

DATA ANALYSIS

Week 7: Sampling and hypothesis testing

logistics: midterm 1

- exam grades will be made available latest by Thursday morning
- review my comments in rubric
- please come see me if you have questions!

mid-semester check-in

- everyone schedules a 15-minute meeting post spring break
[[calendly link](#)]
- fill out **anonymous** mid semester survey [opens on Friday]

7	T: March 4, 2025	W7: Sampling and Hypothesis Testing
7	Th: March 6, 2025	W7 continued...
7	F: March 7, 2025	PS3 revision due
7	F: March 7, 2025	Week 7 Quiz due
8	T: March 11, 2025	Spring Break!
8	Th: March 13, 2025	Spring Break!
9	T: March 18, 2025	Spring Break!
9	Th: March 20, 2025	Spring Break!
10	T: March 25, 2025	W10: Modeling Relationships
10	Th: March 27, 2025	W10 continued...
10	Su: March 30, 2025	Week 10 Quiz due
11	M: March 31, 2025	PS4 due / Opt-out Deadline 2
11	T: April 1, 2025	W11: Special Cases
11	Th: April 3, 2025	W11 continued...
12	M: April 7, 2025	PS5 + PS4 revision due
12	T: April 8, 2025	W12: Loose Ends / Exam 2 review
12	Th: April 10, 2025	Exam (Midterm) 2

today's agenda



sampling



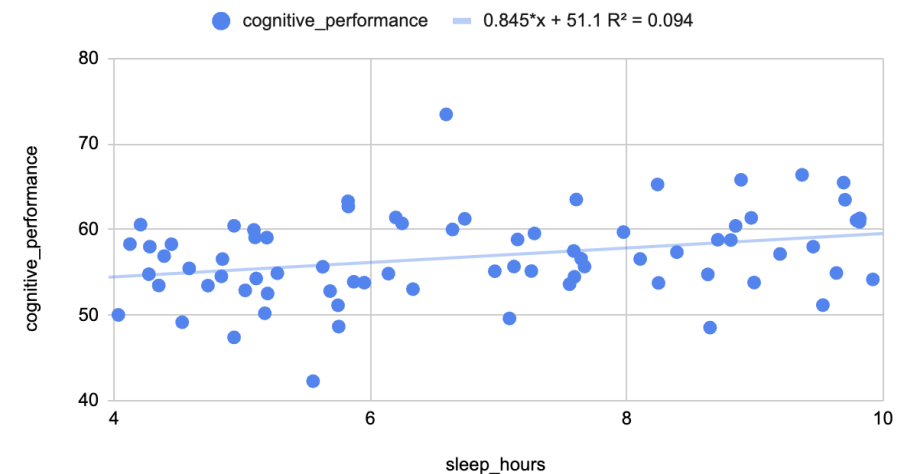
hypothesis testing

an experiment

- **hypothesis**: sleep predicts cognitive performance
- recorded number of hours of sleep via sleep tracker
- cognitive performance via phone game
- model formulation
 - cognitive performance \sim sleep + error
- sample correlation, $r = .31$

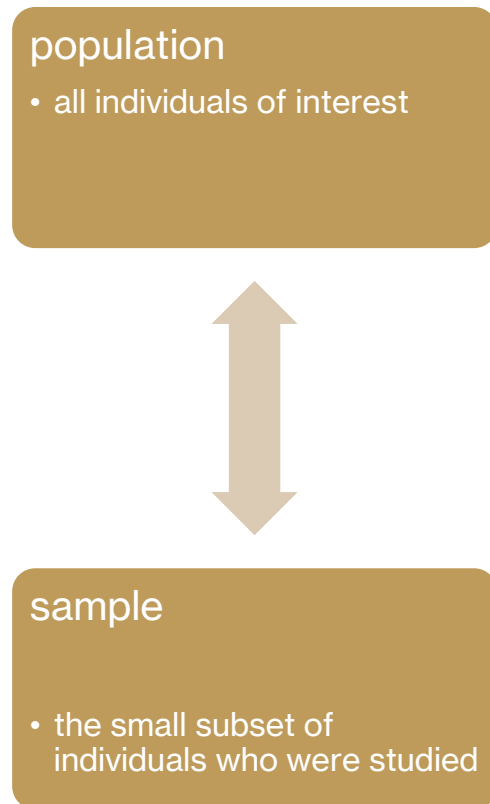
sleep_hours	cognitive_performance
6.24724071	60.7189024
9.70428584	63.4792748
8.39196365	57.3561342
7.59195091	54.4745639
4.93611184	47.3953454
4.93596712	60.434202
4.34850167	53.4598411
9.19705687	57.1451403
7.60669007	63.5052718
8.24843547	53.7573958
4.12350697	58.2974841

cognitive_performance vs. sleep_hours



from samples to populations

- we have a sample statistic (known: r)
- our population parameter (unknown ρ): it could be close or very far from r
- we can **simulate** what the population parameter would look like using our sample statistic
- basic **idea**:
 - samples are small subsets of the population
 - we mimic collecting MANY such samples of the same size and look at the distribution of all possible sample statistics we could obtain



W7 Activity 1

- [Jupyter notebook with data](#)
- confirm correlation, slope, and intercept

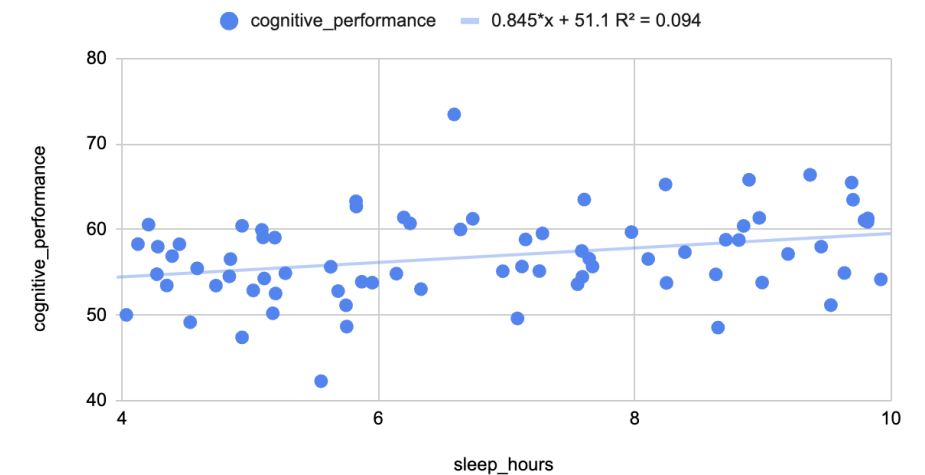
Actual sample slope: 0.84

Actual sample intercept: 51.08

Correlation between sleep_hours and cognitive_performance: 0.31

	sleep_hours	cognitive_performance
0	6.247241	60.718902
1	9.704286	63.479275
2	8.391964	57.356134
3	7.591951	54.474564
4	4.936112	47.395345

cognitive_performance vs. sleep_hours



W7 Activity 1

- what if there was no true relationship between sleep and cognitive performance in the population?
- $\rho = 0$
- how can we mimic this “no relationship” using the sample we have?
- we could keep the **sleep_hours** column the same but **shuffle** the **cognitive_performance** column

original

	sleep_hours	cognitive_performance
0	6.247241	60.718902
1	9.704286	63.479275
2	8.391964	57.356134
3	7.591951	54.474564
4	4.936112	47.395345

shuffled

	sleep_hours	cognitive_performance
0	6.247241	47.395345
1	9.704286	54.845377
2	8.391964	58.297484
3	7.591951	60.718902
4	4.936112	60.422949

