

submission1

Antara Sengupta

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Loading in necessary packages

```
library(readr)
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

library(ggplot2)
library(tibble)
library(tidyr)
library(dplyr)
```

Loading Data

```
# loading in the two separate datasets provided to us (metadata & gene expression data)
series <- read.csv("data/QBS103_GSE157103_series_matrix.csv")
genes <- read.csv("data/QBS103_GSE157103_genes.csv")

# displaying first few rows of each dataframe to get familiarized with the data
head(series)
```

```
##           participant_id geo_accession           status
## 1 COVID_01_39y_male_NonICU   GSM4753021 Public on Aug 29 2020
## 2 COVID_02_63y_male_NonICU   GSM4753022 Public on Aug 29 2020
## 3 COVID_03_33y_male_NonICU   GSM4753023 Public on Aug 29 2020
## 4 COVID_04_49y_male_NonICU   GSM4753024 Public on Aug 29 2020
## 5 COVID_05_49y_male_NonICU   GSM4753025 Public on Aug 29 2020
## 6 COVID_06_:y_male_NonICU    GSM4753026 Public on Aug 29 2020
## X.Sample_submission_date last_update_date type channel_count
## 1           Aug 28 2020      Aug 29 2020  SRA             1
## 2           Aug 28 2020      Aug 29 2020  SRA             1
## 3           Aug 28 2020      Aug 29 2020  SRA             1
## 4           Aug 28 2020      Aug 29 2020  SRA             1
## 5           Aug 28 2020      Aug 29 2020  SRA             1
## 6           Aug 28 2020      Aug 29 2020  SRA             1
```

```

##          source_name_ch1 organism_ch1          disease_status age  sex
## 1 Leukocytes from whole blood Homo sapiens disease state: COVID-19 39  male
## 2 Leukocytes from whole blood Homo sapiens disease state: COVID-19 63  male
## 3 Leukocytes from whole blood Homo sapiens disease state: COVID-19 33  male
## 4 Leukocytes from whole blood Homo sapiens disease state: COVID-19 49  male
## 5 Leukocytes from whole blood Homo sapiens disease state: COVID-19 49  male
## 6 Leukocytes from whole blood Homo sapiens disease state: COVID-19   :  male
##      icu_status apacheii charlson_score mechanical_ventilation
## 1          no         15              0              yes
## 2          no  unknown              2              no
## 3          no  unknown              2              no
## 4          no  unknown              1              no
## 5          no         19              1              yes
## 6          no  unknown              1              no
## ventilator.free_days hospital.free_days_post_45_day_followup ferritin.ng.ml.
## 1              0              0              946
## 2             28             39             1060
## 3             28             18             1335
## 4             28             39             583
## 5             23             27             800
## 6             28             36             563
## crp.mg.l. ddimer.mg.l_feu. procalcitonin.ng.ml.. lactate.mmol.l. fibrinogen
## 1      73.1              1.3              36              0.9             513
## 2   unknown              1.03             0.37          unknown          unknown
## 3      53.2              1.48             0.07          unknown             513
## 4     251.1              1.32             0.98             0.87             949
## 5     355.8              0.69             4.92             1.48             929
## 6     129.1            unknown             0.67             0.86             769
##      sofa
## 1         8
## 2   unknown
## 3   unknown
## 4   unknown
## 5         7
## 6   unknown

```

```
head(genes)
```

```

##          X COVID_01_39y_male_NonICU COVID_02_63y_male_NonICU
## 1      A1BG              0.49              0.29
## 2      A1CF              0.00              0.00
## 3       A2M              0.21              0.14
## 4     A2ML1              0.04              0.00
## 5   A3GALT2              0.07              0.00
## 6    A4GALT              0.00              0.00
## COVID_03_33y_male_NonICU COVID_04_49y_male_NonICU COVID_05_49y_male_NonICU
## 1              0.26              0.45              0.17
## 2              0.00              0.01              0.00
## 3              0.03              0.09              0.00
## 4              0.02              0.07              0.05
## 5              0.00              0.00              0.07
## 6              0.00              0.00              0.00
## COVID_06_.y_male_NonICU COVID_07_38y_female_NonICU COVID_08_78y_male_ICU
## 1              0.21              0.49              0.12
## 2              0.00              0.01              0.00

