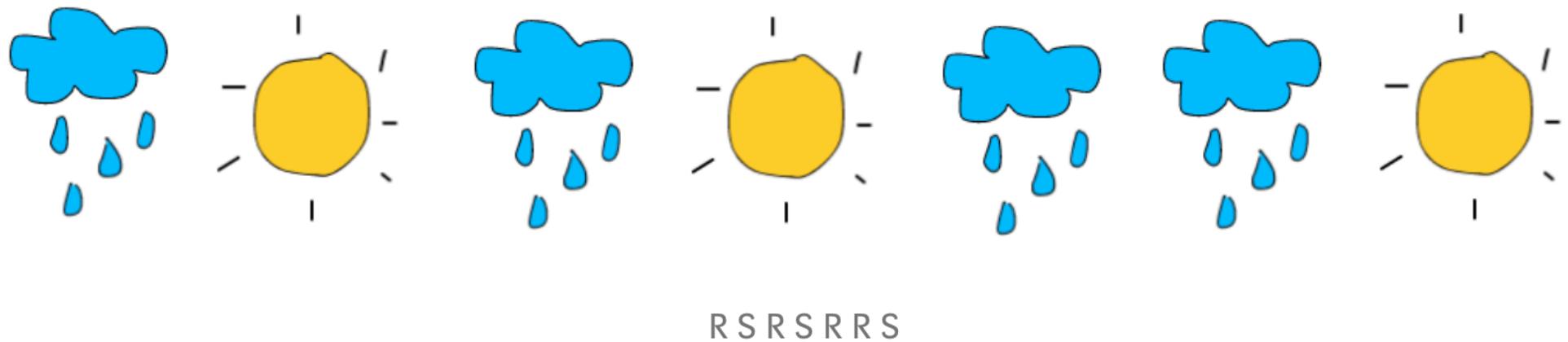
- Well known example: predict weather

In our simplified universe, the weather can only be in one of 2 possible states, “sunny” or “rainy”.

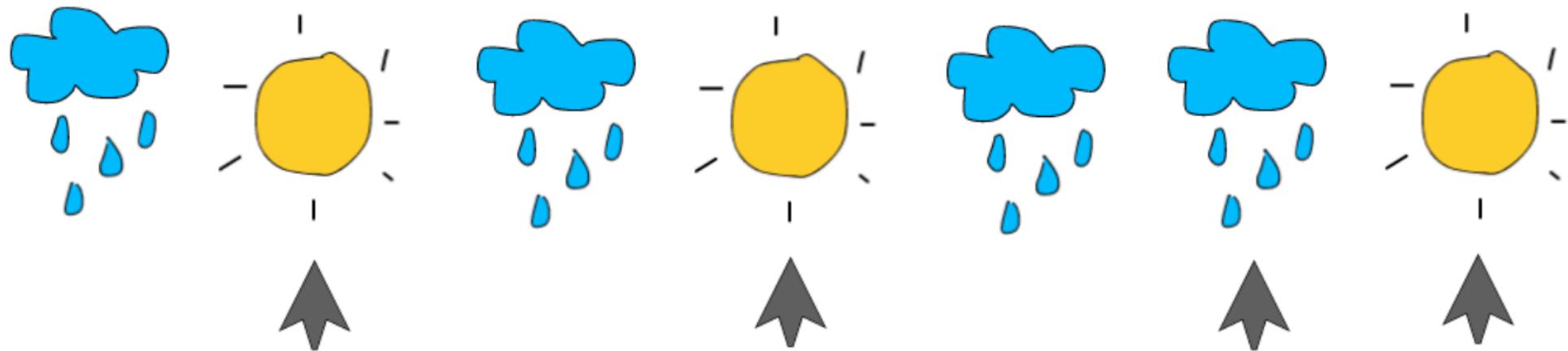
- The catch (in the context of Markov chains) is that the probability of it being sunny or rainy **tomorrow**, depends on whether it is sunny or rainy **today**.
- We'll derive these probabilities from past data, and construct a **transition matrix**.

Using the historic data to build a transition matrix

- Here we use 7 days of historical data on which to “train” our Markov chain. The days are:
[rain, sun, rain, sun, rain, rain, sun]

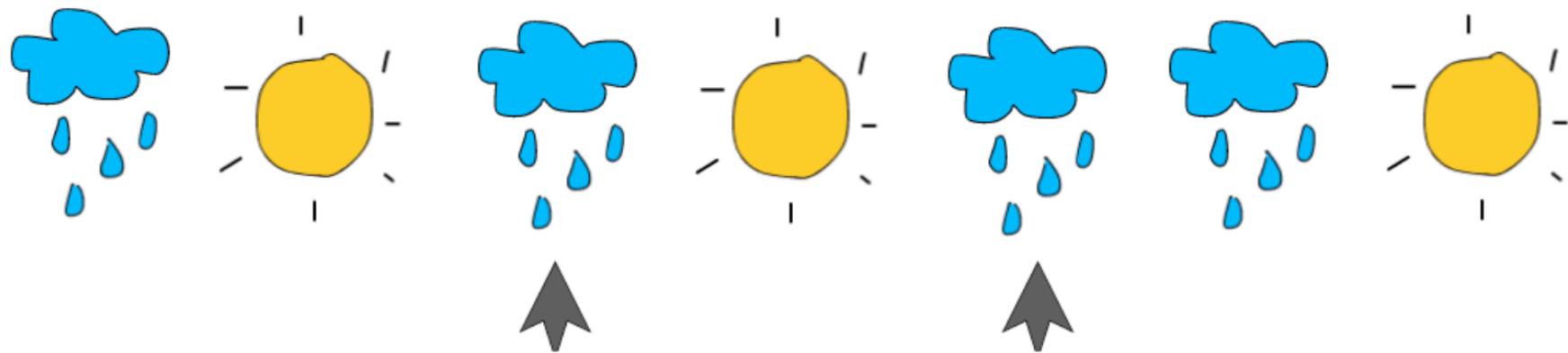


Now calculate the percentage of instances its sunny on days directly following rainy days.



3/4 so 75%.

Now calculate the percentage of instances its rainy on days directly following sunny days.

