# CS167: Machine Learning

k-Nearest Neighbor (k-NN) Handling Missing Data Data Normalization

Thursday, February 15th, 2024



#### Announcements

- <u>Notebook #2: kNN and Normalization</u> is released today
  - due Wednesday 02/21 by 11:59pm
  - to submit, download the ipynb file from Colab
  - directly upload to CodePost
- Heads up that Quiz #1
  - will be released on Tuesday 02/20 after class
  - will be due Tuesday 02/27 by 11:59pm

Before we get started, let's load in our datasets:

- Make sure you change the path to match your Google Drive.
  - Load the titanic.csv file from your Google Drive

[2] #run this cell if you're using Colab: from google.colab import drive drive.mount('/content/drive')



# Today's Agenda

- Topics:
  - kNN Implementation using Pandas

Missing Data

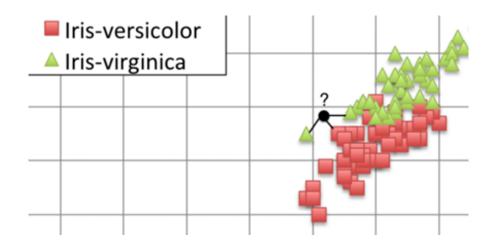
Normalization

#### 3-Nearest Neighbor (3-NN)

- 3-Nearest-Neighbor Algorithm: predict the most commonly appearing class among the 3 closest training examples
  - In other words, k=3
- Let's assume this subset of Iris has only 2 classes (even number): Iris-versicolor

Iris-virginica

• What class will a **3NN** algorithm predict?



#### k-Nearest Neighbor (kNN)

- **k-Nearest-Neighbor** predict the most commonly occurring class of the *k nearest neighbors*.
  - 1. Calculate the distance between the new point (e.g. the Iris we would like to make a prediction on), and the existing training examples.
  - 2. Sort the data by the newly calculated distance so that the nearest training examples are first
  - 3. Take the top k neighbors:
    - if the problem is a *classification*, **take the mode of the target variable** to find the most commonly appearing class and return that as your prediction
    - if the problem is a *regression*, **take the average of the target variables** for the k closest neighbors and return that as your prediction

### k-NN Implementation in Python/Pandas

- Let's build a 5-Nearest-Neighbor Iris classifier from scratch using our Pandas/Python skills:
- To implement this 5NN, we need to do 3 things:
  - 1. Calculate the distances from each of the rows to the new instance
  - 3. Sort the data by these distances
  - 5. Select the k closest training examples and use them to predict the most commonly occurring class of the closest neighbors.

#### Step 1: Calculate the Distances

- Let's start by adding a new column to our iris DataFrame that is the distance from each existing row to the new instance with:
  - 5.1 petal length, 7.2 sepal length, 1.5 petal width, and 2.5 sepal width
  - The syntax for adding a new column is as follows:

