## **Discussion 4**

**DSC 80** 

2024-04-26

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## Section 1

## FA23 Midterm Problem 3

### **Problem**

We have a DataFrame donkeys with 544 rows of measurements. The WeightAlt column has values for 30 donkeys that were **reweighed** a day later.

		id	BCS	Age	Weight	WeightAlt
	0	d01	3.0	<2	77	NaN
	1	d02	2.5	<2	100	NaN
	2	d03	1.5	<2	74	NaN

1d	A unique identifier for each donkey (d01, d02,
	etc.).
BCS	Body condition score: from 1 (emaciated) to 3
	(healthy) to 5 (obese) in increments of 0.5.
Age	Age in years: <2, 2–5, 5–10, 10–15, 15–20, and
	over 20 years.
Weight	Weight in kilograms.

Second weight measurement taken for 30 donkeys. NaN if the donkey was not reweighed.

We have to look at the following scenarios and tell if the missingess for WeightAlt is NMAR, MAR, or MCAR.

WeightAlt

# Refresher on Missingness Types

- Missing By Design (MD): Missingness is exactly determined by other columns, for intended reasons
- **Not missing at random (NMAR)**: Missingness depends on the values of the column that's missing
- Missing at random (MAR): Missingness depends on the values of other columns that are present
- Missing completely at random (MCAR): Missingness in a column does not depend on the values in that column, or in other columns

If you're curious, the terminology comes from Rubin, Donald B. "Inference and missing data." Biometrika 63.3 (1976): 581-592.

### Part A

• The researchers chose the 30 donkeys with the largest Weight values to reweigh.

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#### Part A

- The researchers chose the 30 donkeys with the largest Weight values to reweigh.
- This is MAR: The missingness depends on the values in the Weight column.

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#### Part B

• The researchers drew 30 donkeys uniformly at random without replacement from the donkeys with BCS score of 4 or greater.

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- The researchers drew 30 donkeys uniformly at random without replacement from the donkeys with BCS score of 4 or greater.
- This is MAR: The missingness depends on the values in the BCS column. The pattern of that dependence is different, but it's still MAR.

#### Part C

• The researchers set i as a number drawn uniformly at random between 0 and 514, then reweighed the donkeys in donkeys.iloc[i:i+30].

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- This one... could go either way. The intended answer was MCAR because this process means we're selecting a random length-30 slice of the data to reweigh.

### Part C

- The researchers set i as a number drawn uniformly at random between 0 and 514, then reweighed the donkeys in donkeys.iloc[i:i+30].
- This one... could go either way. The intended answer was MCAR
  because this process means we're selecting a random length-30 slice of
  the data to reweigh.
- But you might notice that with this procedure, not every value has an equal likelihood of being in that slice, so you could also say it's MAR based on index/position.

#### Part D

• The researchers reweighed all the donkeys, but deleted all the values in WeightAlt except for the 30 lowest values.

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- The researchers reweighed all the donkeys, but deleted all the values in WeightAlt except for the 30 lowest values.
- This is **NMAR**: The missingness depends on the values in the column that is missing!

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- The researchers split up the donkeys into the 6 different age groups, then sampled 5 donkeys uniformly at random without replacement within each age group.
- This one... also depends on some assumptions about the data.
- If the age groups are not evenly distributed, then this is definitely
   MAR: the missingness will be more likely for some values than others, depending on age.
- If the age groups are evenly distributed, then every value has the same probability of being chosen, so it's MCAR.

## Section 2

# Scatterplots (1)

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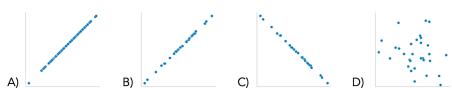
For this next question, assume that the researchers chose the 30 donkeys to reweigh by drawing a simple random sample of 30 underweight donkeys: donkeys with BCS values of 1, 1.5, or 2. The researchers weighed these 30 donkeys one day later and stored the results in WeightAlt.

First: what kind of missingness is this?

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### **Problem**

Which of the following shows the scatter plot of WeightAlt - Weight on the y-axis and 'Weight' on the x-axis? Assume that missing values are not plotted.



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Note that this is the scatter plot of WeightAlt - Weight, not just WeightAlt!

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- Can we say anything from the problem description about the relationship between WeightAlt Weight?

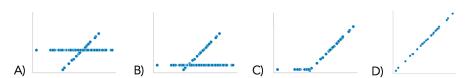
- Note that this is the scatter plot of WeightAlt Weight, not just WeightAlt!
- Can we say anything from the problem description about the relationship between WeightAlt - Weight?
- The answer is, pretty much, no for available values, WeightAlt is just Weight but measured on the next day. So the difference between the two should be pretty much random, and the answer is D.

## Section 3

# Scatterplots (2)

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Suppose we use mean imputation to fill in the missing values in WeightAlt. Select the scatter plot WeightAlt on Weight after imputation.



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- So, what should a scatterplot for a mean-imputed column of data look like?