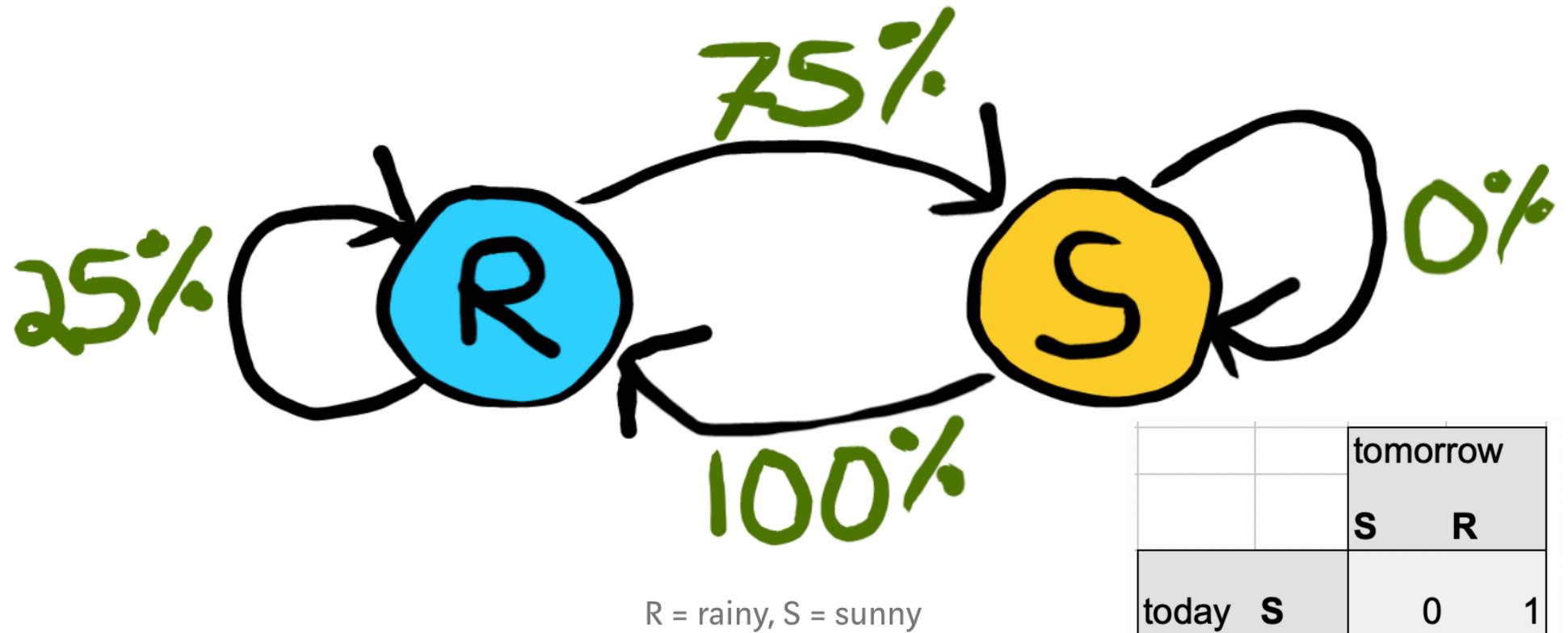


We'll build our transition matrix with that information, inferring missing percentages from the information we've already derived (rain-after-rain = 25% and sun-after-sun = 0%).

Transition Matrix

		tomorrow	
		S	R
today	S	0	1
	R	0.75	0.25

The Markov Chain

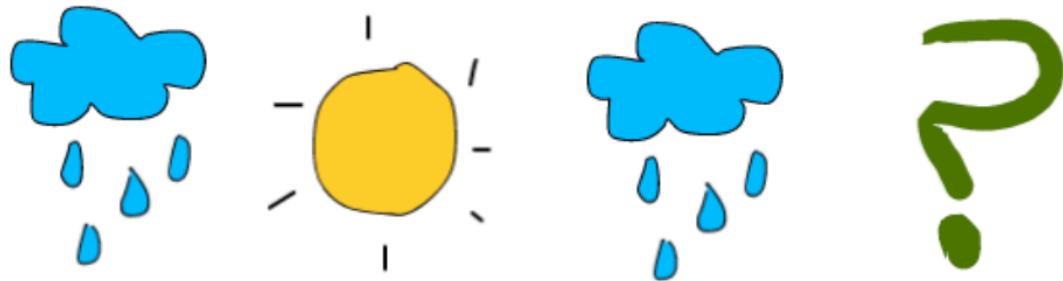


This diagram hits home the fact that probabilities are completely dependent on the current state, not the weather yesterday or the day before that.

Example 1:

The previous 3 days are [rainy, sunny, rainy].

What's the probability of rainy weather tomorrow?



R, S, R

Based on our previously trained model. Tomorrow has a 75% chance of sun and 25% chance of rain.

Example 2:

The previous 2 days are [rainy, rainy].

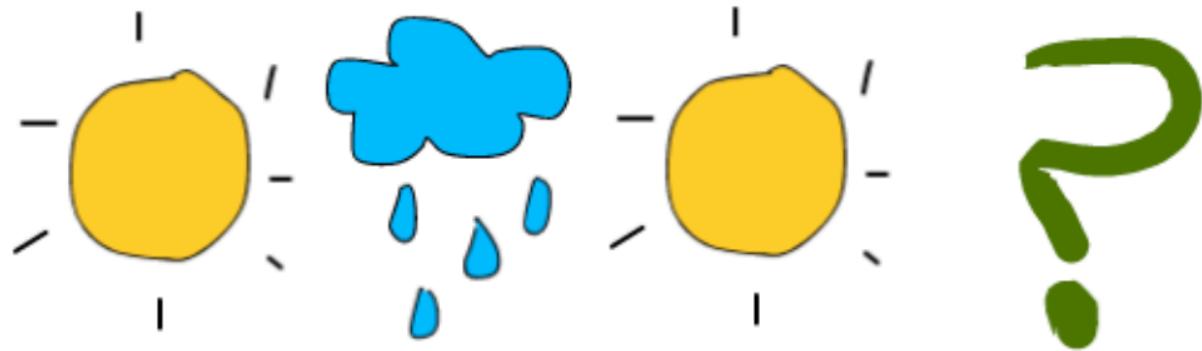


R R

Again, tomorrow has a 75% chance of sun and 25% chance of rain.

Example 3:

The previous 3 days are [sunny, rainy, sunny].