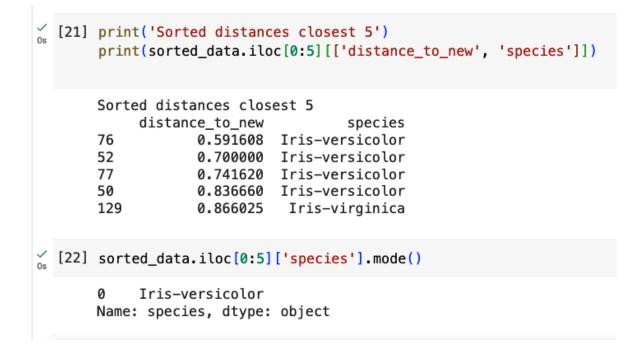
```
• df['new col name'] = _____
```

#### Step 2: Sort the data by the Distances

- Let's now sort our data using the built in sort\_values() function.
   [documentation]
- We want to find the nearest k neighbors, so sorting them in ascending order (which is the default setting for sort\_values() will work nicely.

```
[10] k=15
     sorted_data = iris.sort_values(['distance_to_new'])
     sorted data.head() #shortest distances first
                                                                                                           Ħ
           sepal length sepal width petal length petal width
                                                                            species distance to new
      76
                       6.8
                                     2.8
                                                     4.8
                                                                    1.4 Iris-versicolor
                                                                                                0.591608
                                                                                                           ıl.
       52
                       6.9
                                     3.1
                                                                        Iris-versicolor
                                                     4.9
                                                                    1.5
                                                                                                0.700000
                       6.7
                                                                    1.7 Iris-versicolor
      77
                                     3.0
                                                     5.0
                                                                                                0.741620
       50
                      7.0
                                     3.2
                                                     4.7
                                                                    1.4 Iris-versicolor
                                                                                                0.836660
      129
                      7.2
                                     3.0
                                                     5.8
                                                                    1.6
                                                                          Iris-virginica
                                                                                                0.866025
```

# Step 3: Display the most common species among these 5



• And Viola! We have successfully implemented our first machine learning model from scratch.

#### k-NN All Steps

iris.head()

[10] k=**15** 

sorted\_data = iris.sort\_values(['distance\_to\_new'])
sorted\_data.head() #shortest distances first

Sorted d	istances close	est 5
dis	tance_to_new	species
76	0.591608	Iris-versicolor
52	0.700000	Iris-versicolor
77	0.741620	Iris-versicolor
50	0.836660	Iris-versicolor
129	0.866025	Iris-virginica

```
[22] sorted_data.iloc[0:5]['species'].mode()
```

0 Iris-versicolor
Name: species, dtype: object

#### **Programming Exercise:**

- Rewrite k-NN code so that it's a function.
- Pass the iris measurements (specimen), DataFrame, and k as parameters and return the predicted class.

```
def kNN(specimen, data, k):
    # write your code in here to make this function work
    # 1. calculate distances

    # 2. sort
    # 3. predict
    return prediction
```

#### **Programming Exercise:**

- Rewrite k-NN code so that it's a function.
- Pass the iris measurements (specimen), DataFrame, and k as parameters and return the predicted class.

```
new_iris = {}
new_iris['petal length'] = 5.1
new_iris['sepal length'] = 7.2
new_iris['petal width'] = 1.5
new_iris['sepal width'] = 2.5
# call the function you just wrote
kNN(new_iris, iris, 15)
```

# Today's Agenda

- Topics:
  - kNN Implementation using Pandas

Missing Data

Normalization

- Most datasets you will work with will not be in perfect shape
  - you'll need to "clean" the data before you can run any machine learning algorithms on it.
- Missing data is a pretty common thing so much so that there's a special value for missing data:
  - NaN, or not a number.

- The steps of cleaning data normally include:
  - Step 1: Detecting which columns have missing data
  - Step 2: Determining how much data is missing in each column
  - Step 3: Deciding what to do with the missing data:
    - drop it
    - fill it
    - let it be

- Notice, in the deck column, there are 3 instances of NaN we can see...
- But what about the other 800 or so rows? Do we have to go through and find them manually?

ti	tanic.head	()										$\frown$	
	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town
0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton
1	1	1	female	38.0	1	0	71.2833	С	First	woman	False	С	Cherbourg
2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampton
3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	С	Southampton
4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampton

- In order to identify missing data, we will use a combination of three Pandas functions:
  - **isna()** <u>https://pandas.pydata.org/docs/reference/api/pandas.isna.html</u>
  - **notna()** <u>https://pandas.pydata.org/docs/reference/api/pandas.notna.html</u>
  - **any()** <u>https://pandas.pydata.org/docs/reference/api/pandas.DataFrame.any.html</u>

- Using isna() and notna() to find missing data:
  - isna(): will return a boolean series where it is **True** if the element is **NaN**
  - notna(): will return a boolean series where it is True if the element is not NaN