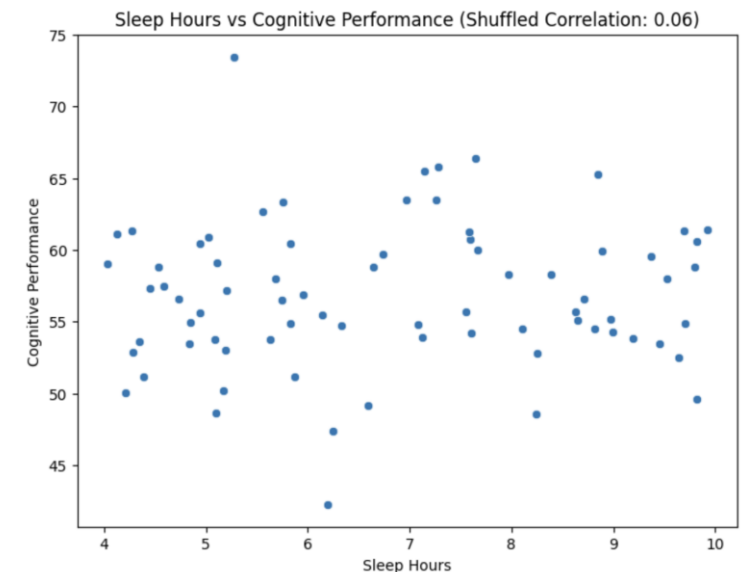
W7 Activity 1

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shuffled



W7 Activity 1

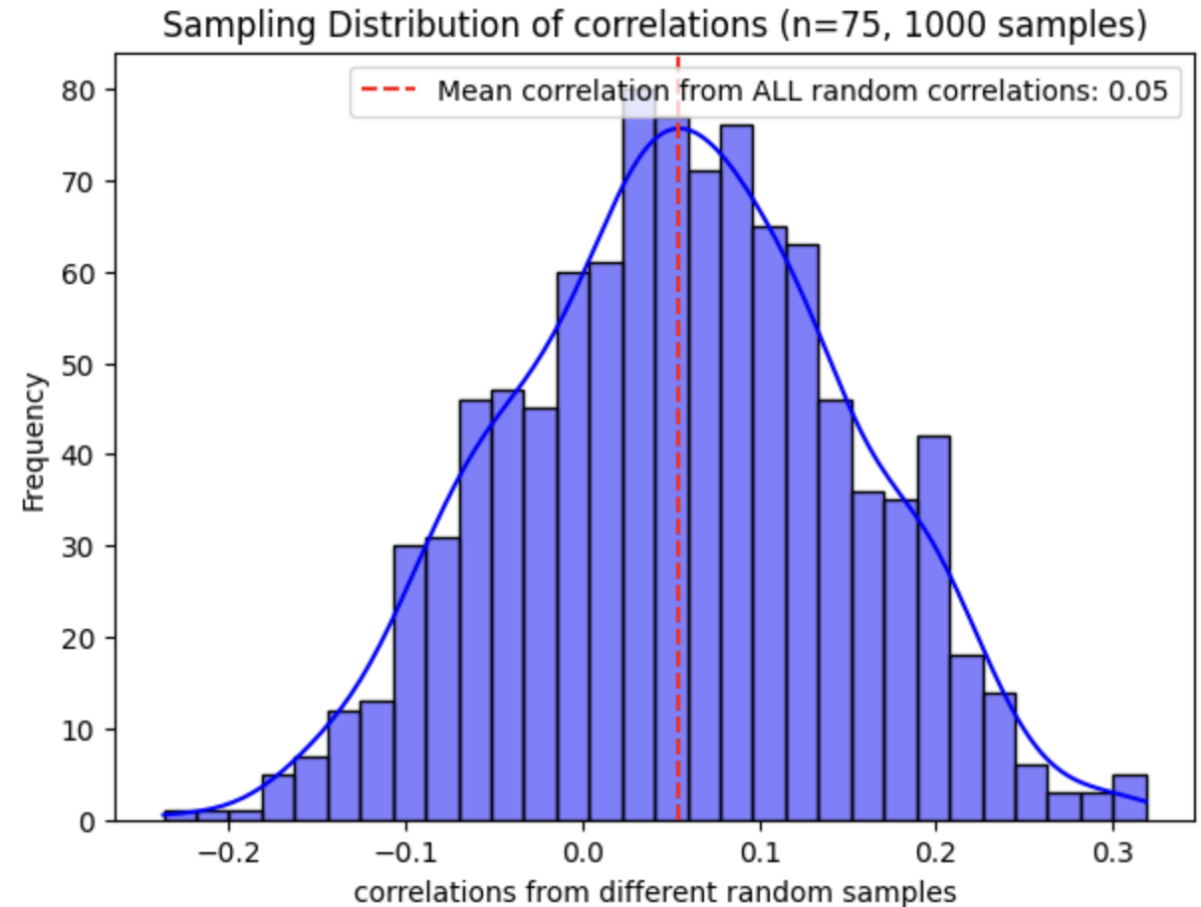
- in this shuffled dataset, what does a random sample look like?
- we repeat this 1000 times, i.e., we take 1000 random samples with replacement
 - **random sample**: each outcome has **an equal chance** of being selected
 - **sampling with replacement** = putting back each observation so that the probability of being selected remains constant on the second draw

```
random sample slope: -0.05  
random sample intercept: 56.53  
random sample correlation: -0.02
```

```
random sample slope: 0.06  
random sample intercept: 55.57  
random sample correlation: 0.03
```