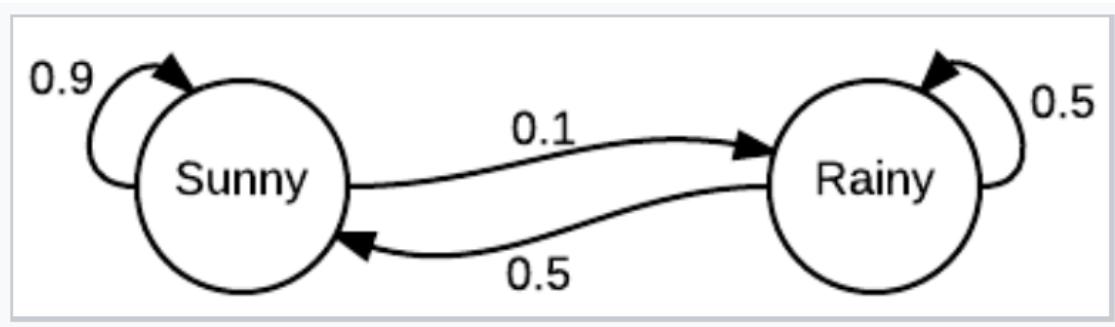
S R S

There is a 100% chance of rain tomorrow. It always rains on days after sun... sad I know...

Suppose we get the transition matrix with lots lots of data

- Say

$$P = \begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix}$$



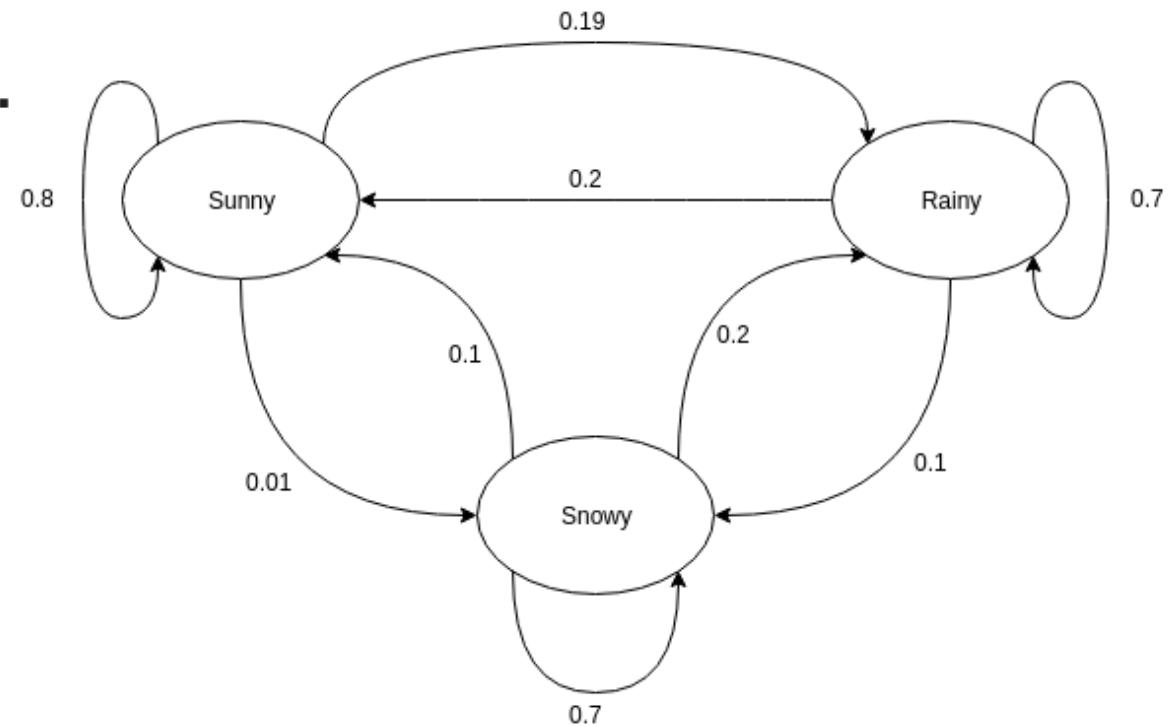
The above matrix as a graph.

The matrix P represents the weather model in which a sunny day is 90% likely to be followed by another sunny day, and a rainy day is 50% likely to be followed by another rainy day. The columns can be labelled "sunny" and "rainy", and the rows can be labelled in the same order.

Definition: stochastic matrix

$(P)_{ij}$ is the probability that, if a given day is of type i , it will be followed by a day of type j .

Notice that the rows of P sum to 1: this is because P is a **stochastic matrix**.



Predicting the weather

The weather on day 0 (today) is known to be sunny. This is represented by a vector in which the "sunny" entry is 100%, and the "rainy" entry is 0%:

$$\mathbf{x}^{(0)} = [1 \quad 0]$$

The weather on day 1 (tomorrow) can be predicted by:

$$\mathbf{x}^{(1)} = \mathbf{x}^{(0)} P = [1 \quad 0] \begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix} = [0.9 \quad 0.1]$$

Thus, there is a 90% chance that day 1 will also be sunny.

The weather on day 2 (the day after tomorrow) can be predicted in the same way:

$$\mathbf{x}^{(2)} = \mathbf{x}^{(1)} P = \mathbf{x}^{(0)} P^2 = [1 \quad 0] \begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix}^2 = [0.86 \quad 0.14]$$

Iterative process

or

$$\mathbf{x}^{(2)} = \mathbf{x}^{(1)} P = [0.9 \quad 0.1] \begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix} = [0.86 \quad 0.14]$$

General rules for day n are:

$$\mathbf{x}^{(n)} = \mathbf{x}^{(n-1)} P$$

$$\mathbf{x}^{(n)} = \mathbf{x}^{(0)} P^n$$

The steady state vector is defined as:

$$\mathbf{q} = \lim_{n \rightarrow \infty} \mathbf{x}^{(n)}$$

but converges to a strictly positive vector only if P is a regular transition matrix (that is, there is at least one P^n with all non-zero entries).