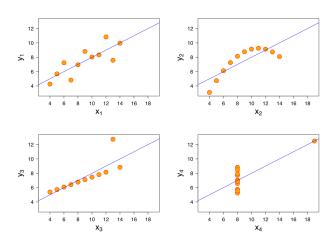
- First: what is mean imputation?
- So, what should a scatterplot for a mean-imputed column of data look like?
- The answer is A: A noisy line of values that were present in the dataset, and then a bunch of values that, no matter the Weight value, now all have the value of the mean.

## Section 4

# **Choosing Test Statistics**

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## **Anscombe's Quartet**



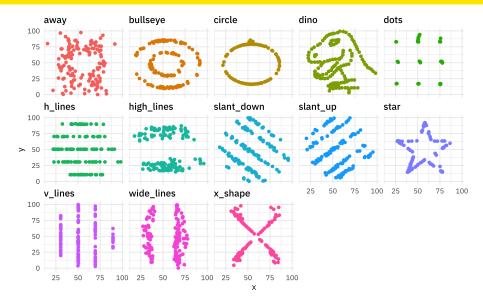
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## **Anscombe's Quartet Stats**

Property	Value	Accuracy
Mean of x	9	exact
Sample variance of $x$ : $s_x^2$	11	exact
Mean of y	7.50	to 2 decimal places
Sample variance of $y$ : $s_y^2$	4.125	±0.003
Correlation between x and y	0.816	to 3 decimal places
Linear regression line	y = 3.00 + 0.500x	to 2 and 3 decimal places, respectively
Coefficient of determination of the linear regression: ${\cal R}^2$	0.67	to 2 decimal places

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## **Datasaurus Dozen**



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## Datasaurus Stats

```
datasaurus dozen %>%
 group_by(dataset) %>%
  summarise(across(c(x, y), list(mean = mean, sd = sd)),
    x_y_cor = cor(x, y)
```

```
## # A tibble: 13 x 6
##
     dataset x_mean x_sd y_mean y_sd x_y_cor
##
  <chr>
              <dbl> <dbl> <dbl> <dbl>
                                       <dbl>
               54.3 16.8 47.8 26.9 -0.0641
##
  1 away
##
   2 bullseye 54.3 16.8 47.8 26.9 -0.0686
               54.3 16.8 47.8 26.9 -0.0683
##
   3 circle
##
   4 dino
               54.3 16.8 47.8 26.9 -0.0645
##
   5 dots
             54.3 16.8 47.8 26.9 -0.0603
             54.3 16.8 47.8 26.9 -0.0617
##
   6 h_lines
  7 high_lines 54.3 16.8 47.8 26.9 -0.0685
##
##
  8 slant_down 54.3 16.8 47.8 26.9 -0.0690
##
             54.3 16.8 47.8 26.9 -0.0686
   9 slant_up
  10 star
              54.3 16.8 47.8 26.9 -0.0630
##
## 11 v_lines
                54.3 16.8 47.8
                                26.9 -0.0694
```

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## Section 5

## FA23 Final Problem 3

#### **Data**

The bus table (left) records bus arrivals over 1 day for all the bus stops within a 2 mile radius of UCSD. The data dictionary (right) describes each column.

	time	line	stop	late
0	12pm	201	Gilman Dr & Mandeville Ln	-1.1
1	1:15pm	30	Gilman Dr & Mandeville Ln	2.8
2	11:02am	101	Gilman Dr & Myers Dr	-0.8
3	8:04am	202	Gilman Dr & Myers Dr	NaN
4	9am	30	Gilman Dr & Myers Dr	-3.0

time	Time of arrival (str). Note that the			
	times are inconsistently entered (e.g.			
	12pm vs. 1:15pm).			

line Bus line (int). There are multiple
 buses per bus line each day.
stop Bus stop (str).

umn are missing.

late The number of minutes the bus arrived after its scheduled time. Negative numbers mean that the bus arrived early (float). Some entries in this col-

## Problem 1

**Question**: Are buses equally likely to be early or late?

How would we generate one sample from the null hypothesis?

- np.random.choice([-1, 1], bus.shape[0])
- np.random.choice(bus['late'], bus.shape[0], replace =
  True)
- Randomly permute the late column

• Is this a (standard) hypothesis test or a permutation test?

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- What is the null hypothesis? Which option generates from the null hypothesis?
- The answer is np.random.choice([-1, 1], bus.shape[0])

# Problem 1 (cont.)

What test statistic should we use?

Note: while the problem says there is only one solution, post-exam two options for the test statistic were given credit. Pick one of the two.

- Number of values below 0
- np.mean
- np.std
- TVD
- K-S statistic

• What information do we want to capture about each sample?

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- Which test statistic captures it?

- What information do we want to capture about each sample?
- Which test statistic captures it?
- The answer is either # values below 0, or np.mean

## **Problem 2**

**Question**: Is the late column MAR dependent on the line column?

How would we generate one sample from the null hypothesis?

- np.random.choice([-1, 1], bus.shape[0])
- np.random.choice(bus['late'], bus.shape[0], replace =
  True)
- Randomly permute the late column

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- Is this a (standard) hypothesis test or a permutation test?
- What is the null hypothesis? Which option generates from the null hypothesis?
- The answer is to randomly permute the late column

### Section 6

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- Good luck!

## Section 7

## **Attendance**

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Once I give you a number, fill out the following Google form: https://forms.gle/wz1r6G2oiwn8cigU8



