My Journey with Sleep: Powered by WHOOP

Code **▼**

Hide

Sys.setenv(RETICULATE_PYTHON = "/home/tliggett/.pyenv/versions/3.9.4/bin/python3.9")
library(reticulate)

Hide

reticulate::repl python()

```
Error in `$.python.builtin.object`(x, name) :
   promise already under evaluation: recursive default argument reference or earlier prob
lems?
Python 3.9.4 (/home/tliggett/.pyenv/versions/3.9.4/bin/python3.9)
Reticulate 1.19 REPL -- A Python interpreter in R.
```

Hide

```
import requests # for getting URL
import json # for parsing json
from datetime import datetime # datetime parsing
import pytz # timezone adjusting
import csv # for making csv files
import os
import pandas as pd
import numpy as np

save_directory = "~/workspace/coursework/data_visualization__COSC322/final_project" # ke
ep trailing slash
```

This is where the API requests from WHOOP are made.

```
def get_user_data_raw(username: str, password: str, start_date='2000-01-01T00:00:00.000
Z', end date='2030-01-01T00:00:00.000Z', url='https://api-7.whoop.com/users/{}/cycles'):
 # GET ACCESS TOKEN
 # Post credentials
 r = requests.post("https://api-7.whoop.com/oauth/token", json={
 "grant type": "password",
  "issueRefresh": False,
 "password": password,
 "username": username })
 # Exit if fail
 if r.status code != 200:
   print("Fail - Credentials rejected.")
   return 1
 print("Success - Credentials accepted")
 # Set userid/token variables
 userid = r.json()['user']['id']
 access token = r.json()['access token']
 # GET DATA
 # Download data
 url = url.format(userid)
 params = {
  'start': start date,
  'end': end date
 }
 headers = {
 'Authorization': 'bearer {}'.format(access token)
 r = requests.get(url, params=params, headers=headers)
 # Check if user/auth are accepted
 if r.status code != 200:
   print("Fail - User ID / auth token rejected.")
   return 1
 print("Success - User ID / auth token accepted")
 # print(json.dumps(r.json()))
 data_raw = r.json()
 return data raw
def get user data df(username: str,
                      password: str,
                      start date='2000-01-01T00:00:00.000Z',
                      end date='2030-01-01T00:00:00.000Z',
                      url='https://api-7.whoop.com/users/{}/cycles'):
```

```
# get raw whoop data
data_raw = get_user_data_raw(username, password, start_date, end_date, url)
# convert json to pandas df

# df = pd.DataFrame.from_dict(data_raw)
df = pd.json_normalize(data_raw)
return df
```

Pull data from both me and Izzy (password are redacted)

Hide

```
tj_data = get_user_data_df("trevor.liggett@gmail.com", "StrongestHalls43")
```

```
Success - Credentials accepted
Success - User ID / auth token accepted
```

Hide

```
izzy_data = get_user_data_df("sommersizzy@gmail.com", "ilovetj")
```

```
Success - Credentials accepted
Success - User ID / auth token accepted
```

This data wrangling is necessary to get both the dates and many of the sleep statistics formatted correctly.

Hide

```
def first_date(row):
    return row['days'][0]

tj_data['date'] = tj_data.apply (lambda row: first_date(row), axis=1)
```

```
def get sleep stat(row, sleep stat: str):
  if row['sleep.sleeps'] == []:
    return np.nan
  return row['sleep.sleeps'][0][sleep_stat]
tj_data['sleep.sws.duration'] = tj_data.apply (lambda row: get_sleep stat(row, 'slowWave
SleepDuration'), axis=1)
tj data['sleep.quality.duration'] = tj data.apply (lambda row: get sleep stat(row, 'qual
ityDuration'), axis=1)
tj_data['sleep.light.duration'] = tj_data.apply (lambda row: get_sleep_stat(row, 'lightS
leepDuration'), axis=1)
tj_data['sleep.rem.duration'] = tj_data.apply (lambda row: get_sleep_stat(row, 'remSleep
Duration'), axis=1)
tj data['sleep.wake.duration'] = tj data.apply (lambda row: get sleep stat(row, 'wakeDur
ation'), axis=1)
tj data['respiratoryRate'] = tj data.apply (lambda row: get sleep stat(row, 'respiratory
Rate'), axis=1)
tj data['sleep.efficiency'] = tj data.apply (lambda row: get sleep stat(row, 'sleepEffic
iency'), axis=1)
tj data['sleep.consistency'] = tj data.apply (lambda row: get sleep stat(row, 'sleepCons
istency'), axis=1)
```

```
quit
library(ggplot2)
library(tidyverse)
```

```
Registered S3 methods overwritten by 'dbplyr':
  method
                  from
  print.tbl lazy
  print.tbl sql
— Attaching packages -
                                                      — tidyverse 1.3.0 —

√ tibble 3.0.6

√ dplyr

                               1.0.5

✓ tidyr

✓ stringr 1.4.0

          1.1.2
                     ✓ forcats 0.5.1

√ readr

          1.4.0
✓ purrr
          0.3.4
— Conflicts -
                                                 — tidyverse conflicts() —
x dplyr::filter() masks stats::filter()
x dplyr::lag()
                  masks stats::lag()
```

```
library(lubridate)
```

```
Attaching package: 'lubridate'

The following objects are masked from 'package:base':

date, intersect, setdiff, union
```

```
library(patchwork) # To display 2 charts together
```

