# INT104 ARTIFICIAL INTELLIGENCE

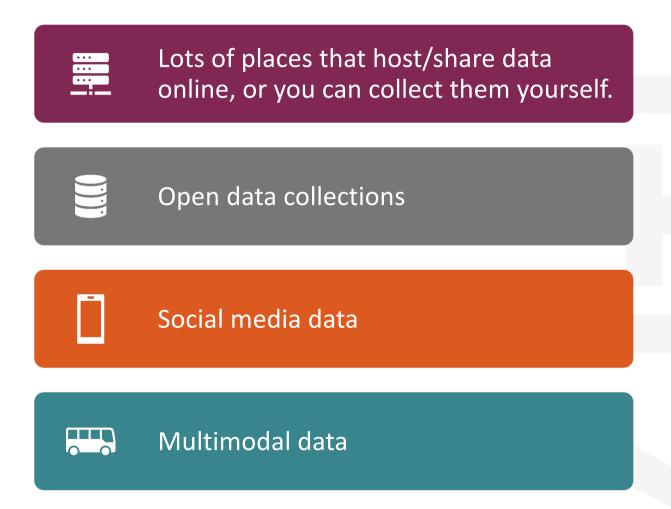
## **REVIEW – WEEK 14**

Sichen Liu Sichen.Liu@xjtlu.edu.cn





## **Data Collection**





## **Data Cleaning**

- Handling Missing Data
  - Get rid of the corresponding instance.
  - Get rid of the whole column.
  - Set the values to some value (zero, the mean, the median, etc.).

- Smooth Noisy Data
  - Identify or remove the outliners
  - Try to resolve the inconsistent

(there is no one way to remove noise, or smooth out the noisiness in the data)



# **Practice: Data Cleaning**

#	Country	Alcohol	Deaths	Heart	Liver	Free
		(L/person)	(Per 100k)	(Per 100k)	(Per 100k)	healthcare
1	Australia	2.5	785	211	15.30000019	Υ
2	Austria	3.00000095	863	167	45.59999847	Υ
3	Belg/Lux	2.900000095	883	131	20.70000076	N
4	Canada	2.400000095	793	NA	16.39999962	Υ
5	Denmark	2.900000095	971	220	23.89999962	Υ
6	Finland	0.80000012	970	297	19	N
7	France	9.100000381	751	11	37.90000153	N
8	Iceland	-0.80000012	743	211	11.19999981	Υ
9	Ireland	0.69999988	1000	300	6.5	Υ
10	Israel	0.600000024	-834	183	13.69999981	Υ
11	Italy	27.900000095	775	107	42.20000076	Υ
12	Japan	1.5	680	36	23.20000076	N
13	Netherlands	1.79999952	773	167	9.199999809	N
14	New Zealand	1.899999976	916	266	7.699999809	Υ
15	Norway	0.0800000012	806	227	12.19999981	N
16	Spain	6.5	724	NA	NA	Υ
17	Sweden	1.600000024	743	207	11.19999981	N
18	Switzerland	5.800000191	693	115	20.29999924	N
19	UK	1.29999952	941	285	10.30000019	Y
20	US	1.200000048	926	199	22.10000038	N
21	West Germany	2.70000048	861	172	36.70000076	Υ



## **Data Transformation**

- Normalization
  - Min–max normalization.

$$x_{scaled} = rac{x - x_{min}}{x_{max} - x_{min}}$$

Z-score normalization.

Normalizing every value in a dataset such that the mean of all of the values is 0 and the standard deviation is 1

$$x_{scaled} = rac{x-mean}{sd}$$

Normalization by decimal scaling.

$$x_{scaled} = \frac{x}{10^j}$$



## **Data Reduction**

Data reduction is a key process in which a reduced representation of a dataset that produces the same or similar analytical results is obtained.