	API reference > General functions > pandas.isna
Input/output	
General functions ^	pandas.isna
pandas.melt	
pandas.pivot	<pre>pandas.isna(obj) [source]</pre>
pandas.pivot_table	Detect missing values for an array-like object.
pandas.crosstab	This function takes a scalar or array-like object and indicates whether values are missing
pandas.cut	(NaN in numeric arrays, None or NaN in object arrays, NaT in datetimelike).
pandas.qcut	Parameters:
pandas.merge	<b>obj</b> : scalar or array-like
pandas.merge_ordered	Object to check for null or missing values.

https://pandas.pydata.org/docs/reference/api/pandas.isna.html

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titanic.loc[0:4]

S	urvived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town
0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton
1	1	1	female	38.0	1	0	71.2833	С	First	woman	False	С	Cherbourg
2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southamptor
3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	С	Southamptor
4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	NaN	Southamptor

• Now, let's call isna(), and see what we get as an output

0		anic.loc[ ook at the			n									
		survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town
	0	False	False	False	False	False	False	False	False	False	False	False	True	False
	1	False	False	False	False	False	False	False	False	False	False	False	False	False
	2	False	False	False	False	False	False	False	False	False	False	False	True	False
	3	False	False	False	False	False	False	False	False	False	False	False	False	False
	4	False	False	False	False	False	False	False	False	False	False	False	True	False

• **isna()** is pretty nifty but there should be better way to summarize this.

• any()

pandas.DataFrame.any	
DataFrame. <mark>any</mark> (*, axis=0, bool_only=False, skipna=True, **kwargs)	[source
Return whether any element is True, potentially over an axis.	

Returns False unless there is at least one element within a series or along a Dataframe axis

https://pandas.pydata.org/docs/reference/api/pandas.DataFrame.any.html

#### Step 1: Identifying Missing Data

 Let's use any() on the call to isna() we just did to let us know which columns have missing data:

	titanic.isna()	.any()	
	survived	False	
	pclass	False	
	sex	False	
(	age	True	
-	sibsp	False	
	parch	False	
	fare	False	
C	embarked	True	$\bigcirc$
	class	False	
	who	False	
_	adult_male	False	
(	deck	True	
(	embark_town	True	$\bigcirc$
	alive	False	
	alone	False	
	dtype: bool		
	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		

- Several columns are missing data: age, embarked, deck, and embark\_town.
- Wouldn't it be great to know how much data is missing in each of those columns?

- The steps of cleaning data normally include:
  - Step 1: Identifying which columns have missing data
  - Step 2: Determining how much data is missing in each column
  - Step 3: Deciding what to do with the missing data:
    - drop it
    - fill it
    - let it be

- To decide how to handle our missing data, it's important to know how much missing data each column has:
  - If the missing data is a small proportion of the data, we choose to drop those rows completely from the dataset
  - However, if most of the rows are missing data for a specific column, maybe it's a sign that we don't need to use that column
- There are multiple ways of doing this, but one of the quickest/easiest is using value\_counts()

 Great, so now that we know which columns are missing data, let's check to see how much data they are missing using value\_counts()

#### pandas.Series.value\_counts

Series.value\_counts(normalize=False, sort=True, ascending=False,

#### bins=None, dropna=True)

[source]

Return a Series containing counts of unique values.

The resulting object will be in descending order so that the first element is the most frequently-occurring element. Excludes NA values by default.