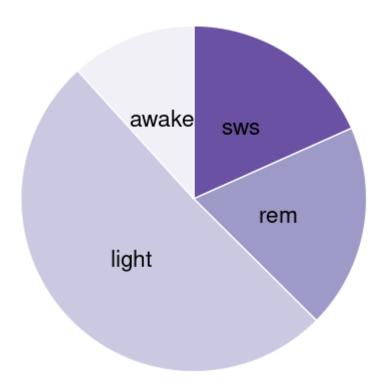
Hide

```
tj_data <- py$tj_data
tj_data$date <- as.Date(tj_data$date, format = "%Y-%m-%d")
```

SLEEP

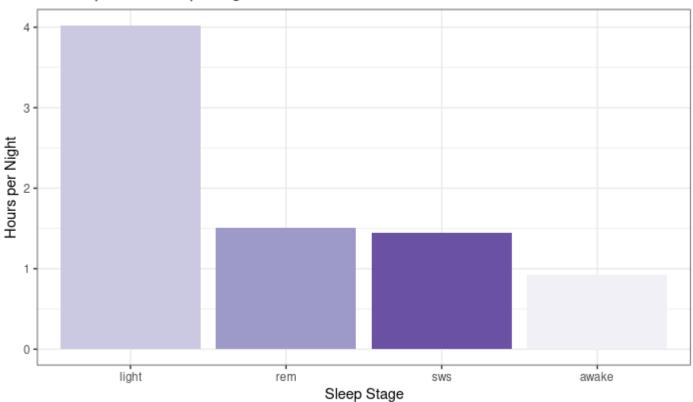
These graphs display summaries of how much time I spent in the different stages of sleep.

```
labels <- c("sws", "rem", "light", "awake")</pre>
sleep_stats <- tj_data %>%
  # filter(!is.na(as.numeric(sleep.sws.duration))) %>%
  summarize(sws = mean(sleep.sws.duration, na.rm = TRUE) / 3600000,
          rem = mean(sleep.rem.duration, na.rm = TRUE) / 3600000,
          light = mean(sleep.light.duration, na.rm = TRUE) / 3600000,
          wake = mean(sleep.wake.duration, na.rm = TRUE) / 3600000)
# Create Data
data <- data.frame(</pre>
  group=c("sws", "rem", "light", "awake"),
  value=c(sleep stats$sws, sleep stats$rem, sleep stats$light, sleep stats$wake)
)
# Compute the position of labels
data <- data %>%
  arrange(desc(group)) %>%
  mutate(prop = value / sum(data$value) *100) %>%
  mutate(ypos = cumsum(prop) - 0.5*prop)
# Basic piechart
ggplot(data, aes(x="", y=prop, fill=group)) +
  geom_bar(stat="identity", width=1, color="white") +
  coord polar("y", start=0) +
  theme void() +
  theme(legend.position="none") +
  geom text(aes(y = ypos, label = group), color = "black", size=6) +
  scale fill brewer(palette="Purples")
```



```
# Bar chart version
ggplot(data = data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(group, -value), y = value, fill=
group)) +
    labs(title = "Time Spent in Sleep Stages", x="Sleep Stage", y="Hours per Night") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="Purples")
```





STRAIN

Here I wrangled the workout data from both me and Izzy in python.