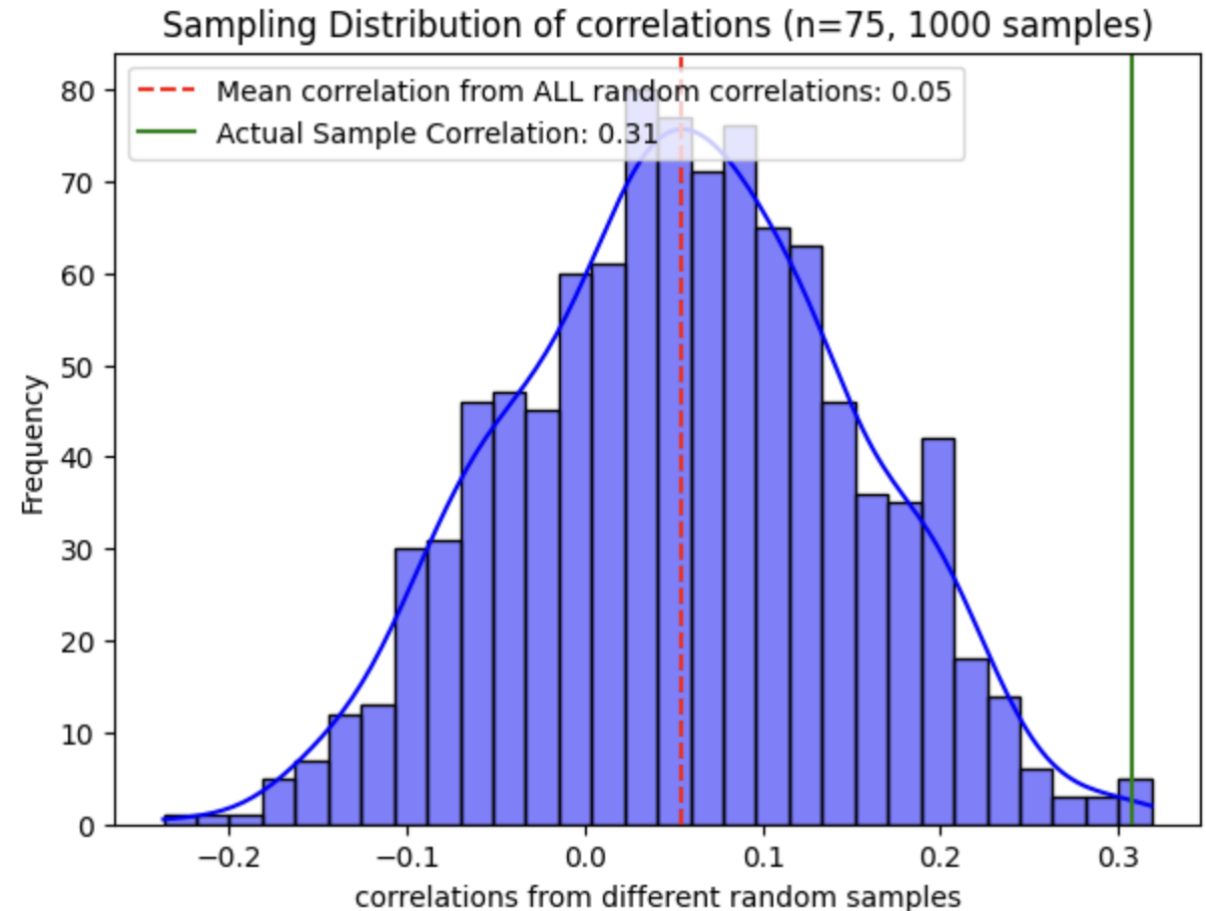
W7 Activity 1

- what does the distribution of correlations look like for MANY such random samples?
- **sampling distribution**: distribution of all possible values of the **sample statistic** obtained from multiple samples of a given size



W7 Activity 1

- what does the distribution of correlations look like for MANY such random samples?
- **sampling distribution**: distribution of all possible values of the **sample statistic** obtained from multiple samples of a given size
- now compare this sampling distribution of random slopes from the shuffled dataset to the **correlation in the actual sample**



W7 Activity 1

- **if there was no relationship between sleep and cognitive performance:**

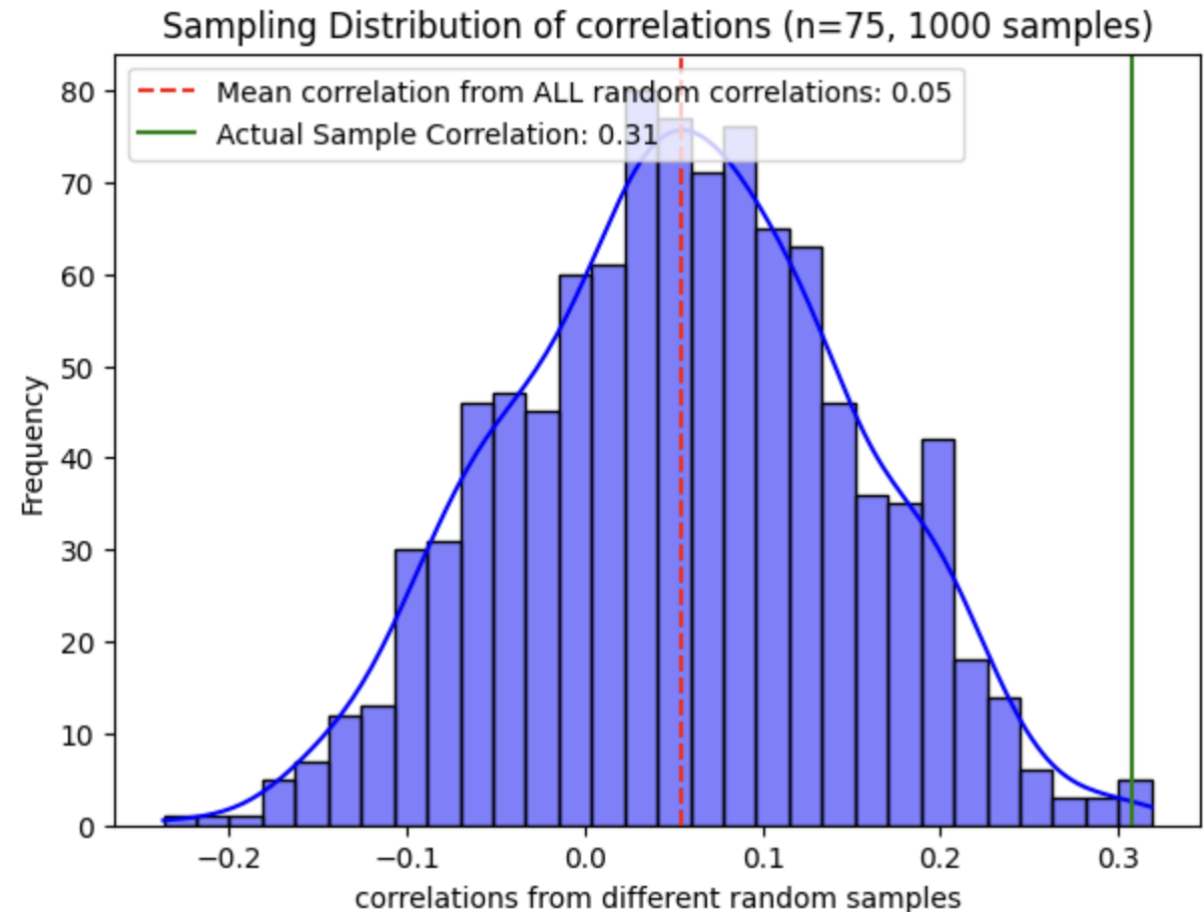
- there is an expected distribution of correlations
- what would be the average distance of a random correlation from the mean correlation?

- standard deviation of sampling distribution of correlations \cong standard error of correlation