Since the \mathbf{q} is independent from initial conditions, it must be unchanged when transformed by P .^[4] This makes it an **eigenvector** (with **eigenvalue** 1), and means it can be derived from P .^[4] For the weather example:

$$P = \begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix}$$

$$\begin{aligned} \mathbf{q}P &= \mathbf{q} \\ &= \mathbf{q}I \end{aligned} \quad (\mathbf{q} \text{ is unchanged by } P.)$$

$$\mathbf{q}(P - I) = \mathbf{0}$$

$$\mathbf{q} \left(\begin{bmatrix} 0.9 & 0.1 \\ 0.5 & 0.5 \end{bmatrix} - \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \right) = \mathbf{0}$$

$$\mathbf{q} \begin{bmatrix} -0.1 & 0.1 \\ 0.5 & -0.5 \end{bmatrix} = \mathbf{0}$$

$$[q_1 \quad q_2] \begin{bmatrix} -0.1 & 0.1 \\ 0.5 & -0.5 \end{bmatrix} = [0 \quad 0]$$

$$-0.1q_1 + 0.5q_2 = 0$$

and since they are a probability vector we know that

$$q_1 + q_2 = 1.$$

Solving this pair of simultaneous equations gives the steady state distribution:

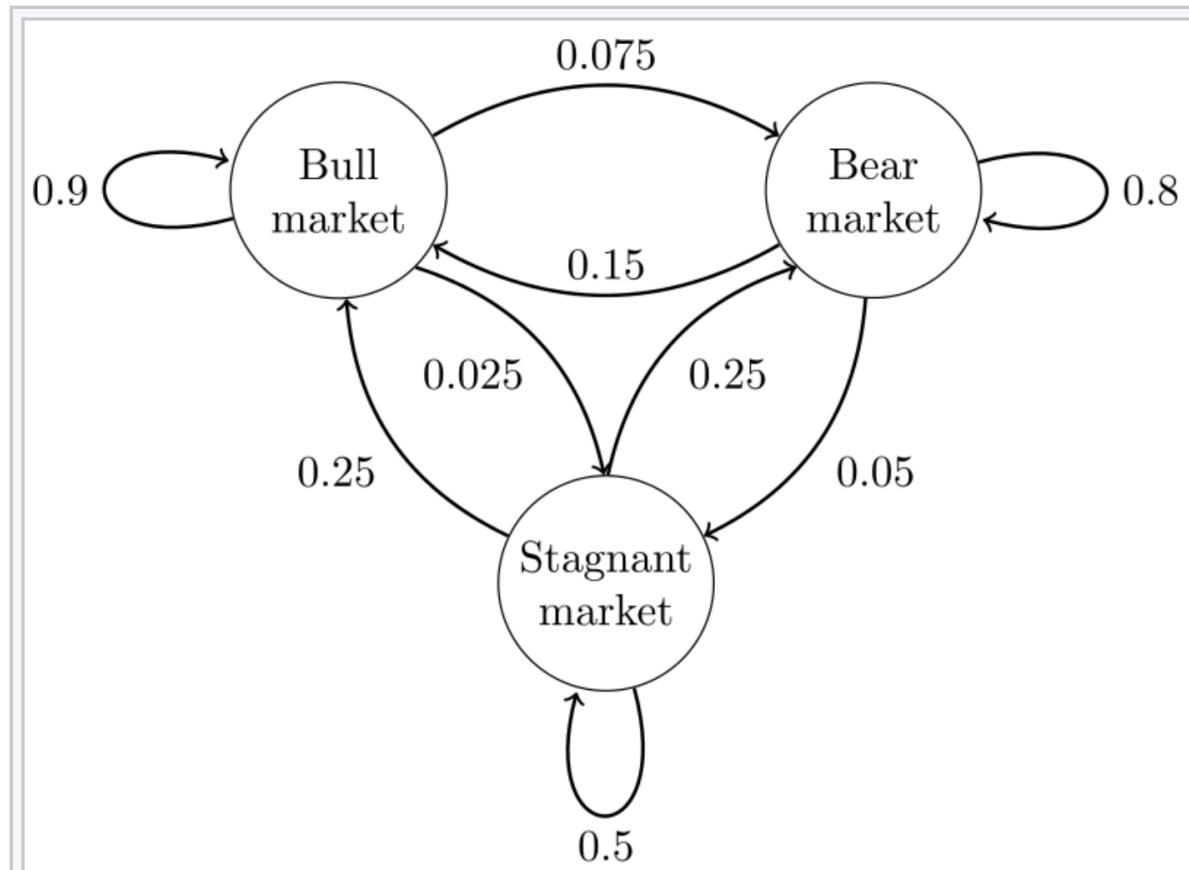
$$[q_1 \quad q_2] = [0.833 \quad 0.167]$$

In conclusion, in the long term, about 83.3% of days are sunny.

Similarly you can use Markov Chains to predict stock trends

Stock market [edit]

A state diagram for a simple example is shown in the figure on the right, using a directed graph to picture the state transitions. The states represent whether a hypothetical stock market is exhibiting a bull market, bear market, or stagnant market trend during a given week.



Using a directed graph, the probabilities of the possible states a hypothetical stock market can exhibit is represented. The matrix on the left shows how probabilities corresponding to different states can be arranged in matrix form.