 Let's apply value\_counts() on the various columns (eg, deck) of Titanic dataset

0			k.value g value	_	lropna= <mark>False</mark> )
	NaN C B D E A	688 59 47 33 32 15			
	F G	13 4			
	Name:	deck,	dtype:	int64	

 Let's apply value\_counts() on the various columns (eg, age) of Titanic dataset

0			e.value_ ng value		ts <mark>(</mark> drop	na=False)
Đ	NaN 24.00 22.00 18.00 28.00 36.50 55.50 0.92 23.50 74.00		77 30 27 26 25 1 1 1 1 1			
		age,	Length:	89,	dtype:	int64

https://pandas.pydata.org/docs/reference/api/pandas.Series.value\_counts.html

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 Let's apply value\_counts() on the various columns (eg, embarked) of Titanic dataset

0		ic.embarked.value_counts(dropna=False) ssing values
	S C Q NaN Name:	644 168 77 2 embarked, dtype: int64

 Let's apply value\_counts() on the various columns (eg, embark\_town) of Titanic dataset

0	<pre>titanic.embark_town.value_counts(dropna=False) #2 missing values</pre>
	Southampton 644 Cherbourg 168 Queenstown 77 NaN 2 Name: embark_town, dtype: int64

• So, here is our results using value\_counts()

Column	Num Rows Missing
deck	688
age	177
embarked	2
embark_town	2

• Now with this new information, it's up to us to decide what to do with these missing values

- The steps of cleaning data normally include:
  - Step 1: Identifying which columns have missing data
  - Step 2: Determining how much data is missing in each column

- Step 3: Deciding what to do with the missing data:
  - **drop it:** drop the missing data from the dataset (either col or row)
  - fill it: fill the missing data with a suitable replacement
  - let it be: let it be and cross our fingers

### Option 1: Drop it using dropna()

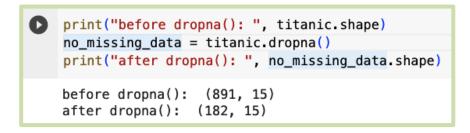
• If there isn't much missing data, and/or you have a very large dataset, dropping the row that includes the missing data is a viable option.

```
print("before: ", titanic.shape)
titanic.dropna()
print("after: ", titanic.shape)
before: (891, 15)
after: (891, 15)
```

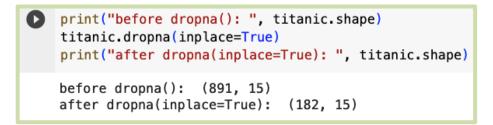
• We know that there's missing data, why didn't the shape change?

#### Option 1: Drop it using dropna()

- Pandas is trying to protect you, and rather than dropping the rows "in place", it is returning a DataFrame with the rows dropped--as written, we're just not saving it's return. There are two ways to fix this:
  - save what dropna() is returning in a variable



• add the parameter inplace=True to the function call, and it will drop the rows in the original dataset (be careful with this one)



#### Option 1: Drop it using dropna()

• That's better, but wow, most of our dataset is gone now if we drop all of the rows that have missing data. If this happens to you, you'll probably want to re-load your data to have the full dataset to work with.

# if that happens, you'll want to re-run your data loading code: path = '/content/drive/MyDrive/cs167\_fall23/datasets/titanic.csv' titanic = pd.read\_csv(path)

### Option 2: Fill it using fillna()

- If dropping all of the data will make your dataset too sparse, consider filling the missing values with something else.
- What do you think we should use to fill in the missing data in the age column?
  - we probably don't want to throw off our statistics...

ιII															
	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town	alive	alone
0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton	no	False
1	1	1	female	38.0	1	0	71.2833	С	First	woman	False	С	Cherbourg	yes	False
2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampton	yes	True
3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	С	Southampton	yes	False
4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampton	no	True
5	0	3	male	NaN	0	0	8.4583	Q	Third	man	True	NaN	Queenstown	no	True
6	0	1	male	54.0	0	0	51.8625	S	First	man	True	E	Southampton	no	True

#### titanic.head(7)

#### Option 2: Fill it using fillna()

- What do you think we should use to fill in the missing data in the age column?
  - we probably don't want to throw off our statistics...

```
print("before: ", titanic['age'].isna().any())
age_mean = titanic['age'].mean()
titanic['age'].fillna(age_mean, inplace=True)
print("after: ", titanic['age'].isna().any())
titanic.head(7)
```

