

Lab 2 Work

BSTA 512/612

2024-02-08

Directions

Please turn in your `.html` file [on Sakai](#). Please let me know if you greatly prefer to submit a physical copy.

[You can download the `.qmd` file for this lab here.](#)

The rest of this lab's instructions are embedded into the lab activities.

Purpose

The main purpose of this lab is to introduce our dataset, codebook, and variables. We will continue to think about the context of our research question, but our main focus is to become familiar with the data.

Grading

This lab is graded out of 12 points. Nicky will use the following rubric to assign grades.

Rubric

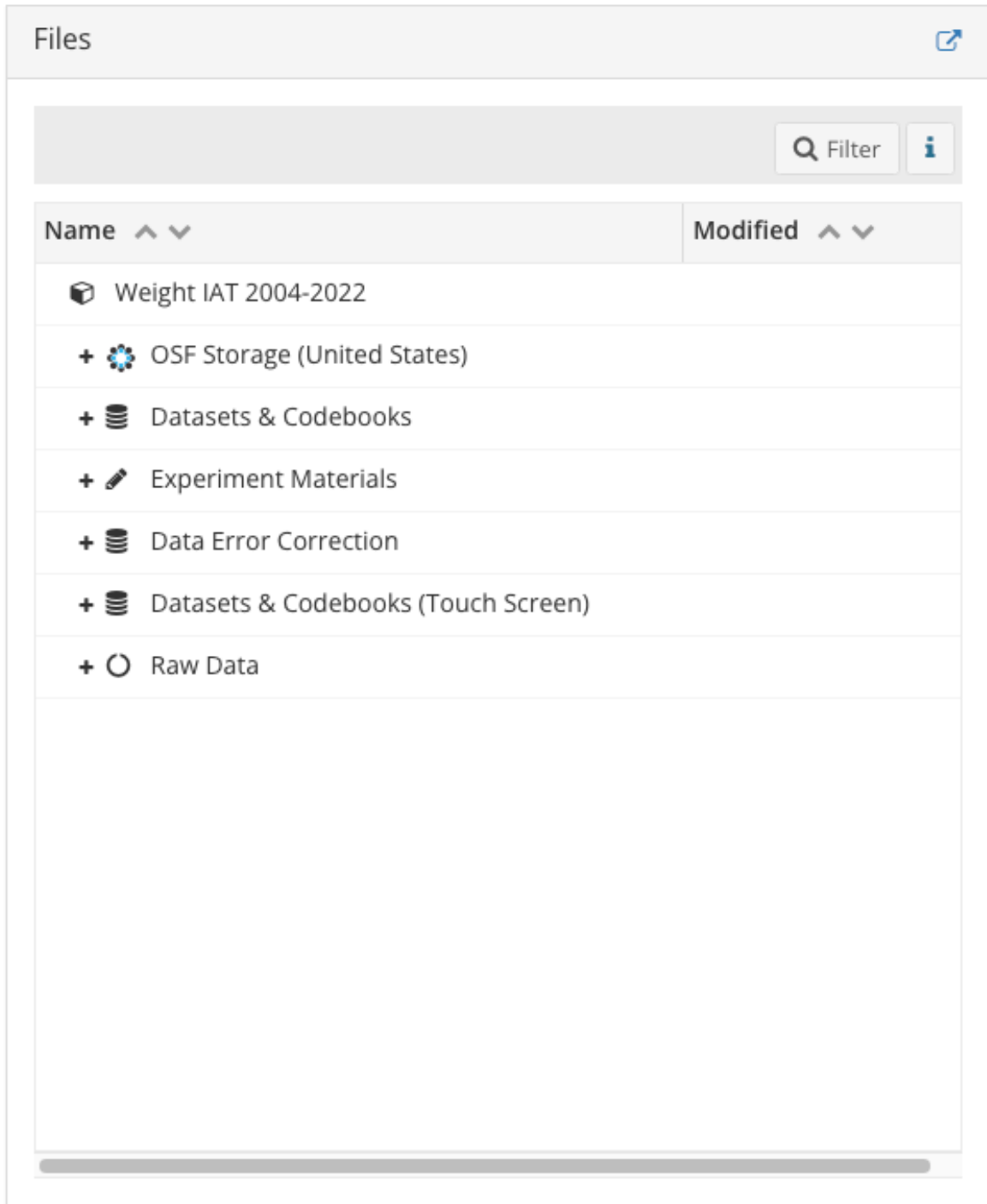
4 points	3 points	2 points	1 point	0 points
Answers demonstrate completion and understanding of the needed activity*. Answers are thoughtful and can be easily integrated into the final report.	Answers demonstrate completion and understanding of the needed activity*. Answers are thoughtful, but lack the clarity needed to easily integrate into the final report.	Answers demonstrate completion and minimal understanding of the needed activity*. Answers are fairly thoughtful, but lack connection to the research.	Answers demonstrate completion of needed activities*, although evidently rushed through. Answers seem rushed and with minimal thought.	It is evident that the needed activities* were not completed. Answers seem rushed and without thought.
Lab submitted on Sakai with .html file. Answers are written in complete sentences with no major grammatical nor spelling errors. With little editing, the answer can be incorporated into the project report.	Lab submitted on Sakai with .html file. Answers are written in complete sentences with grammatical or spelling errors. With editing, the answer can be incorporated into the project report.	Lab submitted on Sakai with .html file. Answers are written in complete sentences with major grammatical or spelling errors. With major editing, the answer can be incorporated into the project report.	Lab submitted on Sakai with .html file. Answers are bulleted or do not use complete sentences.	Lab <i>not</i> submitted on Sakai with .html file.
Code				
Reasoning				