```

## 3	0.08	0.23	0.08
## 4	0.04	0.03	0.01
## 5	0.00	0.07	0.00
## 6	0.00	0.00	0.00
## COVID_09_64y_female_ICU COVID_10_62y_male_ICU COVID_11_52y_female_NonICU			
## 1	0.51	0.10	0.38
## 2	0.01	0.00	0.02
## 3	0.88	0.13	0.47
## 4	0.02	0.01	0.03
## 5	0.79	0.15	0.08
## 6	0.00	0.00	0.00
## COVID_12_50y_male_ICU COVID_13_37y_male_NonICU COVID_14_55y_male_ICU			
## 1	0.45	0.18	0.23
## 2	0.00	0.00	0.00
## 3	0.16	0.07	0.22
## 4	0.00	0.01	0.04
## 5	1.75	0.00	0.93
## 6	0.00	0.00	0.00
## COVID_15_68y_male_ICU COVID_16_48y_male_NonICU COVID_17_54y_male_NonICU			
## 1	0.42	0.41	0.63
## 2	0.00	0.01	0.02
## 3	0.07	0.58	0.15
## 4	0.00	0.00	0.02
## 5	0.15	0.19	0.00
## 6	0.03	0.00	0.00
## COVID_18_70y_female_NonICU COVID_19_51y_male_NonICU COVID_20_62y_male_ICU			
## 1	0.47	0.33	0.32
## 2	0.00	0.02	0.00
## 3	0.30	0.11	0.07
## 4	0.02	0.02	0.00
## 5	0.06	0.00	0.22
## 6	0.03	0.00	0.00
## COVID_21_66y_male_ICU COVID_22_43y_male_ICU COVID_23_76y_male_ICU			
## 1	0.18	0.09	0.18
## 2	0.00	0.00	0.01
## 3	0.00	0.06	0.03
## 4	0.00	0.00	0.00
## 5	0.37	0.06	0.07
## 6	0.03	0.00	0.03
## COVID_24_55y_male_ICU COVID_25_55y_male_ICU COVID_26_41y_female_ICU			
## 1	0.22	0.29	0.42
## 2	0.01	0.00	0.00
## 3	0.11	0.09	0.18
## 4	0.02	0.03	0.00
## 5	0.15	0.00	0.87
## 6	0.00	0.00	0.00
## COVID_27_71y_female_ICU COVID_28_63y_male_ICU COVID_29_63y_female_ICU			
## 1	0.16	0.18	0.35
## 2	0.01	0.00	0.00
## 3	0.23	0.18	0.03
## 4	0.01	0.05	0.03
## 5	0.18	0.45	0.15
## 6	0.00	0.00	0.03
## COVID_30_54y_male_ICU COVID_31_50y_male_ICU COVID_32_72y_male_ICU			

## 1	0.23	0.15	0.34
## 2	0.00	0.00	0.01
## 3	0.11	0.47	0.04
## 4	0.01	0.00	0.00
## 5	0.00	0.00	0.29
## 6	0.00	0.03	0.00
## COVID_33_81y_male_NonICU	COVID_34_64y_female_NonICU		
## 1	0.35	0.36	
## 2	0.00	0.00	
## 3	0.30	0.11	
## 4	0.06	0.00	
## 5	0.26	0.12	
## 6	0.00	0.00	
## COVID_35_58y_female_NonICU	COVID_36_68y_male_NonICU	COVID_37_87y_male_NonICU	
## 1	0.26	0.18	0.20
## 2	0.00	0.01	0.00
## 3	0.51	0.09	0.09
## 4	0.02	0.00	0.07
## 5	0.16	0.08	0.31
## 6	0.00	0.00	0.00
## COVID_38_68y_male_ICU	COVID_39_80y_female_ICU	COVID_40_66y_male_ICU	
## 1	0.29	0.19	0.22
## 2	0.00	0.00	0.00
## 3	0.10	0.27	0.17
## 4	0.02	0.00	0.00
## 5	0.35	0.00	0.08
## 6	0.00	0.07	0.00
## COVID_41_74y_male_ICU	COVID_42_21y_female_ICU	COVID_43_83y_female_ICU	
## 1	0.19	0.24	0.29
## 2	0.00	0.01	0.00
## 3	0.14	0.33	0.00
## 4	0.00	0.01	0.00
## 5	0.19	0.39	0.11
## 6	0.00	0.00	0.00
## COVID_44_46y_male_ICU	COVID_45_62y_female_ICU	COVID_46_62y_male_ICU	
## 1	0.22	0.14	0.53
## 2	0.00	0.00	0.01
## 3	0.14	0.15	0.10
## 4	0.00	0.03	0.00
## 5	0.00	0.19	0.06
## 6	0.04	0.00	0.00
## COVID_47_78y_male_ICU	COVID_48_72y_female_ICU	COVID_49_73y_male_ICU	
## 1	0.08	0.19	0.48
## 2	0.01	0.00	0.00
## 3	0.04	0.06	0.09
## 4	0.03	0.01	0.03
## 5	0.60	0.23	0.00
## 6	0.00	0.06	0.00
## COVID_50_37y_male_ICU	COVID_51_58y_female_NonICU	COVID_52_71y_male_NonICU	
## 1	0.08	0.21	0.25
## 2	0.00	0.00	0.01
## 3	0.01	0.13	0.00
## 4	0.00	0.00	0.03
## 5	0.00	0.00	0.00