- $SE_r = s_r = \sqrt{\frac{1-r^2}{n-2}}$

Mean of ALL random correlations: 0.05

Standard deviation of ALL random correlations: 0.10

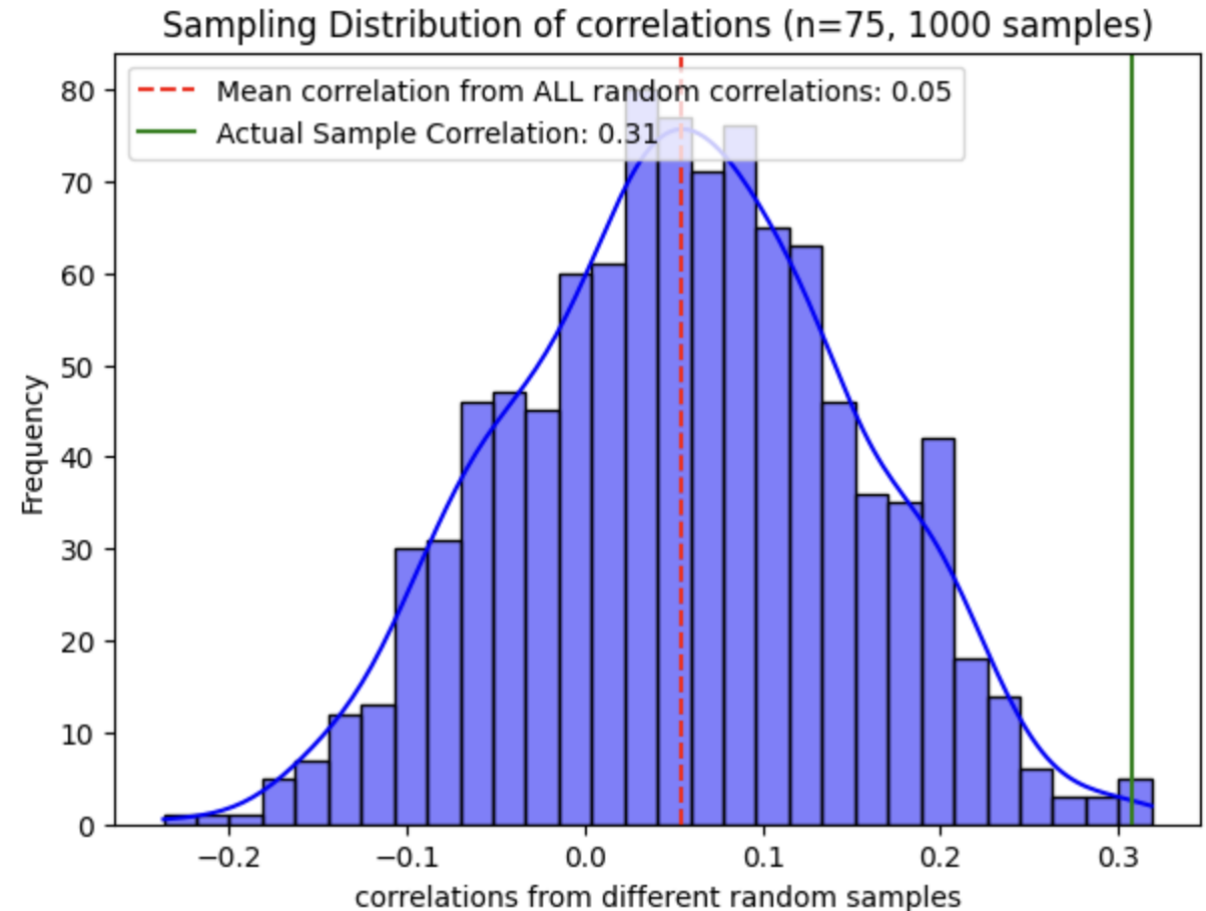


W7 Activity 1

- if there was no relationship between sleep and cognitive performance: is obtaining a sample correlation of 0.31 typical?
- ratio of **observed difference** vs. **expected difference**
- $\frac{\text{observed difference}}{\text{expected difference}} = \frac{r - \rho}{SE_r}$
- like a z-score!

Mean of ALL random correlations: 0.05

Standard deviation of ALL random correlations: 0.10

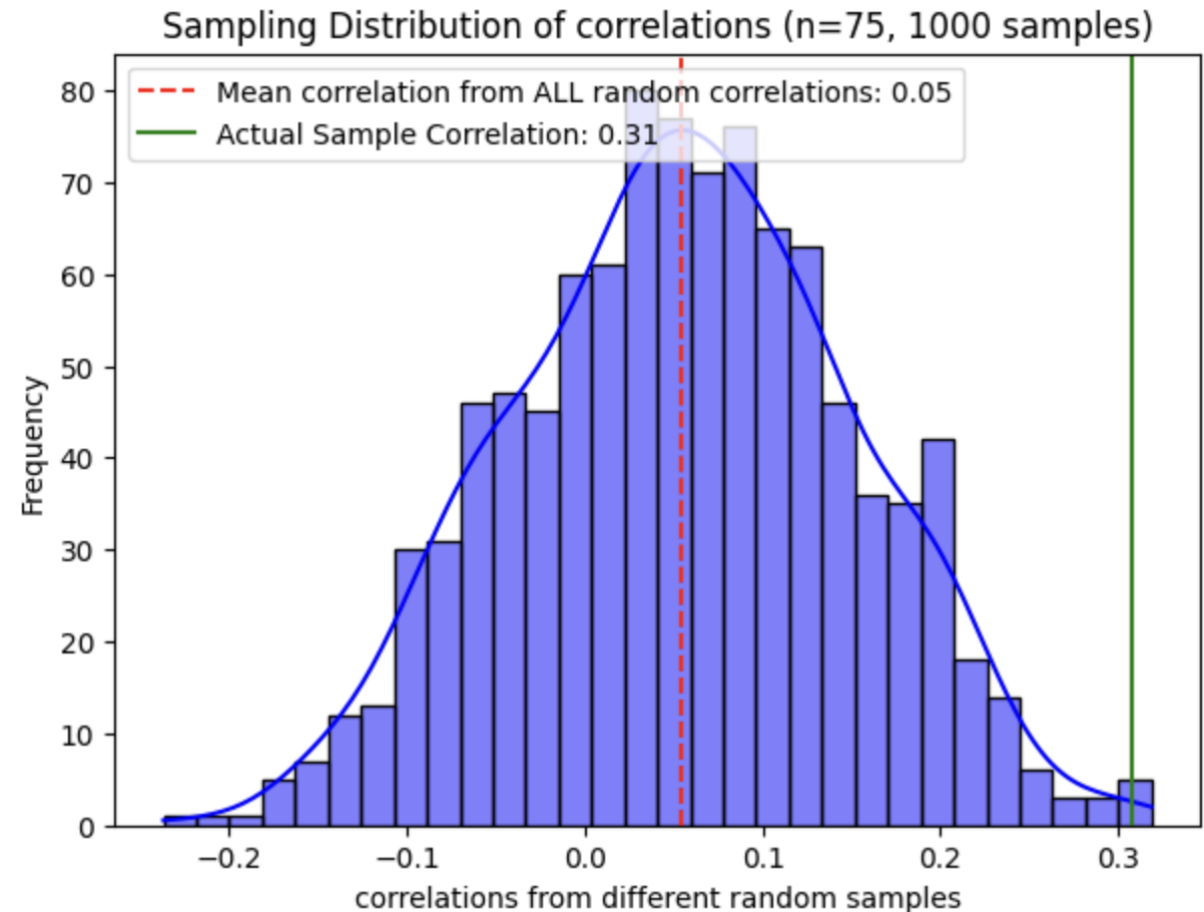


W7 Activity 1

- $\frac{\text{observed difference}}{\text{expected difference}} = \frac{r - \rho}{SE_r}$
- what if we wanted an EXACT probability for our observed correlation?
- we would need to know the exact form of the sampling distribution of the sample statistic we are calculating

Mean of ALL random correlations: 0.05

Standard deviation of ALL random correlations: 0.10



Student's t distribution



BIOMETRIKA.

THE PROBABLE ERROR OF A MEAN.

By STUDENT.

Introduction.

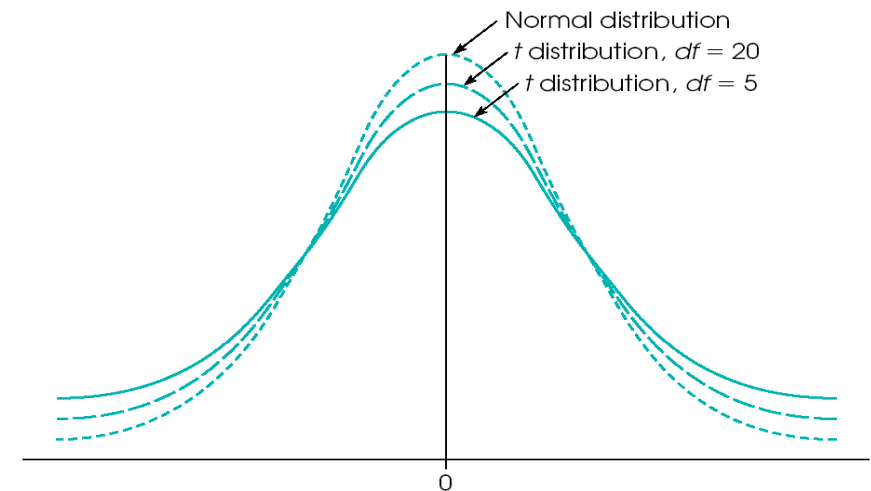
ANY experiment may be regarded as forming an individual of a "population" of experiments which might be performed under the same conditions. A series of experiments is a sample drawn from this population.

Now any series of experiments is only of value in so far as it enables us to form a judgment as to the statistical constants of the population to which the experiments belong. In a great number of cases the question finally turns on the value of a mean, either directly, or as the mean difference between the two quantities.