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reticulate::repl_python()

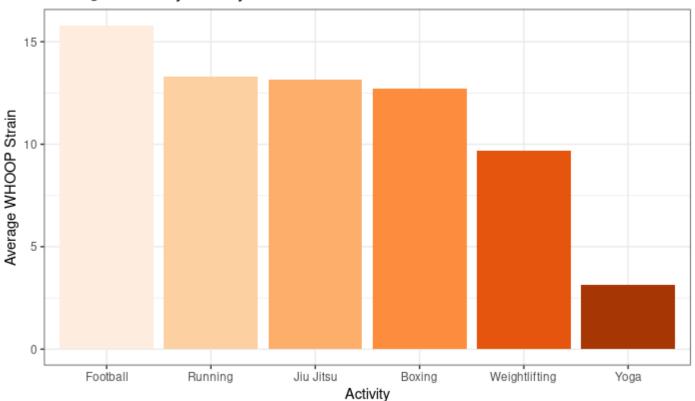
Python 3.9.4 (/home/tliggett/.pyenv/versions/3.9.4/bin/python3.9)
Reticulate 1.19 REPL -- A Python interpreter in R.

```
workouts = pd.DataFrame(columns=['sportId', 'cumulativeWorkoutStrain', 'score'])
for row in tj data['strain.workouts']:
  for workout in row:
    workouts = workouts.append({'sportId': workout['sportId'],
                                 'cumulativeWorkoutStrain': workout['cumulativeWorkoutStr
ain'l,
                                'score': workout['score']
                                }, ignore index=True)
izzy workouts = pd.DataFrame(columns=['sportId', 'cumulativeWorkoutStrain', 'score'])
for row in izzy data['strain.workouts']:
  for workout in row:
    izzy workouts = izzy workouts.append({'sportId': workout['sportId'],
                                 'cumulativeWorkoutStrain': workout['cumulativeWorkoutStr
ain'],
                                'score': workout['score']
                                }, ignore index=True)
```

Now to convert and pipe the data into meaningful summaries, and then graph the findings. Done in R

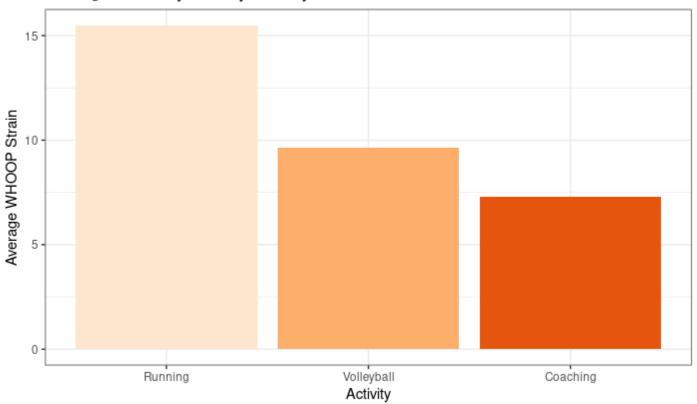
```
workouts <- py$workouts %>%
  filter(sportId %in% c(21, 0, 98, 39, 45, 44)) %>%
  group by(sportId) %>%
  summarize(strain = mean(score, na.rm = TRUE)) %>%
  arrange(desc(strain))
izzy workouts <- py$izzy workouts %>%
  filter(sportId %in% c(0, 36, 87)) %>%
  group by(sportId) %>%
  summarize(strain = mean(score, na.rm = TRUE)) %>%
  arrange(desc(strain))
workouts$sport <- c("Football", "Running", "Jiu Jitsu", "Boxing", "Weightlifting", "Yog</pre>
a")
izzy workouts$sport <- c("Running", "Volleyball", "Coaching")</pre>
ggplot(data = workouts) +
    geom bar(stat = "identity", mapping = aes(x=reorder(sport, -strain), y = strain, fil
l=reorder(sport, -strain))) +
    labs(title = "Average Strain by Activity for T.J.", x="Activity", y="Average WHOOP S
train") +
    theme bw() +
    theme(legend.position = "none") +
    scale fill brewer(palette="0ranges")
```

Average Strain by Activity for T.J.



```
ggplot(data = izzy_workouts) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(sport, -strain), y = strain, fil
l=reorder(sport, -strain))) +
    labs(title = "Average Strain by Activity for Izzy", x="Activity", y="Average WHOOP S
train") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="Oranges")
```