	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town	alive	alone
0	0	3	male	22.000000	1	0	7.2500	S	Third	man	True	NaN	Southampton	no	False
1	1	1	female	38.000000	1	0	71.2833	С	First	woman	False	С	Cherbourg	yes	False
2	1	3	female	26.000000	0	0	7.9250	S	Third	woman	False	NaN	Southampton	yes	True
3	1	1	female	35.000000	1	0	53.1000	S	First	woman	False	С	Southampton	yes	False
4	0	3	male	35.000000	0	0	8.0500	S	Third	man	True	NaN	Southampton	no	True
5	0	3	male	29.699118	0	0	8.4583	Q	Third	man	True	NaN	Queenstown	no	True
6	0	1	male	54.000000	0	0	51.8625	S	First	man	True	Е	Southampton	no	True

## Option 3: Let it be

- What's so bad about missing data? Why do we care if some data is missing?
- What happens if we try to do math with NaN? Try it out for yourself:
  - Go to the bottom of the Day05\_Missing\_Data\_Normalization.ipynb and try out

# Summary: Missing Data

- The steps of cleaning data normally include:
  - Step 1: Detecting which columns have missing data
  - Step 2: Determining how much data is missing in each column
  - Step 3: Deciding what to do with the missing data:
    - drop it
    - fill it
    - let it be

# Summary: Missing Data Functions

- isna(): returns True for any missing data
- notna(): returns True for any data that is not NaN
- any(): returns true if any of the elements in a Series is True
- value\_counts(): returns a list of the values in a Series, use dropna=False to see NaN values
- dropna(): drops rows or columns (specify which axis, 1 or 0) that have missing data. Don't forget to either save the result of the call or add inplace=True as a parameter
- fillna(): replaces missing data with a given value (generally 0 or the mean)

# Today's Agenda

- Topics:
  - kNN Implementation using Pandas

Missing Data

• Normalization

#### Normalization

- Normalizing data:
  - rescale attribute values so they're about the same
  - adjusting values measured on different scales to a common scale

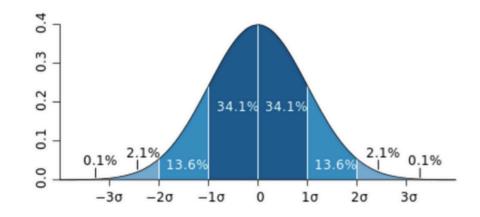
## A Simple Normalization:

- One simple method of normalizing data is to replace each value with a proportion relative to the max value.
- For example, the oldest person on the Titanic was 80, so:

age	replaced by
80	80/80 = 1
50	50/80 = 0.625
48	48/80 = 0.6
25	25/80 = 0.3125
4	4/80 = 0.05

# **Z-Score: Another Normalization Method**

- Idea: rather than normalize to proportion of max, normalize based on how many standard deviations they are away from the mean
- Standard Deviation: usually represented as  $\sigma$ (sigma), a kind of 'average' distance from the average value
  - a low standard deviation indicates that the values tend to be close to the mean
  - a high standard deviation indicates that the values are spread out over a wider range



Standard Deviation:

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#### **Standard Deviation Calculation**

- Standard Deviation: usually represented as  $\sigma$ (sigma), a kind of 'average' distance from the average value
  - Find the mean, represented as  $\mu$ :**MU**
  - Then, for each number, subtract the mean and square the result
  - Then, find the mean of those squared differences
  - Take the square root of that and we are done

• Let  $\mu$  be the mean, then standard deviation of  $x_1$ ,  $x_2$ ,...,  $x_N$  is:

$$\sigma = \sqrt{\frac{(x_1 - \mu)^2 + (x_2 - \mu)^2 + \dots + (x_N - \mu)^2}{N}}$$