If the number of experiments be very large, we may have precise information

$$t = \frac{\text{sample statistic} - \text{population parameter}}{\text{standard error}} = \frac{\text{observed}}{\text{expected}}$$

- t distribution approximates the normal distribution
- how good is this approximation?
 - depends on the sample size (n)
 - each t-curve is defined by *degrees of freedom (df)* which depend on the sample size and statistic
 - for large dfs, the t distribution approximates the normal distribution

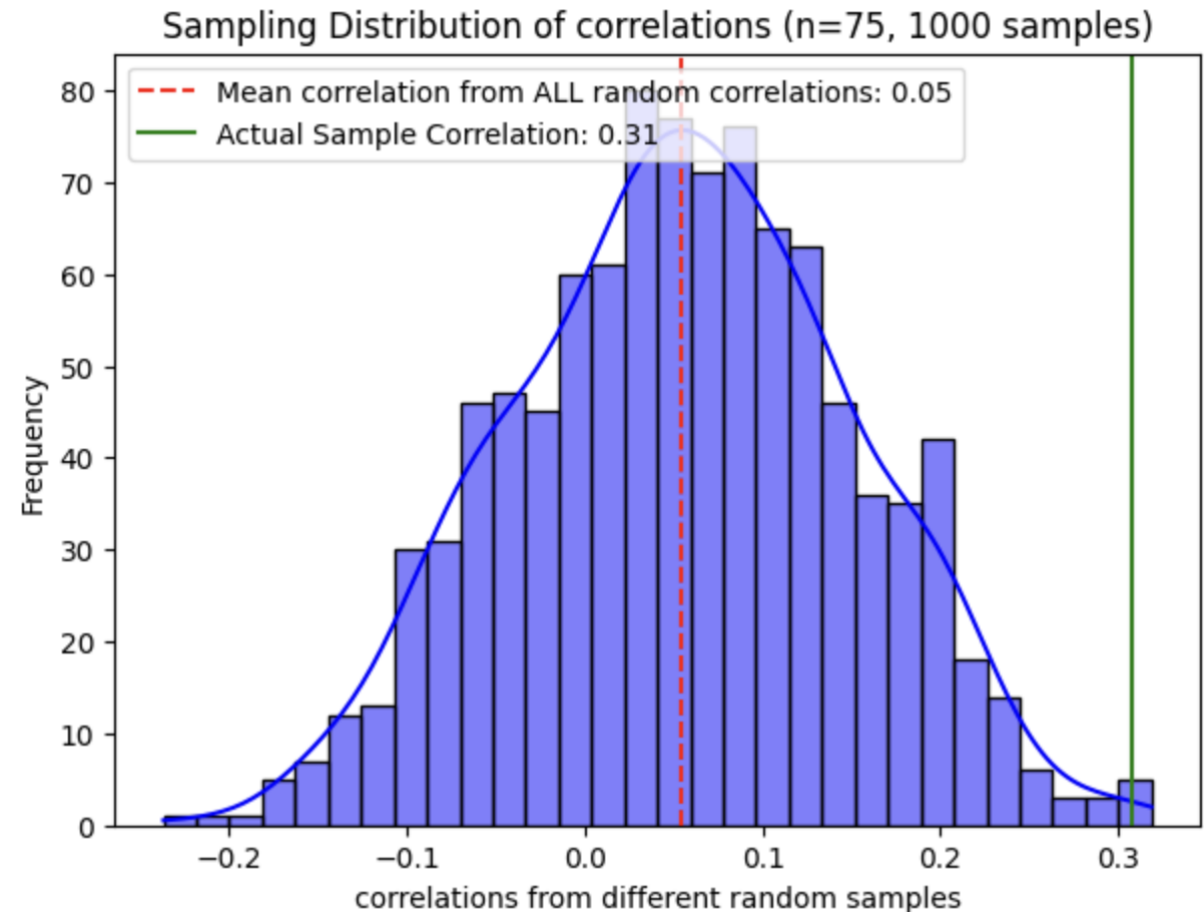


typical sampling distributions

sample statistic	sampling distribution
correlation / slopes	Student's t distribution
ratio of squared errors	F-distribution
means	normal (central limit theorem)

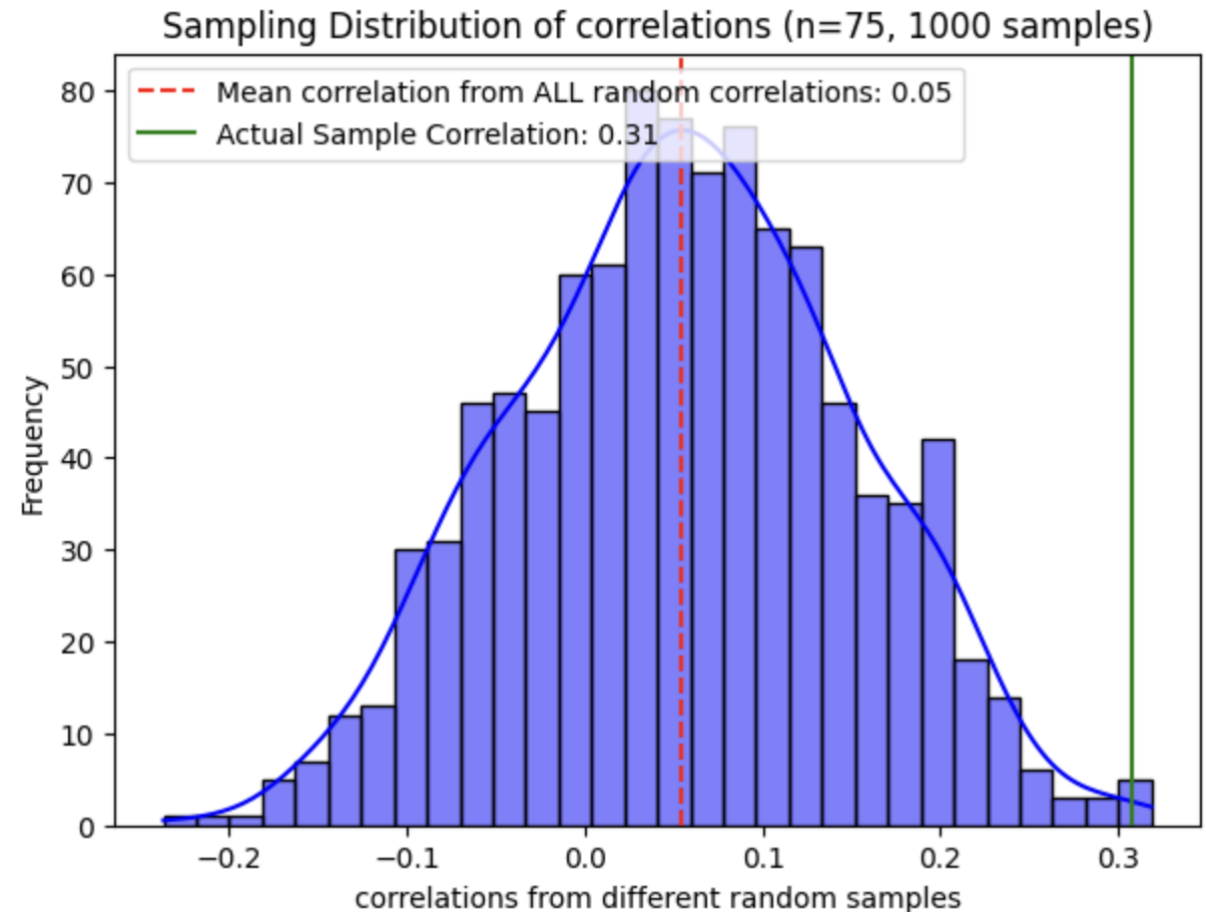
null hypothesis significance testing (NHST)