Average Strain by Activity for Izzy



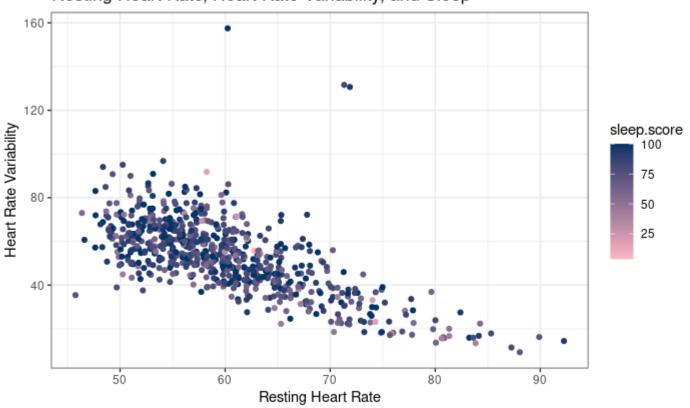
RECOVERY

Charts to show the relationships between recovery statistics.

```
tj_data$recovery.hrv <- tj_data$recovery.heartRateVariabilityRmssd * 1000
tj_data$sleep.hours <- tj_data$sleep.qualityDuration / 3600000

ggplot(data = tj_data, aes(x = recovery.restingHeartRate, y = recovery.hrv, color=sleep.
score)) +
    geom_jitter() +
    scale_color_gradient(low="lightpink", high="#003366") +
    labs(title = "Resting Heart Rate, Heart Rate Variability, and Sleep", x="Resting Heart
Rate", y="Heart Rate Variability") +
    theme_bw()</pre>
```

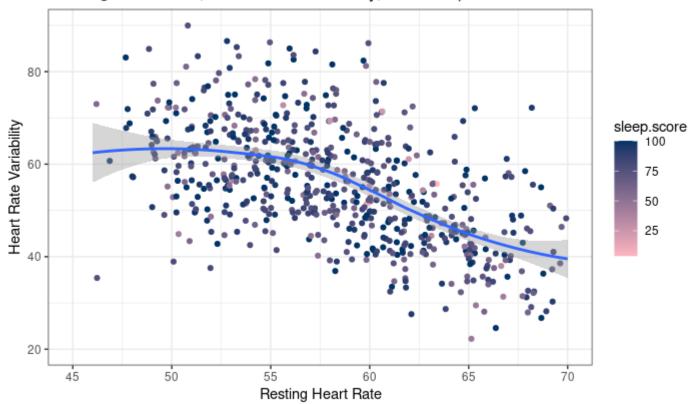
Resting Heart Rate, Heart Rate Variability, and Sleep



```
ggplot(data = tj_data, aes(x = recovery.restingHeartRate, y = recovery.hrv, color=sleep.
score)) +
   geom_jitter() +
   geom_smooth() +
   scale_color_gradient(low="lightpink", high="#003366") +
   labs(title = "Resting Heart Rate, Heart Rate Variability, and Sleep", x="Resting Heart
Rate", y="Heart Rate Variability") +
   theme bw() +
```

xlim(45, 70) + ylim(20, 90)

Resting Heart Rate, Heart Rate Variability, and Sleep



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NA

DAY OF WEEK SUMMARIES

The data wrangling for the dow summaries

```
tj_data$day0fWeek <- weekdays(as.Date(tj_data$date))
tj_data$day0fWeekNum <- wday(as.Date(tj_data$date))

weekdays <- c("Sunday", "Monday", "Tuesday", "Wednesday", "Thursday", "Friday", "Saturda y")

weekday_data <- tj_data %>%
    group_by(day0fWeekNum) %>%
    summarize(recovery = median(recovery.score, na.rm = TRUE), strain = median(strain.scor e, na.rm = TRUE), sleep = mean(sleep.qualityDuration, na.rm = TRUE) / 3600000, sleep_need = mean(sleep.needBreakdown.total, na.rm = TRUE) / 3600000)

weekday_data$dow <- weekdays

weekday_data</pre>
```

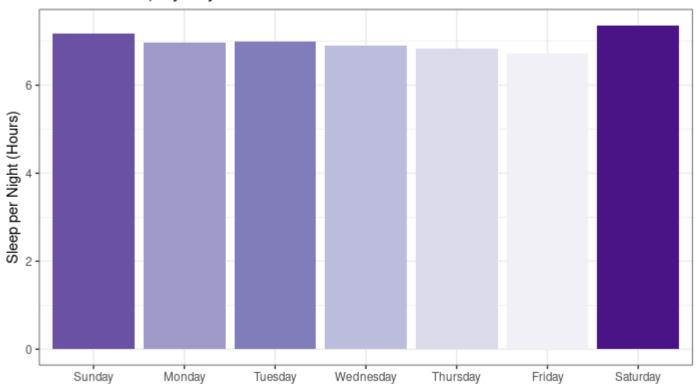
dayOfWeekNum	recovery	strain	sleep	sleep_need dow
<dpl></dpl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl> <chr></chr></dbl>

dayOfWeekNum <dbl></dbl>	recovery <dbl></dbl>	strain <dbl></dbl>	sleep <dbl></dbl>	sleep_need <dbl></dbl>	dow <chr></chr>
1	55	6.152754	7.171962	8.085589	Sunday
2	75	12.958233	6.960574	8.235998	Monday
3	61	13.097235	6.984664	8.546159	Tuesday
4	56	12.552008	6.901612	8.559127	Wednesday
5	58	12.930203	6.826156	8.789615	Thursday
6	57	11.712450	6.726550	8.619979	Friday
7	52	8.908119	7.360054	8.515442	Saturday