- https://en.wikipedia.org/wiki/Examples_of_Markov_chains

Exercise

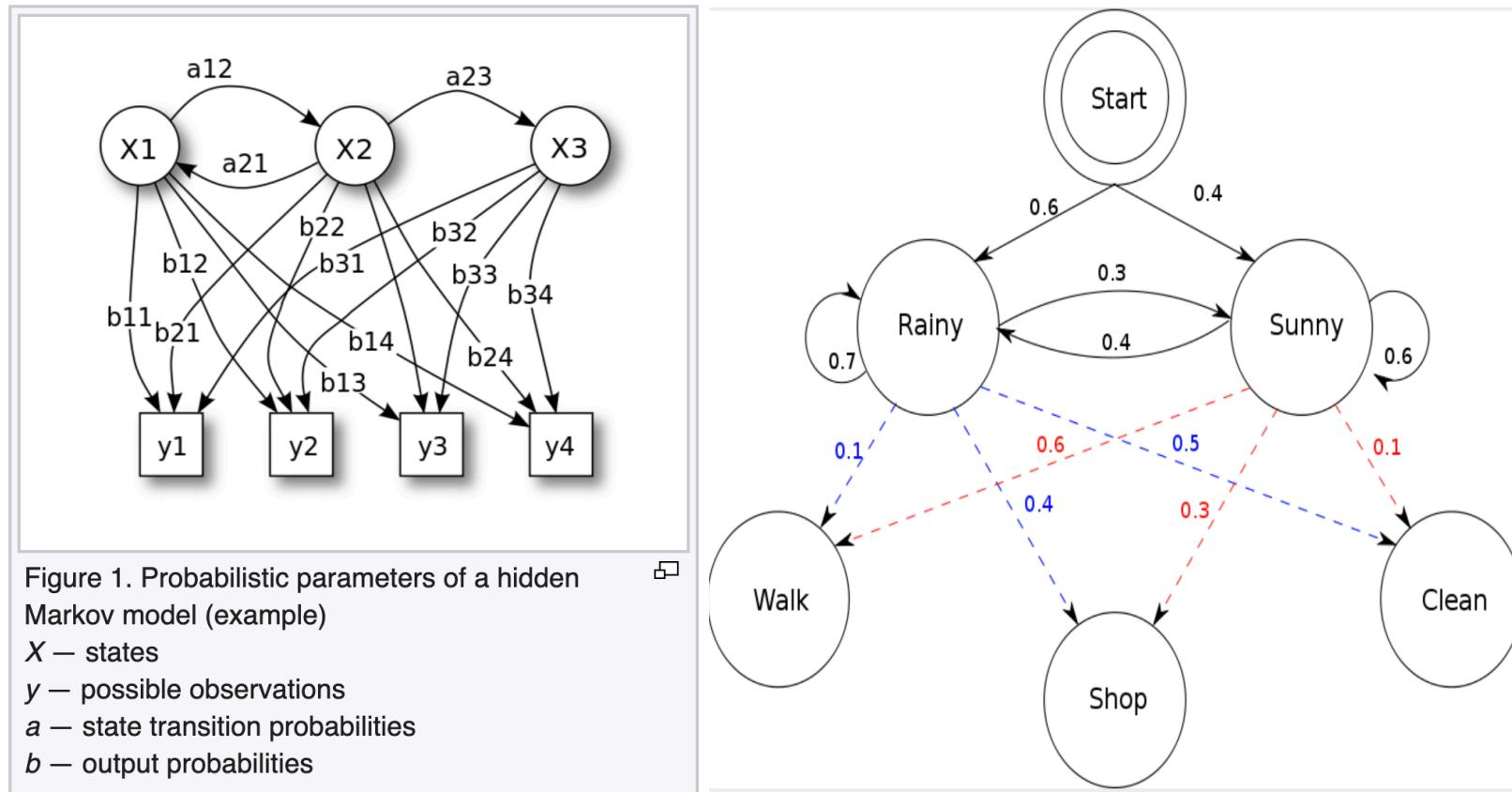
- Please write out the stochastic matrix using the above graph (**called a probability graphic model**).

In real life modeling, often the situation is much more complicated

- We need to consider global economic environment.
- There are a lot of **hidden things which are not directly observable**.

Study two examples on Wikipedia

https://en.wikipedia.org/wiki/Hidden_Markov_model



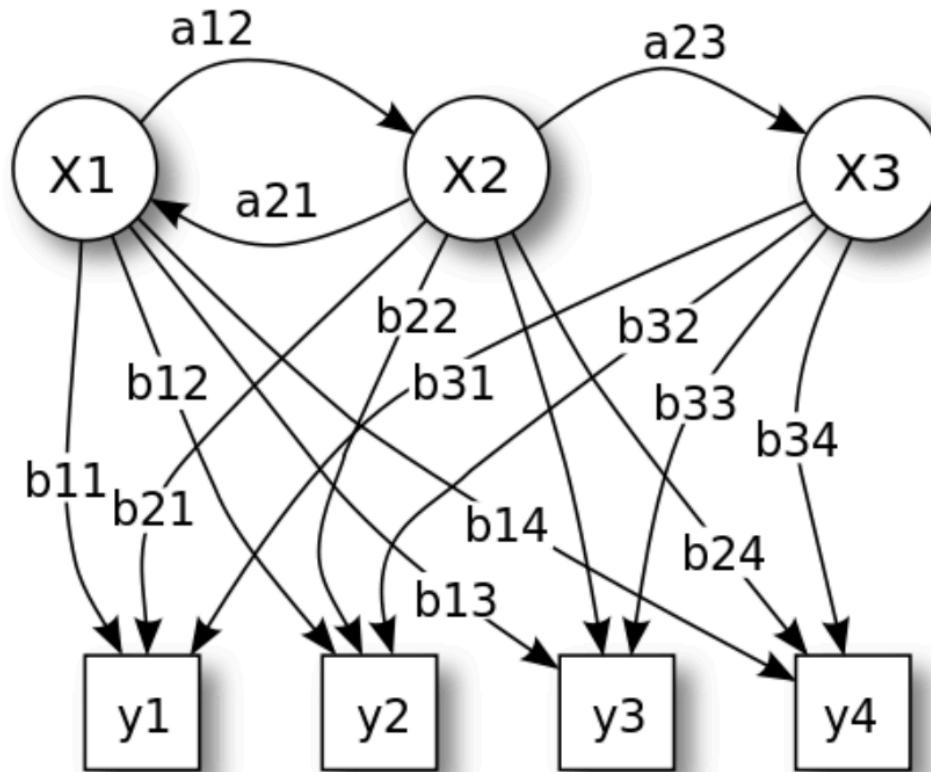
Hidden Markov Models

- Look an example on Wikipedia:

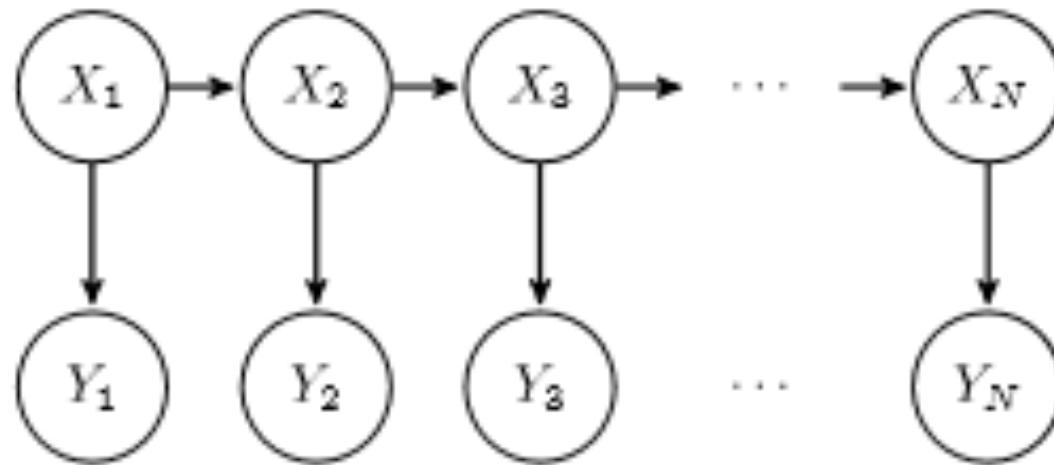
https://en.wikipedia.org/wiki/Hidden_Markov_model

$A = (a_{ij})$
= Transition Matrix

$B = (b_{kl})$
= Emission Matrix.