Lab activities


1. Access and download the data




This serves as good practice for accessing data that is online or needs to be downloaded from a collaborator.















Data can be accessed [here](#). Under “Weight IAT 2004-2022” there are several drop down menus:



I opened the first “Datasets & Codebooks,” then selected “OSF Storage (United States).” Once selected, the “Download as zip” option pops up in the top right part of the Files section.


Files 





 Download as zip
  Filter
 















Name ^ v	Modified ^ v
 Weight IAT 2004-2022	
+  OSF Storage (United States)	
-  Datasets & Codebooks	
-  OSF Storage (United States)	
 Weight_IAT.public.2004.zip	2014-04-17 02:48 PM
 Weight_IAT.public.2005.zip	2014-06-09 02:27 PM
 Weight_IAT.public.2006.zip	2014-04-17 02:49 PM
 Weight_IAT.public.2007.zip	2014-04-17 02:50 PM
 Weight_IAT.public.2008.zip	2014-04-17 02:51 PM
 Weight_IAT.public.2009.zip	2014-04-17 02:52 PM
 Weight_IAT.public.2010.revised.zip	2016-10-14 03:36 PM
 Weight_IAT.public.2010.zip	2014-04-17 02:53 PM
 Weight_IAT.public.2011.revised.zip	2016-10-14 03:38 PM
 Weight_IAT.public.2011.revised3320...	2017-03-03 09:46 AM

We will be working with the `Weight_IAT.public.2021.csv` dataset. Please locate the zip file called `Weight IAT.public.2021-CSV.zip` . To download, you need to click the row of

the zip file, but you can't click the name of the zip file. If a link opens, then you clicked the name. If the row is highlighted blue and clickable "Download" and "View" buttons appear on the top right, then you selected it correctly! (See below image for what it should look like.)

Files 

 Download
  View
  Filter
 

Name ^ v	Modified ^ v
 Weight IAT.public.2012.revised-CSV....	2022-06-04 09:22 AM
 Weight IAT.public.2013-CSV.zip	2022-06-04 09:33 AM
 Weight IAT.public.2013.revised-CSV....	2022-06-04 09:30 AM
 Weight IAT.public.2014-CSV.zip	2022-06-04 09:51 AM
 Weight IAT.public.2014.revised-CSV....	2022-06-04 09:40 AM
 Weight IAT.public.2015-CSV.zip	2022-06-04 10:05 AM
 Weight IAT.public.2016-CSV.zip	2022-06-04 03:34 PM
 Weight IAT.public.2017-CSV.zip	2022-06-04 05:28 PM
 Weight IAT.public.2018-CSV.zip	2022-06-04 05:38 PM
 Weight IAT.public.2019-CSV.zip	2022-06-04 05:51 PM
 Weight IAT.public.2019.zip	2021-02-19 02:06 PM
 Weight IAT.public.2020-CSV.zip	2022-06-05 12:14 PM
 Weight IAT.public.2021-CSV.zip	2022-06-05 01:06 PM
-  Amazon S3: weightiat:/ (US Standard)	

Then click the “Download” button to download! Note that the name does not have an underscore between “Weight” and “IAT.” I like to have my datasets named without spaces, so I will

replace the space with an underscore.