Pretty graphs to show trends in dow

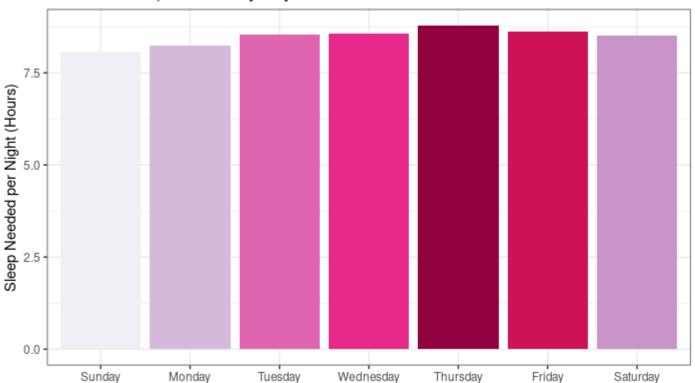
```
ggplot(data = weekday_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(dow,day0fWeekNum), y = sleep, fi
ll=reorder(dow, sleep))) +
    labs(title = "Hours of Sleep by Day of the Week",x = "", y="Sleep per Night (Hours)"
) +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="Purples")
```

Hours of Sleep by Day of the Week



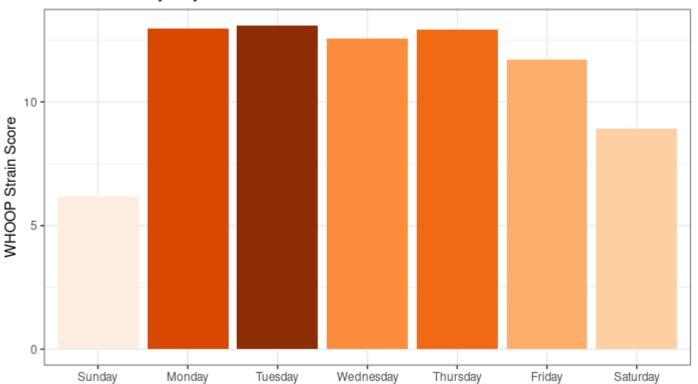
```
ggplot(data = weekday_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(dow,day0fWeekNum), y = sleep_nee
d, fill=reorder(dow,sleep_need))) +
    labs(title = "Hours of Sleep Needed by Day of the Week",x="", y="Sleep Needed per Ni
ght (Hours)") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="PuRd")
```

Hours of Sleep Needed by Day of the Week



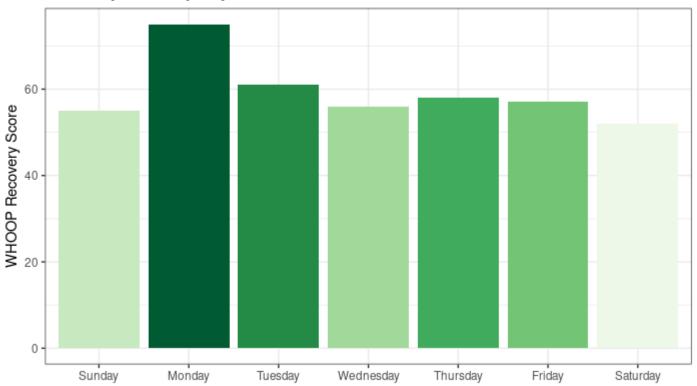
```
ggplot(data = weekday_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(dow,day0fWeekNum), y = strain, f
ill=reorder(dow,strain))) +
    labs(title = "Strain Score by Day of the Week", x="", y="WHOOP Strain Score") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="Oranges")
```

Strain Score by Day of the Week



```
ggplot(data = weekday_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(dow,day0fWeekNum), y = recovery,
fill = reorder(dow,recovery))) +
    labs(title = "Recovery Score by Day of the Week", x="", y="WHOOP Recovery Score") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_brewer(palette="Greens")
```