Hidden Markov Models



- Work out details with students on iPAD.
- Please see the detailed notes of HMM that I sent to you in email.

HMM is a typical example of a Probabilistic Graphical Model

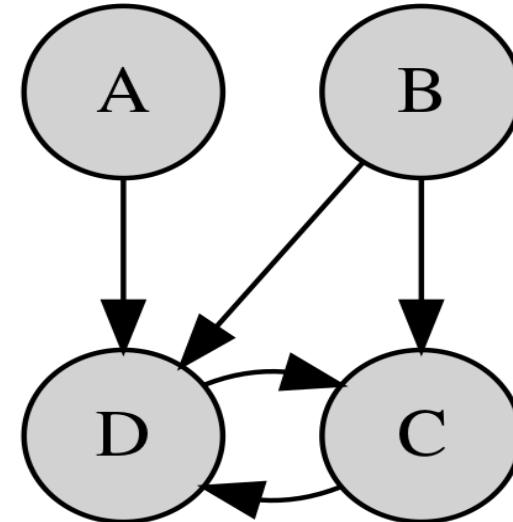
- What is a probabilistic Graphical Model?
- A probabilistic graphical model (PGM) is a probabilistic model for which a graph expresses the conditional dependence structure between random variables.

They are commonly used in probability theory, statistics—particularly Bayesian statistics—and machine learning.

Recall: $F: X \rightarrow Y$.

We say Y is a function of X , i.e. Y depends on X .

Note: the arrow starts from X and ends on Y .



An example of a graphical model. Each arrow indicates a dependency. In this example: D depends on A, B, and C; and C depends on B and D; whereas A and B are each independent.

Note: there are 3 arrows starts from A, B, C and ends on D. This means D depends on A, B, and C.

Why using Probabilistic Graphical Models

- Generally, probabilistic graphical models use a graph-based representation as the foundation for encoding a distribution over a multi-dimensional space and a graph that is a compact or factorized representation of a set of independences that hold in the specific distribution.

Recall: Chain rule for random variables

Two random variables [edit]

For two random variables X, Y , to find the joint distribution, we can apply the definition of conditional probability to obtain:

$$P(X, Y) = P(X|Y) \cdot P(Y)$$

More than two random variables [edit]

Consider an indexed collection of random variables X_1, \dots, X_n . To find the value of this member of the joint distribution, we can apply the definition of conditional probability to obtain:

$$P(X_n, \dots, X_1) = P(X_n | X_{n-1}, \dots, X_1) \cdot P(X_{n-1}, \dots, X_1)$$

Repeating this process with each final term creates the product:

$$P\left(\bigcap_{k=1}^n X_k\right) = \prod_{k=1}^n P\left(X_k \mid \bigcap_{j=1}^{k-1} X_j\right)$$

Recall: Probability Chain Rule for Events

The chain rule for two random events A and B says

$$P(A \cap B) = P(B | A) \cdot P(A).$$

For more than two events A_1, \dots, A_n the chain rule extends to the formula

$$P(A_n \cap \dots \cap A_1) = P(A_n | A_{n-1} \cap \dots \cap A_1) \cdot P(A_{n-1} \cap \dots \cap A_1)$$

which by induction may be turned into

$$P(A_n \cap \dots \cap A_1) = \prod_{k=1}^n P\left(A_k \mid \bigcap_{j=1}^{k-1} A_j\right).$$

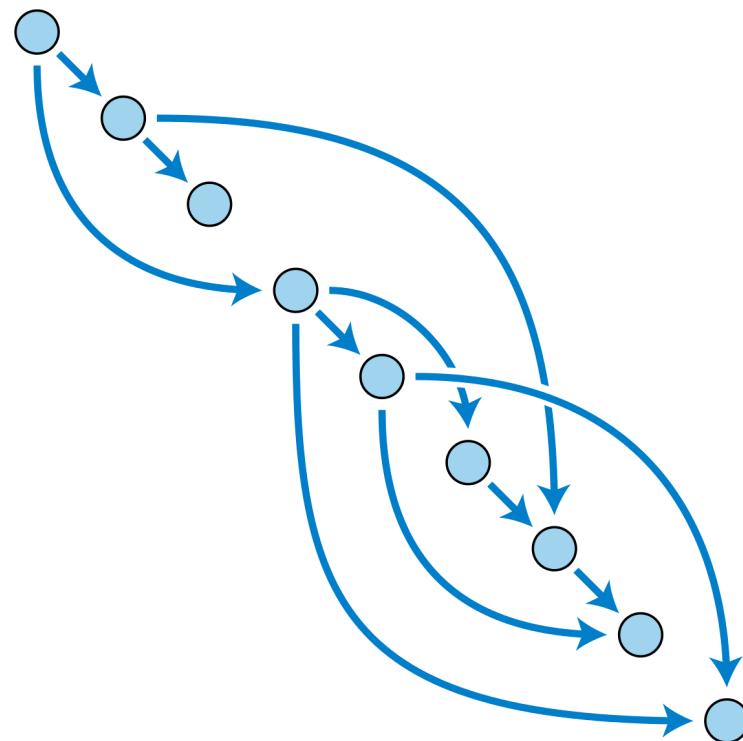
Example [edit]

With four events ($n = 4$), the chain rule is

$$\begin{aligned} P(A_4 \cap A_3 \cap A_2 \cap A_1) &= P(A_4 | A_3 \cap A_2 \cap A_1) \cdot P(A_3 \cap A_2 \cap A_1) \\ &= P(A_4 | A_3 \cap A_2 \cap A_1) \cdot P(A_3 | A_2 \cap A_1) \cdot P(A_2 \cap A_1) \\ &= P(A_4 | A_3 \cap A_2 \cap A_1) \cdot P(A_3 | A_2 \cap A_1) \cdot P(A_2 | A_1) \cdot P(A_1) \end{aligned}$$