For the codebook, perform the same process for the file named: `Weight_IAT_public_2021_codebook.xlsx`

You will need to unzip the actual data.

Move the data to a folder that you can easily access as you work from this document. I like to have a folder named `data` to house my data.

2. Load data and needed packages

First, load the packages that you will need in the remainder of this lab. You can add to this as you need to. At the top of your R code chunk, you can add the following option to repress the messages from the loading packages:

```
{r}
#!/ message: false
```

```
library(tidyverse)
library(gtsummary)
library(here)
```

Using R, load the data (csv file) into this document. Note that this is a csv file that we can load with basic R packages. Name your dataset something that feels intuitive to you and will distinguish it from other datasets that you work with.

```
iat_2021_raw = read.csv(file = here("../TA_files/Project/data/Weight_IAT.public.2021.csv"))
```

Take a glimpse at the data to make sure you loaded it correctly.

```
glimpse(iat_2021_raw)
```

```
Rows: 465,886
Columns: 92
$ session_id      <dbl> 2653543637, 2653543649, 2653543656, 2653543718~
$ session_status  <chr> " ", " ", "C", " ", "C", "C", "C", " ", " ", "~
$ study_name      <chr> "Demo.Weight.0004", "Demo.Weight.0004", "Demo.~
$ date            <chr> "1/1/2021 0:00:49", "1/1/2021 0:02:36", "1/1/2~
$ month           <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1~
$ day             <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1~
$ year            <int> 2021, 2021, 2021, 2021, 2021, 2021, 2021, 2021~
$ hour            <int> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1~
```