Recovery Score by Day of the Week

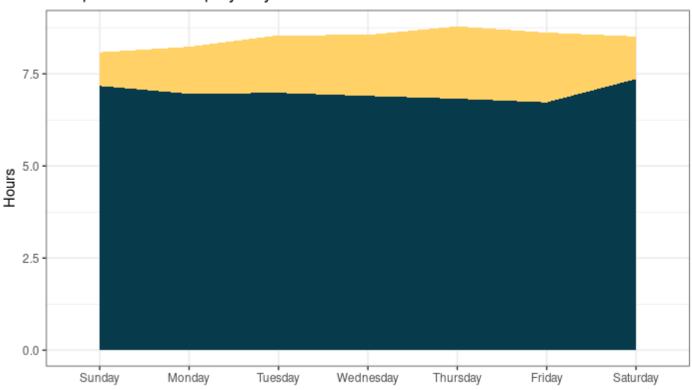


Didn't end up using this but shows the sleep debt incurred in a typical week

```
Hide
```

```
ggplot(data = weekday_data, aes(x=reorder(dow,day0fWeekNum), group = 1)) +
  geom_area(aes(y=sleep_need), fill="#FFD166") +
  geom_area(aes(y=sleep), fill = "#073B4C") +
  labs(title = "Sleep Need vs Sleep by Day of the Week",x="", y="Hours") +
    theme_bw() +
  theme(legend.position = "none")
```

Sleep Need vs Sleep by Day of the Week



MONTHLY SUMMARIES

Summarized data by month. Used in presentation but not the paper.

```
Hide
```

```
# The wrangling of the data
months <- c("Jan", "Feb", "Mar", "Apr", "May", "Jun", "Jul", "Aug", "Sep", "Oct", "Nov",
"Dec")

tj_data$month <- month(as.Date(tj_data$date))

month_data <- tj_data %>%
    filter(date >= "2020-01-01" & date <"2021-01-01") %>%
    group_by(month) %>%
    summarize(recovery = mean(recovery.score, na.rm = TRUE), strain = mean(strain.score, n
a.rm = TRUE), sleep = mean(sleep.qualityDuration, na.rm = TRUE) / 3600000, sleep_need =
    mean(sleep.needBreakdown.total, na.rm = TRUE) / 3600000)

month_data$monthtext <- months

month_data</pre>
```

month	recovery	strain	sleep	sleep_need	monthtext
<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<chr></chr>
1	57.89655	10.311614	6.853161	8.623407	

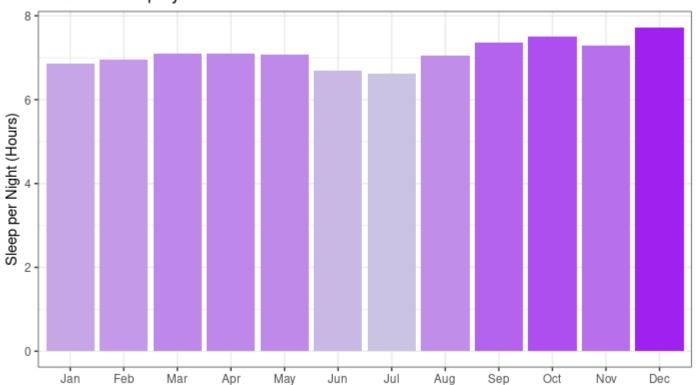
month <dbl></dbl>	recovery <dbl></dbl>	strain <dbl></dbl>	sleep <dbl></dbl>	sleep_need <dbl></dbl>	monthtext <chr></chr>
2	60.79310	11.564513	6.957157	9.060672	Feb
3	64.43478	9.287099	7.089860	8.715013	Mar
4	63.46667	10.555871	7.099221	8.544709	Apr
5	56.50000	9.943054	7.078087	8.599903	May
6	57.16667	12.440945	6.703877	8.430310	Jun
7	57.51613	11.075592	6.613951	8.609364	Jul
8	56.65517	11.536605	7.051709	8.700162	Aug
9	58.90000	10.154535	7.367824	7.831096	Sep
10	59.10000	11.038353	7.497224	8.665091	Oct
1-10 of 12 rows				Prev	ious 1 2 Next

```
# Pretty graphs

ggplot(data = month_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(monthtext, month), y = sleep, fi

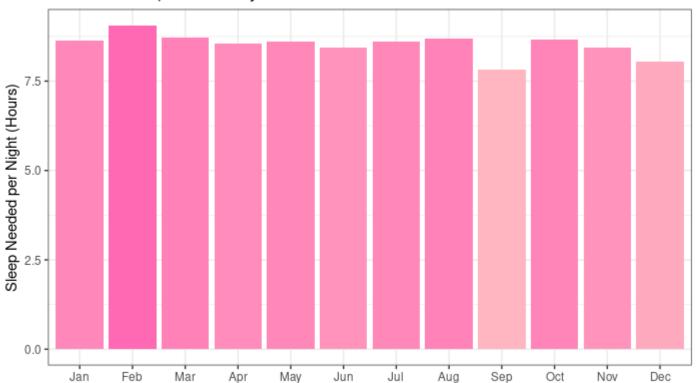
ll=sleep)) +
    labs(title = "Hours of Sleep by Month 2020",x = "", y="Sleep per Night (Hours)") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_continuous(low="#CBC3E3", high="purple")
```