## 6	0.72	0.00	0.00
## COVID_53_35y_female_NonICU	COVID_55_62y_female_ICU	COVID_56_33y_female_NonICU	
## 1	0.25	0.09	0.28
## 2	0.00	0.00	0.00
## 3	0.64	0.09	0.16
## 4	0.10	0.01	0.09
## 5	0.00	0.00	0.23
## 6	0.00	0.00	0.00
## COVID_57_30y_female_NonICU	COVID_58_62y_male_NonICU	COVID_59_55y_male_NonICU	
## 1	0.42	0.39	0.33
## 2	0.00	0.00	0.00
## 3	0.27	0.08	0.10
## 4	0.01	0.00	0.00
## 5	0.19	0.00	0.07
## 6	0.05	0.00	0.00
## COVID_60_49y_male_NonICU	COVID_61_54y_female_NonICU	COVID_62_78y_female_ICU	
## 1	0.22	0.25	0.21
## 2	0.00	0.00	0.00
## 3	0.14	0.10	0.04
## 4	0.00	0.03	0.00
## 5	0.00	0.13	0.05
## 6	0.02	0.00	0.00
## COVID_63_39y_female_ICU	COVID_64_65y_male_ICU	COVID_65_84y_male_NonICU	
## 1	0.29	0.38	0.40
## 2	0.00	0.01	0.01
## 3	0.01	0.04	0.07
## 4	0.00	0.02	0.00
## 5	0.14	0.56	0.58
## 6	0.00	0.00	0.00
## COVID_66_66y_female_NonICU	COVID_67_57y_male_ICU	COVID_68_79y_male_ICU	
## 1	0.64	0.37	0.58
## 2	0.00	0.00	0.00
## 3	0.00	0.35	0.15
## 4	0.00	0.00	0.01
## 5	0.00	0.00	0.00
## 6	0.00	0.00	0.05
## COVID_69_77y_female_NonICU	COVID_70_81y_male_NonICU	COVID_71_37y_male_ICU	
## 1	0.52	0.27	0.07
## 2	0.00	0.00	0.01
## 3	0.29	0.07	0.12
## 4	0.02	0.00	0.01
## 5	0.00	0.00	0.00
## 6	0.00	0.06	0.00
## COVID_72_50y_female_NonICU	COVID_73_82y_male_NonICU	COVID_74_55y_female_ICU	
## 1	0.52	0.46	0.24
## 2	0.00	0.01	0.00
## 3	0.10	0.02	0.12
## 4	0.01	0.02	0.02
## 5	0.00	0.17	0.26
## 6	0.00	0.04	0.00
## COVID_75_55y_male_NonICU	COVID_76_73y_female_ICU	COVID_77_55y_female_ICU	
## 1	0.23	0.17	0.05
## 2	0.01	0.00	0.00
## 3	0.14	0.09	0.01