\$ weekday	<int>	6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, ~
\$ birthmonth	<int>	NA, 3, 1, NA, 1, 3, 5, NA, NA, 11, 4, 1, 4, 1, ~
\$ birthyear	<int>	NA, 1975, 1979, NA, 1972, 2002, 1943, NA, NA, ~
\$ birthSex	<int>	NA, 2, 2, NA, 2, 1, 2, NA, NA, 1, 2, 2, 1, 2, ~
\$ genderIdentity	<chr>	" ", "[2]", "[2]", " ", "[2]", "[1]", "[2]", "~
\$ num_002	<int>	NA, 1, 1, NA, 1, 1, 2, NA, NA, 1, 1, 1, 4, 4, ~
\$ ethnicityomb	<int>	NA, 2, 2, NA, 2, 3, 2, NA, NA, 1, 2, 2, 2, 2, ~
\$ raceomb_002	<int>	NA, 6, 6, -999, 6, 5, 6, NA, NA, 6, 7, 5, 6, 5-~
\$ raceombmulti	<chr>	" ", " ", " ", " ", " ", " ", " ", " ", " ", " ", "~
\$ D_biep.Thin_Good_all	<dbl>	NA, NA, -0.40210378, 0.51834547, 0.67850537, 0-~
\$ Mn_RT_all_3467	<dbl>	NA, NA, 864.4333, 911.1250, 1088.9833, 655.941~
\$ N_3467	<int>	NA, NA, 120, 120, 120, 120, 120, NA, NA, 120, ~
\$ PCT_error_3467	<dbl>	NA, NA, 7.500000, 5.000000, 2.500000, 7.500000~
\$ Order	<int>	NA, NA, 2, 1, 1, 2, 1, NA, NA, 2, 2, NA, 2, 1, ~
\$ Side_Thin_34	<int>	NA, NA, 1, 2, 2, 1, 2, NA, NA, 1, 1, NA, 1, 2, ~
\$ Side_Good_34	<int>	NA, NA, 2, 2, 2, 2, 2, NA, NA, 2, 2, NA, 2, 2, ~
\$ Stimuli	<int>	NA, NA, 3, 3, 3, 3, 3, NA, NA, 3, 3, NA, 3, 3, ~
\$ pct_300	<dbl>	NA, NA, 0.000000, 0.000000, 0.000000, 8.333333~
\$ pct_400	<dbl>	NA, NA, 0.8333333, 0.0000000, 0.0000000, 14.16~
\$ pct_2K	<dbl>	NA, NA, 0.8333333, 4.1666667, 7.5000000, 3.333~
\$ pct_3K	<dbl>	NA, NA, 0.8333333, 0.0000000, 0.8333333, 1.666~
\$ pct_4K	<dbl>	NA, NA, 0.0000000, 0.0000000, 0.0000000, 0.000~
\$ att7	<int>	NA, NA, 4, 4, 5, 4, 5, NA, NA, 6, 5, NA, 5, NA~
\$ tfat	<int>	NA, NA, 7, NA, 5, 5, 3, NA, NA, 6, 5, NA, 3, N~
\$ tthin	<int>	NA, NA, 7, NA, 5, 5, 5, NA, NA, 6, 7, NA, 7, N~
\$ comptomost_001	<int>	NA, NA, 5, NA, 3, 4, 4, NA, NA, 5, 4, NA, 5, N~
\$ controlyou_001	<int>	NA, NA, 3, NA, 2, 3, 3, NA, NA, 3, 2, NA, 2, N~
\$ controlother_001	<int>	NA, NA, 3, NA, 3, 3, 3, NA, NA, 2, 2, NA, 3, N~
\$ easytolose_001	<int>	NA, NA, 4, NA, 3, 3, 4, NA, NA, 3, 4, NA, 3, N~
\$ iam_001	<int>	NA, NA, 5, NA, 4, 4, 4, NA, NA, 6, 5, NA, 5, N~
\$ identfat_001	<int>	NA, NA, 3, NA, 2, 3, 3, NA, NA, 2, 1, NA, 2, N~
\$ identtthin_001	<int>	NA, NA, 3, NA, 3, 3, 2, NA, NA, 2, 1, NA, 2, N~
\$ important_001	<int>	NA, NA, 4, NA, 3, 3, 2, NA, NA, 4, 4, NA, 3, N~
\$ mostpref_001	<int>	NA, NA, 6, NA, 4, 4, 6, NA, NA, 6, 6, NA, 6, N~
\$ othersay_001	<int>	NA, NA, 4, NA, 4, 4, 4, NA, NA, 4, 4, NA, 5, N~
\$ D_biep.Thin_Good_36	<dbl>	NA, NA, -0.65922173, 0.25492457, 0.85758482, 0-~
\$ D_biep.Thin_Good_47	<dbl>	NA, NA, -0.1449858, 0.7817664, 0.4994259, 0.36~
\$ Mn_RT_all_3	<dbl>	NA, NA, 708.40, 773.70, 978.90, 911.90, 1192.6~
\$ Mn_RT_all_4	<dbl>	NA, NA, 864.9500, 758.3750, 888.8000, 623.8250~
\$ Mn_RT_all_6	<dbl>	NA, NA, 890.550, 860.350, 1510.250, 717.200, 2~
\$ Mn_RT_all_7	<dbl>	NA, NA, 928.875, 1157.975, 1133.575, 529.450, ~
\$ SD_all_3	<dbl>	NA, NA, 262.0541, 303.0142, 472.6722, 805.6804~
\$ SD all 4	<dbl>	NA, NA, 537.2739, 317.2618, 569.8885, 296.7929~


```

$ SD_all_6 <dbl> NA, NA, 265.6665, 376.0075, 644.9575, 438.7512~
$ SD_all_7 <dbl> NA, NA, 320.9680, 588.8174, 362.0522, 213.2356~
$ N_3 <int> NA, NA, 20, 20, 20, 20, 20, NA, NA, 20, 20, NA~
$ N_4 <int> NA, NA, 20, 20, 20, 20, 20, NA, NA, 20, 20, NA~
$ N_5 <int> NA, NA, 28, 28, 28, 28, 28, NA, NA, 28, 28, NA~
$ N_6 <int> NA, NA, 20, 20, 20, 20, 20, NA, NA, 20, 20, NA~