Hours of Sleep by Month 2020



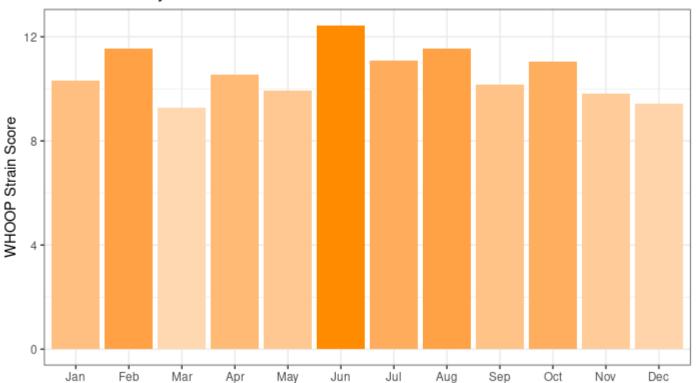
```
ggplot(data = month_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(monthtext, month), y = sleep_nee
d, fill=sleep_need)) +
    labs(title = "Hours of Sleep Needed by Month 2020",x="", y="Sleep Needed per Night
(Hours)") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_continuous(low="lightpink", high="hotpink")
```

Hours of Sleep Needed by Month 2020



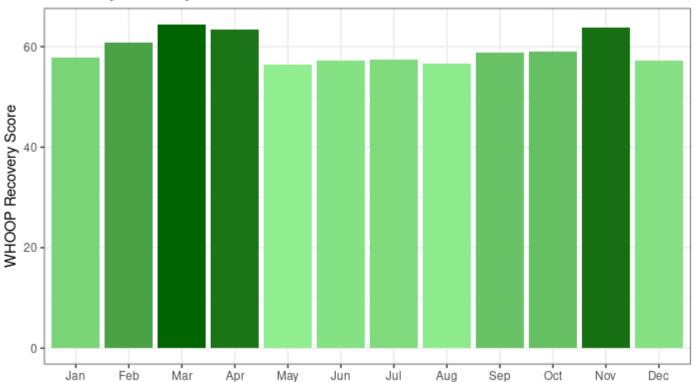
```
ggplot(data = month_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(monthtext, month), y = strain, f
ill=strain)) +
    labs(title = "Strain Score by Month 2020", x="", y="WHOOP Strain Score") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_continuous(low="#fed8b1", high="darkorange")
```

Strain Score by Month 2020



```
ggplot(data = month_data) +
    geom_bar(stat = "identity", mapping = aes(x=reorder(monthtext, month), y = recovery,
fill=recovery)) +
    labs(title = "Recovery Score by Month 2020", x="", y="WH00P Recovery Score") +
    theme_bw() +
    theme(legend.position = "none") +
    scale_fill_continuous(low="lightgreen", high="darkgreen")
```



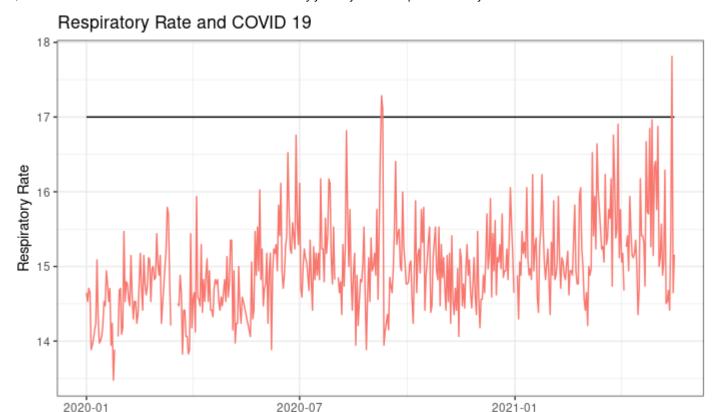


COVID 19 Stuff

For Appendix A. This shows how WHOOP was able to detect my COVID-19 Case as well as my vaccination... Crazy

```
rr_data <- tj_data %>%
  filter(date >= "2020-01-01")

ggplot(data = rr_data, aes(x = date)) +
  geom_line(aes(y=17)) +
  geom_line(aes(y= respiratoryRate, colour='#271033')) +
  labs(title = "Respiratory Rate and COVID 19", x="", y="Respiratory Rate") +
  theme_bw() +
  theme(legend.position = "none")
```