### **Corrected Sample Standard Deviation**

• **Bessel's correction** says that you should divide by N-1 instead of N when working with a sample (as we usually do in machine learning tasks), and your estimate will be a little less biased.

$$\sigma = \sqrt{\frac{(x_1 - \mu)^2 + (x_2 - \mu)^2 + \dots + (x_N - \mu)^2}{N - 1}}$$

# Computing the Z-Score

• After computing the corrected sample standard deviation, to normalize, replace each value  $x_i$  with it's **Z-Score** based on the mean ( $\mu$ ) and standard deviation ( $\sigma$ ) of it's column.

$$Z-score: \frac{x_i-\mu}{\sigma}$$

# Computing the Z-Score

- For example: On the Titanic:
  - sex mean(0:male, 1:female): 0.35
  - sex standard deviation: 0.48
  - age mean: 29.7
  - age standard deviation: 13

$$Z-score: \frac{x_i-\mu}{\sigma}$$

	sex	age
example 1	1	50
example 2	0	48

	sex	age
example 1	1	50
example 3	1	25

Z-Score for male:  $(0 - 0.35)/0.48 \approx -0.73$ Z-Score for female:  $(1 - 0.35)/0.48 \approx 1.35$ Z-Score for age 50:  $(50 - 29.7)/13 \approx 1.56$ Z-Score for age 48:  $(48 - 29.7)/13 \approx 1.41$ Z-Score for age 25:  $(25 - 29.7)/13 \approx -0.36$ 

## **Distance Computation Before** Normalization

	sex	age
example 1	1	50
example 2	0	48

dist

ance: 
$$\sqrt{(1-0)^2 + (50-48)^2} \approx 2.24$$

	sex	age
example 1	1	50
example 3	1	25

distance:  $\sqrt{(1-1)^2 + (50-25)^2} = 25$ 

## Distance Computation After Normalization

	sex	age
example 1	1.35	1.56
example 2	-0.73	1.41

$$\begin{array}{c} \text{distance:} \\ \sqrt{(1.35 - -0.73)^2 + (1.56 - 1.41)^2} \\ \approx 2.09 \end{array}$$

	sex	age
example 1	1.35	1.56
example 3	1.35	-0.36

$$\begin{array}{l} \text{distance:} \\ \sqrt{(1.35-1.35)^2+(1.56--0.36)^2} \\ = 1.92 \end{array}$$

#### Computing the Z-Score on Titanic

 Called on a dataframe, will replace values given in to\_replace with value. Let's use this to make the sex column of the dataset numeric.

titanic['sex'] = titanic['sex'].replace(to\_replace='female', value=1)
titanic['sex'] = titanic['sex'].replace(to\_replace='male', value=0)
titanic.head()

	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town	alive	alone
0	0	3	0	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton	no	False
1	1	1	1	38.0	1	0	71.2833	С	First	woman	False	С	Cherbourg	yes	False
2	1	3	1	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampton	yes	True
3	1	1	1	35.0	1	0	53.1000	S	First	woman	False	С	Southampton	yes	False
4	0	3	0	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampton	no	True

#### Computing the Z-Score on Titanic

 Now that we have the data as 1s and 0s, let's calculate the mean and standard deviation

s\_mean = titanic.sex.mean()
s\_std = titanic.sex.std()

#replace column with each entry's z-score
titanic.sex = (titanic.sex - s\_mean)/s\_std
titanic.head()

 $Z - score : \frac{x_i - \mu}{\sigma}$ 

	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town	alive	alone
0	0	3	-0.734928	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton	no	False
1	1	1	1.359146	38.0	1	0	71.2833	С	First	woman	False	С	Cherbourg	yes	False
2	1	3	1.359146	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampton	yes	True
3	1	1	1.359146	35.0	1	0	53.1000	S	First	woman	False	С	Southampton	yes	False
4	0	3	-0.734928	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampton	no	True