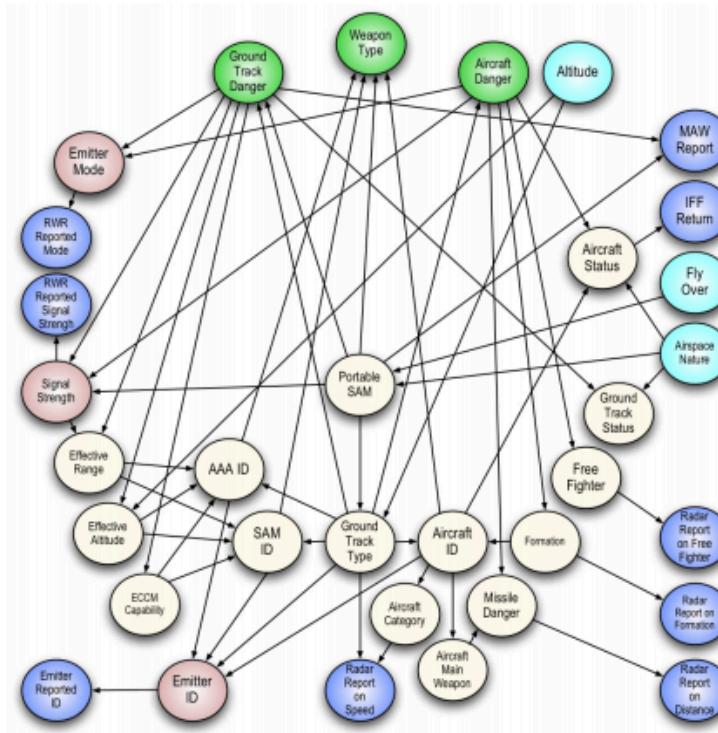
HMM is a typical example of Directed Acyclic Graph (DAG)

- A DAG is a finite directed graph with no directed cycles. That is, it consists of finitely many vertices and edges (also called *arcs*), with each edge directed from one vertex to another, such that there is no way to start at any vertex v and follow a consistently-directed sequence of edges that eventually loops back to v again. Equivalently, a DAG is a directed graph that has a topological ordering, a sequence of the vertices such that every edge is directed from earlier to later in the sequence.



There are many applications of DAG: Radar and Aircraft Control

- Modeling multiple planes and radar signals:



<https://pr-owl.org/basics/bn.php>

HMM and Directed Acyclic Graphical (DAG) Prob. Models

DAG models use a factorization of the joint distribution,

$$p(x_1, x_2, \dots, x_d) = \prod_{j=1}^d p(x_j | x_{\text{pa}(j)}),$$

where $\text{pa}(j)$ are the “parents” of node j .

This assumes a **Markov property** (generalizing Markov property in chains),

$$p(x_j | x_{1:j-1}) = p(x_j | x_{\text{pa}(j)}),$$

Note: Also can factor into blocks

Instead of factorizing by variables j , could **factor into blocks b** :

$$p(x) = \prod_b p(x_b | x_{\text{pa}(b)}),$$

and have the nodes be blocks.

- Usually assuming **full connectivity within the block**.

We will work out an example on HMM using iPAD on how to factor into blocks after the slides.

Review of Independence

- Let A and B are random variables taking values $a \in \mathcal{A}$ and $b \in \mathcal{B}$.
- We say that A and B are **independent** if we have

$$p(a, b) = p(a)p(b),$$

for all a and b .

- To denote independence of x_i and x_j we use the notation

$$x_i \perp x_j.$$

- In a product of Bernoullis, we assume $x_i \perp x_j$ for all i and j .

Review of Independence

- For independent a and b we have

$$p(a | b) = \frac{p(a, b)}{p(b)} = \frac{p(a)p(b)}{p(b)} = p(a).$$

- This gives us a more intuitive definition: A and B are independent if

$$p(a | b) = p(a)$$

for all a and $b \neq 0$.

- In words: knowing b tells us nothing about a (and vice versa).
 - This will tend to simplify calculations involving a .

- Useful fact: $a \perp b$ iff $p(a, b) = f(a)g(b)$ for some functions f and g .

Conditional Independence

- We say that A is **conditionally independent** of B given C if

$$p(a, b | c) = p(a | c)p(b | c),$$

for all a , b , and $c \neq 0$.

- Equivalently, we have

$$p(a | b, c) = p(a | c).$$

- “If you know C , then *also knowing B would tell you nothing about A* ”.
 - In mixture of Bernoullis, given cluster there is no dependence between variables.
- We often write this as
$$A \perp B | C.$$
- In a mixture of Bernoullis, we assume $x_i \perp x_j | z$ for all i and j .
 - This simplifies calculations involving x_i and x_j , provided that we know z .

Extra Conditional Independences in Markov Chains

- In Markov chains, the **Markov assumption** is $x_j \perp x_1, x_2, \dots, x_{j-2} \mid x_{j-1}$,

$$p(x_j \mid x_{j-1}, x_{j-2}, \dots, x_1) = p(x_j \mid x_{j-1}).$$

- But note that this **also implies** the additional conditional independence that

$$p(x_j \mid x_{j-2}, x_{j-3}, \dots, x_1) = p(x_j \mid x_{j-2}).$$

- We can use this property to easily compute $p(x_j \mid x_{j-2}, x_{j-3}, \dots, x_1)$:

$$\begin{aligned} p(x_j \mid x_{j-2}, x_{j-3}, \dots, x_1) &= p(x_j \mid x_{j-2}) \\ &= \sum_{x_{j-1}} p(x_j, x_{j-1} \mid x_{j-2}) \\ &= \sum_{x_{j-1}} p(x_j \mid x_{j-1}, x_{j-2}) p(x_{j-1} \mid x_{j-2}) \\ &= \sum_{x_{j-1}} \underbrace{p(x_j \mid x_{j-1})}_{\text{tran prob}} \underbrace{p(x_{j-1} \mid x_{j-2})}_{\text{tran prob}}. \end{aligned}$$

DAGs and Conditional Independence

- Conditional independences can substantially simplify inference.
- But it's **tedious** to formally show that the above are true.
 - See the last slide, and the EM notes.
- In DAGs we make the **conditional independence assumption** that
$$p(x_j \mid x_{j-1}, x_{j-2}, \dots, x_1) = p(x_j \mid x_{\text{pa}}(j)).$$
- Is there an easy way to find out what other independences are true?
 - If so, we could quickly find out which calculations are easy to do in a given DAG.