Weekly Average Summaries

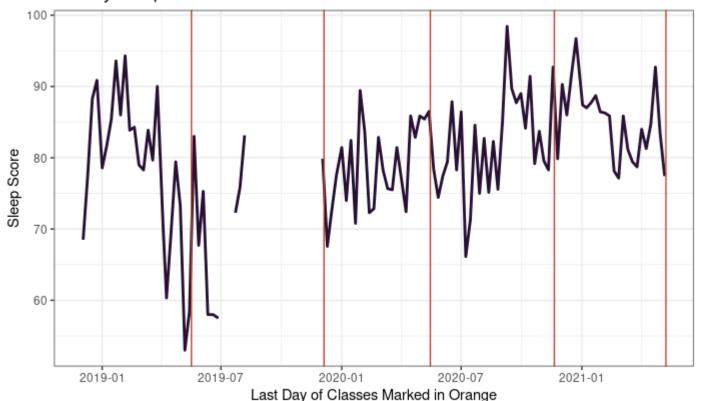
Summarizing some of my major statistics with weekly averages over the years.

`summarise()` has grouped output by 'year'. You can override using the `.groups` argumen t.

```
# These were the last day of classes each of these semesters
finals_dates <- c("2019-05-17","2019-12-05", "2020-05-15","2020-11-20", "2021-05-09")

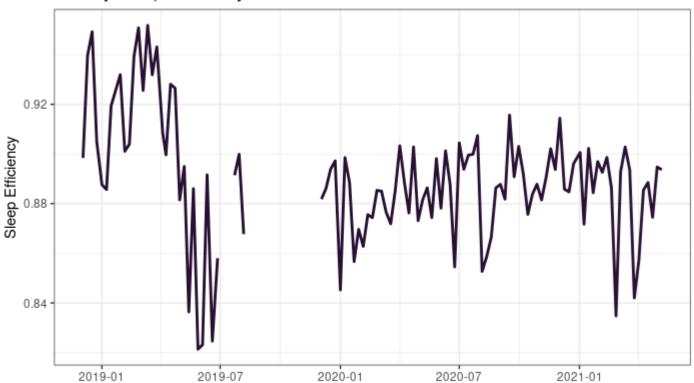
ggplot(weekly_data, aes(x = date, y = sleep)) +
    geom_line(color="#271033", size = 1) +
    geom_vline(xintercept = as.numeric(as.Date(finals_dates, format = "%Y-%m-%d")), color=
"#BB4430") +
    labs(title = "Weekly Sleep Score and Final Exams", x="Last Day of Classes Marked in Or
ange", y="Sleep Score") +
    theme_bw() +
    theme(legend.position = "none")</pre>
```

Weekly Sleep Score and Final Exams



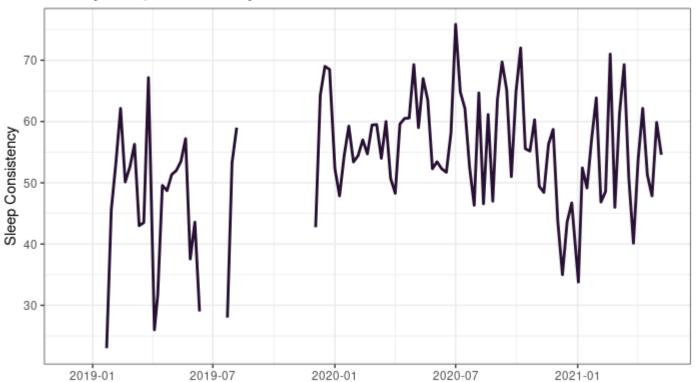
```
ggplot(weekly_data, aes(x = date, y = efficiency)) +
  geom_line(color="#271033", size = 1) +
  labs(title = "Weekly Sleep Efficiency", x="", y="Sleep Efficiency") +
  theme_bw() +
  theme(legend.position = "none")
```

Weekly Sleep Efficiency



```
ggplot(weekly_data, aes(x = date, y = consistency)) +
  geom_line(color="#271033", size = 1) +
  labs(title = "Weekly Sleep Consistency", x="", y="Sleep Consistency") +
  theme_bw() +
  theme(legend.position = "none")
```

Weekly Sleep Consistency



```
ggplot(weekly_data, aes(x = date, y = duration)) +
  geom_line(color="#271033", size = 1) +
  labs(title = "Weekly Sleep Duration", x="", y="Sleep Duration (Hours)") +
  theme_bw() +
  theme(legend.position = "none")
```