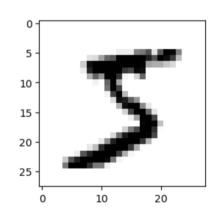
	A1	A2	A3	 A200			A1	A2	А3	 A120
T1						T1				
T2					-	T2				
Т3						T3				
						T150				
T200										

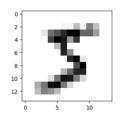


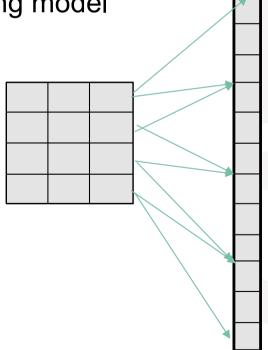
# **Dimensionality Reduction**

## Data with high dimensions:

- High computational complexity
- May contain many irrelevant or redundant features
- Difficulty in visualization
- With high risk of getting an overfitting model









# Principal Component Analysis (PCA)

### Preserving the Variance:

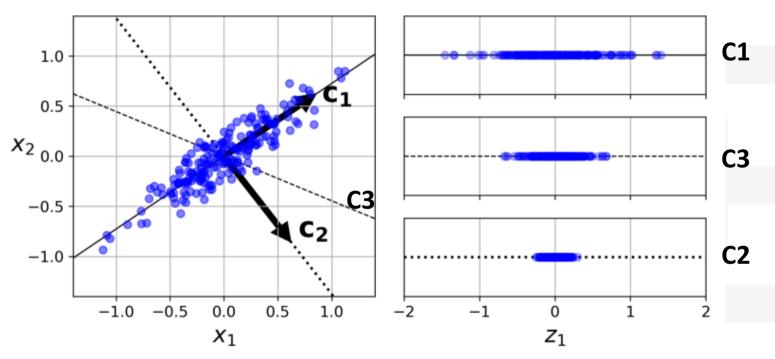
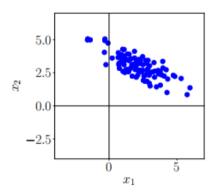


Figure 8-7. Selecting the subspace to project on

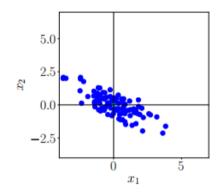
PCA identifies the axis that accounts for the largest amount of variance in the training set.

## **PCA**

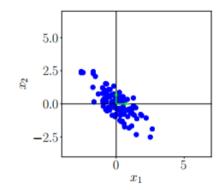
### Key steps of PCA in practice



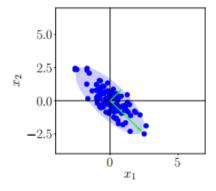
(a) Original dataset.



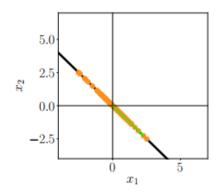
(b) Step 1: Centering by subtracting the mean from each data point.



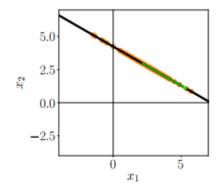
(c) Step 2: Dividing by the standard deviation to make the data unit free. Data has variance 1 along each axis.



(d) Step 3: Compute eigenvalues and eigenvectors (arrows) of the data covariance matrix (ellipse).



(e) Step 4: Project data onto the principal subspace.



(f) Undo the standardization and move projected data back into the original data space from (a).



## **API Information**

## numpy.linalg.norm

linalg.norm(x, axis=None): Matrix or vector norm.

#### **Parameters**

**x**: array like

Input array. If axis is None, x must be 1-D or 2-D.

#### **Returns**

**n**: float or ndarray

Norm of the matrix or vector(s).



## **API Information**

#### sklearn.cluster.KMeans

class sklearn.cluster.KMeans(n\_clusters=8, random\_state=None): KMeans clustering.

#### **Parameters**

n\_clusters: int, default=8

The number of clusters to form as well as the number of centroids to generate random\_state: int, RandomState instance or None, default=None Determines random number generation for centroid initialization. Use an int to make the randomness deterministic.

#### **Attributes**

labels\_: ndarray of shape (n samples,), Labels of each point.

#### **Methods**

fit(X), Compute k-means clustering.

#### **Parameters**

**X:** array-like, sparse matrix of shape (n samples, n features) Training instances to cluster.

#### **Returns**

self: object

Fitted estimator.



## **Indentation matters!**

- Code is grouped by its indentation
- Indentation is the number of whitespace or tab characters before the code.
- If you put code in the wrong block, then you will get unexpected behavior

```
Line 1
Line 2
Line 2
Line 3
Line 4
Line 5
Line 6

Executing else")

print("Executing else")

print("Prints regardless of the if-else block")

Executing if

Prints regardless of the if-else block
```