- start by **assuming that our hypothesis is wrong**
 - **null hypothesis**: there is no meaningful relationship between Y (performance) and X (sleep)
 - H_0 : population parameter $\rho = 0$
 - **alternative hypothesis**: there is a meaningful relationship between Y (performance) and X (sleep)
 - H_1 : population parameter $\rho \neq 0$
- we generate a **sampling distribution** under the null hypothesis using the sample statistic



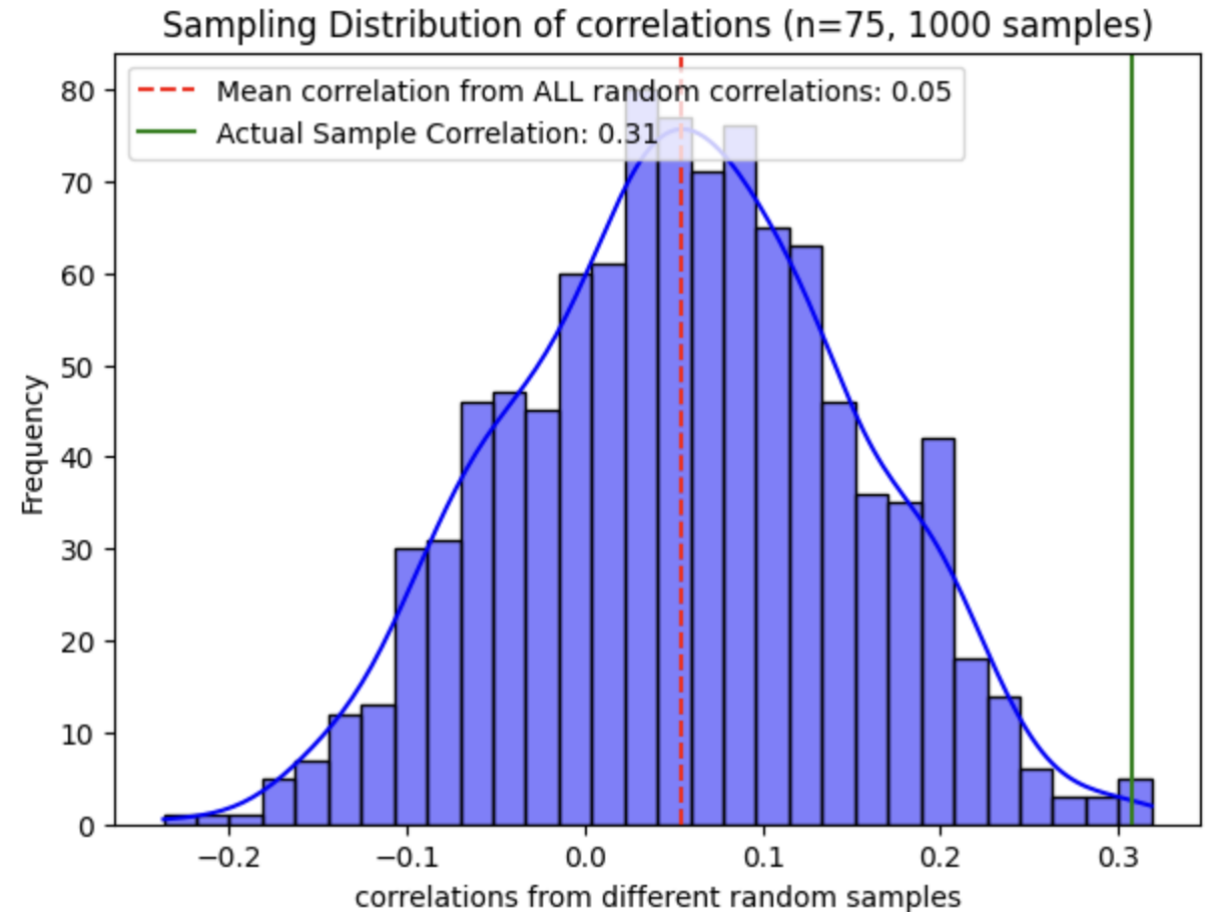
p-value: $P(\text{data} \mid \text{null})$

- once we have a **sampling distribution** under the null hypothesis, we want to know **how likely is the sample statistic you obtained**
- p-value = probability of observing a sample statistic as or more extreme as the one observed if the null hypothesis was true
- if this probability is really low, we can **infer** that the null hypothesis may not be true, and **subsequently infer** that your actual hypothesis may be true!



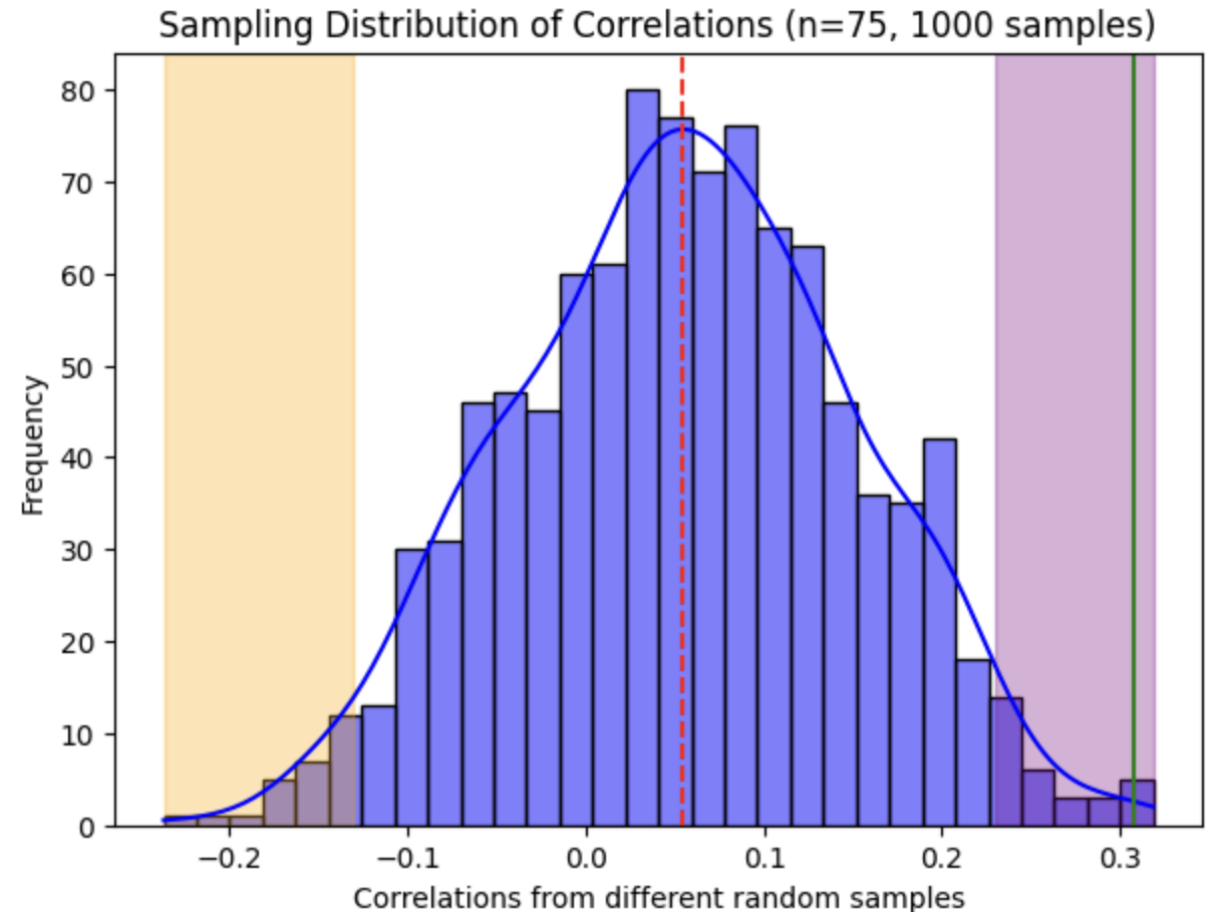
α -level and critical region

- α -level = significance-level criteria
 - typically set to 0.05, i.e., the extreme 5% of the sampling distribution