$ N_7 <int> NA, NA, 40, 40, 40, 40, 40, NA, NA, 40, 40, NA~
$ Mn_RT_correct_3 <dbl> NA, NA, 708.4000, 773.7000, 978.9000, 791.0526~
$ Mn_RT_correct_4 <dbl> NA, NA, 708.8235, 745.5385, 809.2051, 601.8421~
$ Mn_RT_correct_6 <dbl> NA, NA, 890.5500, 840.8947, 1487.1053, 661.176~
$ Mn_RT_correct_7 <dbl> NA, NA, 872.3243, 1121.3333, 1114.1282, 501.05~
$ SD_correct_3 <dbl> NA, NA, 262.0541, 303.0142, 472.6722, 613.9011~
$ SD_correct_4 <dbl> NA, NA, 253.1620, 310.7078, 270.6234, 286.9534~
$ SD_correct_6 <dbl> NA, NA, 265.6665, 375.8263, 654.0420, 446.9360~
$ SD_correct_7 <dbl> NA, NA, 258.7414, 597.8491, 344.9726, 185.6936~
$ N_ERROR_3 <int> NA, NA, 0, 0, 0, 1, 0, NA, NA, 1, 0, NA, 2, 0, ~
$ N_ERROR_4 <int> NA, NA, 6, 1, 1, 2, 0, NA, NA, 10, 3, NA, 1, 0~
$ N_ERROR_6 <int> NA, NA, 0, 1, 1, 3, 1, NA, NA, 1, 2, NA, 2, 1, ~
$ N_ERROR_7 <int> NA, NA, 3, 4, 1, 3, 1, NA, NA, 5, 4, NA, 6, 4, ~
$ myweight_002 <int> NA, NA, 24, NA, 20, 32, 19, NA, NA, 34, 18, NA~
$ myheight_002 <int> NA, NA, 36, NA, 34, 38, 33, NA, NA, 33, 30, NA~
$ countrycit_num <int> NA, 1, 85, NA, 1, 1, 1, NA, NA, 1, 105, 1, 1, ~
$ countryres_num <int> NA, 1, 85, NA, 1, 1, 1, NA, NA, 1, 105, 1, 1, ~
$ edu <int> NA, 7, 7, NA, 9, 4, 11, NA, NA, 5, 11, 5, 13, ~
$ edu_14 <int> NA, 7, 7, NA, 9, 4, 11, NA, NA, 5, 11, 5, 13, ~
$ occuSelf <chr> " ", "11-", "43-", " ", "2931", "15-", "2931", ~
$ occuSelfDetail <chr> " ", "1", "43-6000", " ", "29-1000", "15-1000"~
$ politicalid_7 <int> NA, 6, 4, NA, 5, 4, 6, NA, NA, 5, 4, 5, 6, NA, ~
$ STATE <chr> " ", "NC", " ", " ", "NY", "NC", " ", " ", " " ~
$ CountyNo <int> NA, 129, NA, NA, 55, 105, NA, NA, NA, 37, NA, ~
$ MSANo <int> NA, 48900, NA, NA, 40380, 99032, NA, NA, NA, 3~
$ MSAName <chr> " ", "Wilmington, NC MSA", " ", " ", "Rocheste~
$ religion2014 <int> NA, 7, 2, NA, 7, 2, 1, NA, NA, 2, 6, 7, 7, 3, ~
$ religionid <int> NA, 1, 2, NA, 1, 3, 2, NA, NA, 2, 3, 4, 2, 4, ~
$ iatevaluations001 <int> NA, NA, 4, NA, 3, 3, 3, NA, NA, 4, NA, NA, 3, ~
$ iatevaluations002 <int> NA, NA, 3, NA, 1, 3, 2, NA, NA, 2, NA, NA, 2, ~
$ iatevaluations003 <int> NA, NA, 3, NA, 3, 3, 2, NA, NA, 1, NA, NA, 2, ~
$ broughtwebsite <chr> " ", " ", "Mention or link at a non-news Inter~
$ user_id <int> -1, -1, -1, -1, -1, -1, 11555672, -1, -1, -1, ~
$ previous_session_id <dbl> NA, 2653543637, NA, 2653543685, NA, NA, 265354~
$ previous_session_schema <chr> " ", "s", " ", "s", " ", " ", "s", " ", "s", " ~

```

How many rows and columns are in the dataset? Do you think we will need all these variables

for our analysis?

3. Data wrangling

3.1 Restrict your analysis to 1 outcome and 10 possible covariates/predictors

3.1 Manipulating variables that are coded as numeric variables

3.2 Converting categorical variables to continuous variables

3.3 Make a new dataset with only complete cases

Quickly make sure that we are not introducing bias by using complete cases

4. Some exploratory data analysis

4.1 Peek at your outcome

This serves as a check to make sure we are all looking at the correct outcome: IAT score. Please plot a histogram of the IAT scores.

4.1 Univariate exploratory data analysis

4.2 Bivariate exploratory data analysis

4.3 Multivariate exploratory data analysis

5. Revisit your research question

Please restate the research question that you proposed in Lab 1. What are your thoughts on the research question now that we looked at the data?

5. Make a Table 1