D-Separation: From Graphs to Conditional Independence

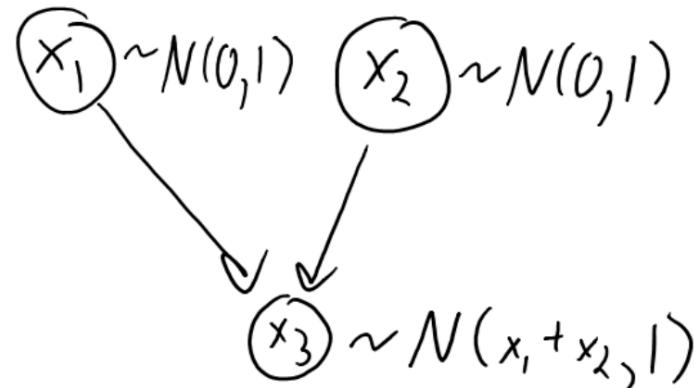
- All conditional independences implied by a DAG can be read from the graph.
- In particular: variables A and B are conditionally independent given C if:
 - “D-separation blocks all undirected paths in the graph from any variable in A to any variable in B .”
- In the special case of product of independent models our graph is:



- Here there are no paths to block, which implies the variables are independent.
- Checking paths in a graph tends to be faster than tedious calculations.
 - We can start connecting properties of graphs to computational complexity.

D-Separation as Genetic Inheritance

- The rules of d-separation are intuitive in a simple model of gene inheritance:
 - Each person has single number, which we'll call a "gene".
 - If you have no parents, your gene is a random number.
 - If you have parents, your gene is a sum of your parents plus noise.
- For example, think of something like this:



- Graph corresponds to the factorization $p(x_1, x_2, x_3) = p(x_1)p(x_2)p(x_3 | x_1, x_2)$.
 - In this model, does $p(x_1, x_2) = p(x_1)p(x_2)$? (Are x_1 and x_2 independent?)

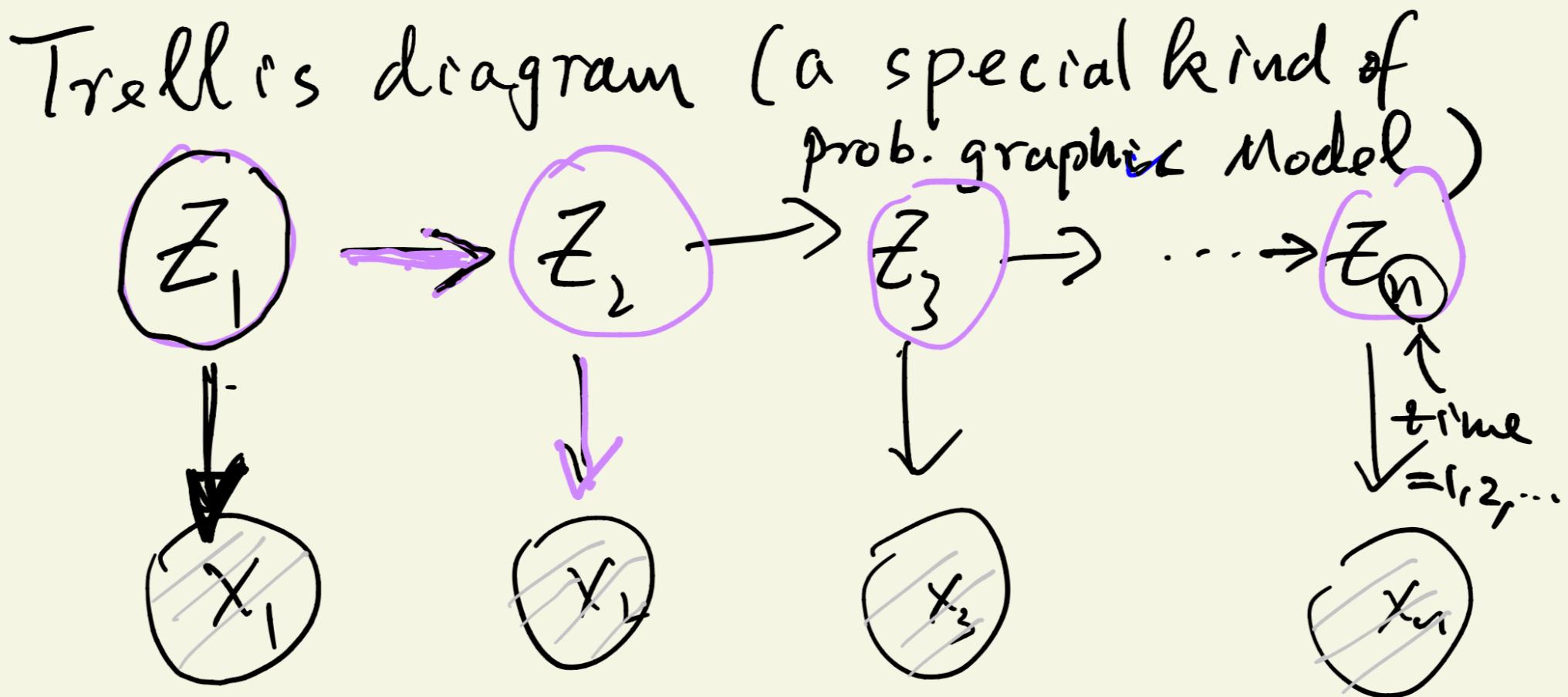
D-Separation as Genetic Inheritance

- Genes of people are **independent** if knowing one says nothing about the other.
- Your gene is **dependent on your parents**:
 - If I know you your parent's gene, I know something about yours.
- Your gene is **independent of your (unrelated) friends**:
 - If know you your friend's gene, it doesn't tell me anything about you.
- Genes of people can be **conditionally independent** given a third person:
 - Knowing your grandparent's gene tells you something about your gene.
 - But grandparent's gene isn't useful if you know parent's gene.

Hidden Markov Models

- Work out details with students on iPAD.
- Please see the detailed notes of HMM that I sent to you in email.

Hidden Markov Model (HMM)



$\underbrace{z_1, \dots, z_n}_{\text{Hidden variables}} \in \{1, 2, \dots, m\}$

$x_1, \dots, x_n \in \mathcal{X} (\text{discrete, } \mathbb{R}, \mathbb{R}^d, \dots)$
 "Observable" data

Key: The Joint prob factorized in following way:

$$\begin{aligned}
 & P(z_1, z_2, \dots, z_n, x_1, \dots, x_n) \\
 & \underset{\text{Initial prob.}}{\approx} P(z_1) \underset{\text{Emission prob.}}{\circled{P(x_1 | z_1)}} \prod_{k=2}^n P(z_k | z_{k-1}) \underset{\text{Transition prob.}}{\circled{P(x_k | z_k)}}
 \end{aligned}$$