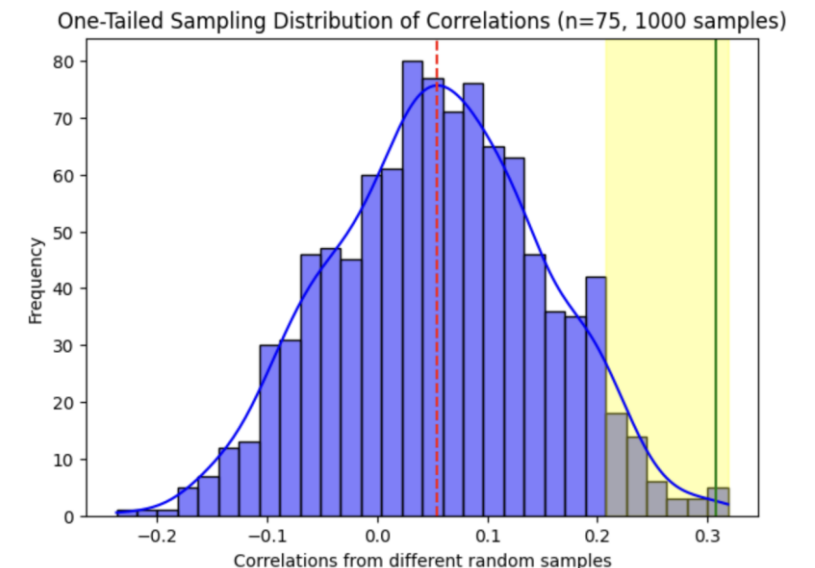
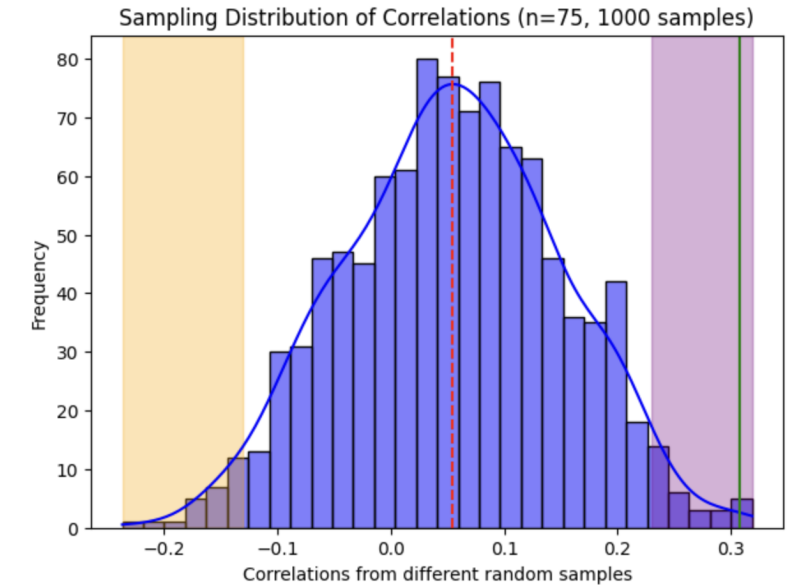
α -level and critical region

- α -level = significance-level criteria
 - typically set to 0.05, i.e., the extreme 5% of the sampling distribution
 - if the observed statistic is within this critical region, we will reject the null hypothesis in favor of the alternative hypothesis
 - commonly referred to as a statistically significant result
- we typically find a critical value based on the underlying sampling distribution (e.g., $t_{critical}$) using a [calculator](#)



one vs. two-tailed tests

- two-tailed tests make **no assumptions about directionality** when discussing the hypotheses
 - $H_0: \rho = 0, H_1: \rho \neq 0$
 - $\alpha = 0.05$ splits the null distribution into two regions (corresponding to $p < .025$ and $p > .025$)
- one-tailed (directional) tests **specify a direction in the hypotheses**, i.e., an increase or decrease in the population parameter
 - $H_0: \rho \leq 0, H_1: \rho > 0$
 - $\alpha = 0.05$ is restricted to only ONE part of the null distribution, leading to a larger area
 - more sensitive but also less conservative





W7 Activity 2

W7 Activity 2 debrief

For a two-tailed hypothesis test evaluating a Pearson correlation, what is stated by the null hypothesis?

- ☐ there is a non-zero correlation for the sample
- ☐ the sample correlation is zero
- ☐ there is a non-zero correlation for the general population
- ☐ the population correlation is zero

Which of the following accurately describes the critical region?

- ☐ outcomes with a very low probability whether or not the null hypothesis is true
- ☐ outcomes with a high probability whether or not the null hypothesis is true
- ☐ outcomes with a high probability if the null hypothesis is true
- ☐ outcomes with a very low probability if the null hypothesis is true

W7 Activity 2 debrief

Which of the following accurately describes a hypothesis test?

- ☐ an inferential technique that uses the data from a sample to draw inferences about a population
- ☐ a descriptive technique that allows researchers to describe a sample
- ☐ a descriptive technique that allows researchers to describe a population
- ☐ an inferential technique that uses information about a population to make predictions about a sample

W7 Activity 2 debrief

If a research report includes the term *significant result*, it means that the null hypothesis was accepted.

- ☐ True
- ☐ False

The null hypothesis is stated in terms of the population, even though the data come from a sample.

- ☐ True
- ☐ False

summary of NHST



step 1:
state the
hypotheses

step 2:
set criteria
for decision

step 3:
collect
data

step 4:
make a
decision!

NHST for correlations

step 1:
state the
hypotheses

$H_0: \rho = 0$
 $H_1: \rho \neq 0$
find the form of the
sampling distribution of
correlations under H_0

step 2:
set criteria
for decision

$\alpha = .05$
find $t_{critical}$ based on
one vs. two tailed
test and **degrees of
freedom = $n - 2$**

step 3:
collect
data

(1) compute SE_r for
sampling distribution of
correlations under H_0
(2) compute $t_{observed} = \frac{r - \rho}{SE_r}$
(3) find p-value for t-score

step 4:
make a
decision!

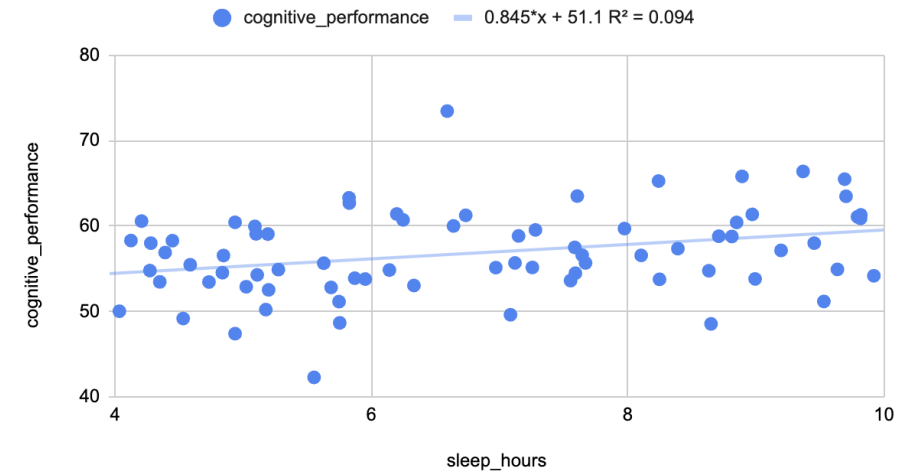
check whether $t_{observed}$
is beyond $t_{critical}$ and
p-value < .05. if so,
reject null hypothesis!

example: NHST for correlation

- hypothesis: sleep predicts cognitive performance

sleep_hours	cognitive_performance
6.24724071	60.7189024
9.70428584	63.4792748
8.39196365	57.3561342
7.59195091	54.4745639
4.93611184	47.3953454
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8.24843547	53.7573958
4.12350697	58.2974841

cognitive_performance vs. sleep_hours



example: NHST for correlation

- **step 1: state the hypotheses**

- $H_0: \rho = 0$
- $H_1: \rho \neq 0$

- **step 2: set criteria for decision**

- $t_{n-2} = t_{73} = t_{critical} = 1.99 \text{ at } \alpha = .05$

- **step 3: collect data**

- compute sample correlation $r = 0.31$
- compute the standard error for correlation

$$SE_r = s_r = \sqrt{\frac{1-r^2}{n-2}} = .11$$

- compute the t-statistic: $t_{observed} = \frac{r-0}{SE_r} = \frac{.31}{.11} = 2.76$
- compute p-value: $p_{observed} = .0073$

- **step 4: decide & report!**

- sleep is significantly correlated with cognitive performance, $r = .31, t(73) = 2.76, p = .007$

t valuez valuechi-square valuef value

r value

Significance Level α : (0 to 0.5)

0.05

Sample Inputs

Degrees of Freedom:

73

Reset

Calculate

Results

t value for Right Tailed Probability:

1.666

t value for Left Tailed Probability:

- 1.666

t value for Two Tailed Probability:

± 1.993

P Value Results

t=2.76 DF=73

The two-tailed P value equals 0.0073

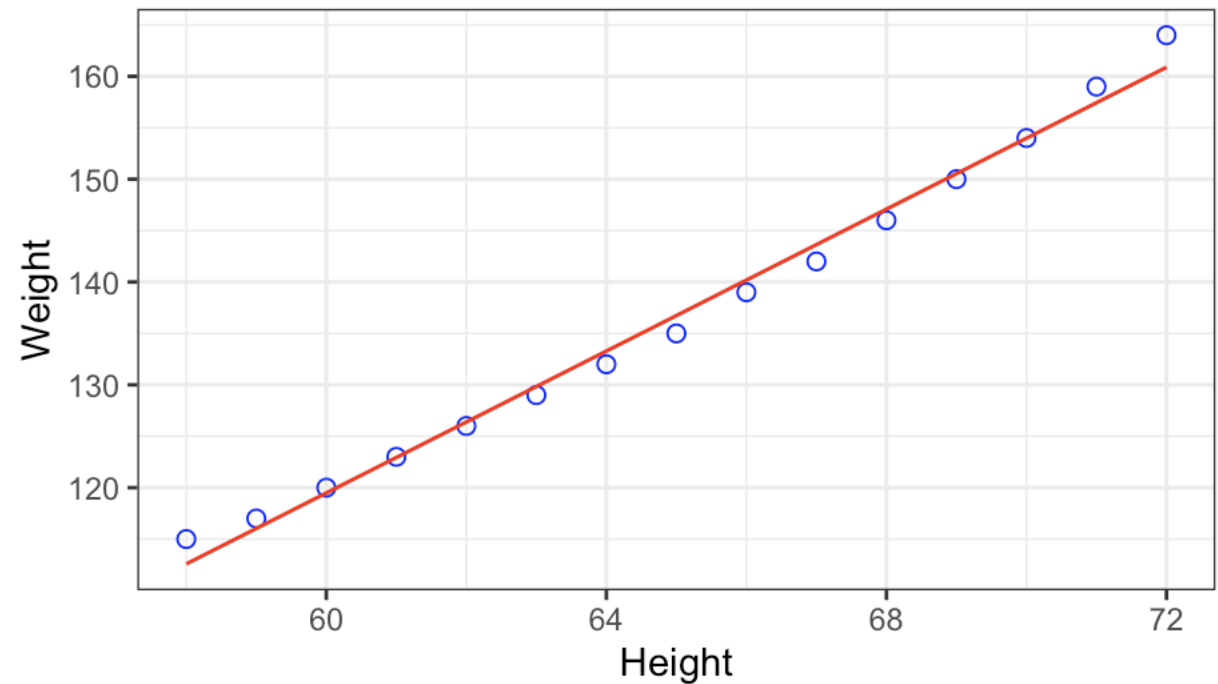
By conventional criteria, this difference is considered to be very statistically significant.

reporting statistical results

- verbal description, sample statistic = $X.XX$, *test statistic* (sample size) = $X.XX$, *p value* = $.XXX$
- statistics are always italicized
- p-values are reported up to third decimal
- everything else reported up to second decimal
- sleep is significantly correlated with cognitive performance,
 $r = .31$, $t(73) = 2.76$, $p = .007$

activity: NHST for correlation

- [women data](#)
- compute correlation and perform a hypothesis test!
- [critical value calculator](#)
- [p-value calculator](#)



activity: NHST for correlation

- **step 1: state the hypotheses**

- $H_0: \rho = 0$
- $H_1: \rho \neq 0$

- **step 2: set criteria for decision**

- $t_{n-2} = t_{13} = t_{critical} = 2.16 \text{ at } \alpha = .05$

- **step 3: collect data**

- compute the correlation $r = 0.995$
- compute the standard error for correlation

$$SE_r = s_r = \sqrt{\frac{1-r^2}{n-2}} = .026$$

- compute the t-statistic: $t_{observed} = \frac{r-0}{SE_r} = \frac{.995}{.026} = 37.855$
- compute p-value: $p_{observed} < .0001$

- **step 4: decide!**

- height significantly correlates with weight,
 $r = .995, t(13) = 37.86, p < .001$

t value z value chi-square value f value r value

Significance Level α : (0 to 0.5) [Sample Inputs](#)

0.05

Degrees of Freedom:

13

[Reset](#) [Calculate](#)

Results

t value for Right Tailed Probability:

1.7709

t value for Left Tailed Probability:

- 1.7709

t value for Two Tailed Probability:

± 2.1604

P from t

t

37.855

DF

13

[Compute P](#)

P Value from Pearson (R) Calculator

This should be self-explanatory, but just in case it's not: your r score goes in the R Score box, the number of pairs in your sample goes in the N box (you must have at least 3 pairs), then you select your significance level and press the button.

If you need to derive a r score from raw data, [you can find a Pearson \(\$r\$ \) calculator here](#).

[How to report Pearson's \$r\$ \(APA\)](#)

R Score: .995

N: 15

Significance Level:

- ☐ 0.01
- ☒ 0.05
- ☐ 0.10

The P-Value is $< .00001$. The result is significant at $p < .05$.

[Calculate](#)

next time

- **more** hypothesis testing

Before Thursday

- Watch: [Hypothesis Testing \(Correlations\)](#).

After Thursday

- Fill out the [Mid-Semester Survey](#) for extra credit!
 - Due after spring break
- See [Apply](#) section.

Here are the to-do's for this week:

- Submit [Week 7 Quiz](#)
- Submit [Problem Set 3 Revision](#)
- Start working on [Problem Set 4](#)
- Submit any lingering questions [here](#)!
- Have a wonderful spring break!!
- Extra credit opportunities:
 - Submit [Mid-Semester Survey](#)
 - Submit [Extra Credit Questions](#)
 - Submit [Optional Meme Submission](#)