## 4	0.00	0.01	0.00
## 5	0.00	0.04	0.00
## 6	0.00	0.00	0.00
## COVID_78_80y_male_NonICU	COVID_79_27y_male_NonICU	COVID_80_71y_male_ICU	
## 1	0.19	0.08	0.28
## 2	0.00	0.01	0.00
## 3	0.20	0.03	0.05
## 4	0.00	0.00	0.00
## 5	0.00	0.00	0.05
## 6	0.00	0.00	0.00
## COVID_82_67y_male_NonICU	COVID_83_85y_female_NonICU		
## 1	0.39	0.47	
## 2	0.01	0.00	
## 3	0.10	0.18	
## 4	0.00	0.05	
## 5	0.00	0.00	
## 6	0.00	0.00	
## COVID_84_75y_female_NonICU	COVID_85_62y_male_ICU	COVID_86_52y_female_NonICU	
## 1	0.35	0.29	0.60
## 2	0.00	0.00	0.00
## 3	0.03	0.04	0.27
## 4	0.00	0.00	0.02
## 5	0.17	0.00	0.00
## 6	0.00	0.00	0.00
## COVID_87_61y_male_ICU	COVID_89_90y_female_NonICU	COVID_90_86y_female_NonICU	
## 1	0.65	0.20	0.40
## 2	0.00	0.00	0.00
## 3	0.15	0.07	0.05
## 4	0.00	0.03	0.01
## 5	0.00	0.14	0.31
## 6	0.00	0.00	0.02
## COVID_91_29y_female_NonICU	COVID_92_82y_female_ICU	COVID_93_81y_female_ICU	
## 1	0.60	0.34	0.37
## 2	0.00	0.00	0.00
## 3	0.03	0.02	0.11
## 4	0.02	0.04	0.00
## 5	0.05	0.58	0.05
## 6	0.00	0.00	0.00
## COVID_94_24y_female_NonICU	COVID_95_49y_male_NonICU	COVID_96_51y_male_NonICU	
## 1	0.81	0.37	1.61
## 2	0.00	0.01	0.00
## 3	0.17	0.20	0.02
## 4	0.02	0.02	0.00
## 5	0.00	0.15	0.00
## 6	0.06	0.00	0.00
## COVID_97_76y_male_ICU	COVID_98_81y_male_NonICU	COVID_99_71y_male_ICU	
## 1	0.19	0.78	0.33
## 2	0.00	0.00	0.00
## 3	0.02	0.26	0.02
## 4	0.05	0.00	0.00
## 5	0.12	0.37	0.04
## 6	0.03	0.00	0.00
## COVID_100_74y_female_NonICU	COVID_101_58y_male_ICU	COVID_102_84y_male_NonICU	
## 1	0.30	0.33	0.12

## 2	0.00	0.00	0.00
## 3	0.09	0.11	0.01
## 4	0.00	0.03	0.01
## 5	0.04	0.05	0.00
## 6	0.00	0.00	0.07
## COVID_103_83y_male_NonICU	NONCOVID_01_54y_female_NonICU		
## 1	0.20	0.89	
## 2	0.00	0.00	
## 3	0.03	0.04	
## 4	0.03	0.00	
## 5	0.04	0.00	
## 6	0.00	0.00	
## NONCOVID_02_65y_male_ICU	NONCOVID_03_65y_male_ICU	NONCOVID_04_90y_male_NonICU	
## 1	0.32	0.44	0.21
## 2	0.00	0.00	0.00
## 3	0.01	0.05	0.05
## 4	0.00	0.02	0.00
## 5	0.04	0.04	0.21
## 6	0.00	0.00	0.00
## NONCOVID_05_83y_female_NonICU	NONCOVID_06_75y_female_ICU		
## 1	0.31	0.89	
## 2	0.00	0.00	
## 3	0.01	0.14	
## 4	0.01	0.01	
## 5	0.00	0.00	
## 6	0.00	0.06	
## NONCOVID_07_50y_male_ICU	NONCOVID_08_53y_female_ICU		
## 1	0.45	0.47	
## 2	0.00	0.01	
## 3	0.07	0.04	
## 4	0.02	0.00	
## 5	0.00	0.15	
## 6	0.00	0.00	
## NONCOVID_09_49y_female_NonICU	NONCOVID_10_67y_male_ICU		
## 1	0.40	0.33	
## 2	0.00	0.00	
## 3	0.04	0.05	
## 4	0.00	0.01	
## 5	0.00	0.23	
## 6	0.00	0.08	
## NONCOVID_11_58y_female_NonICU	NONCOVID_12_82y_male_ICU		
## 1	0.58	0.12	
## 2	0.00	0.00	
## 3	0.03	0.02	
## 4	0.00	0.00	
## 5	0.00	0.00	
## 6	0.00	0.02	
## NONCOVID_13_65y_male_ICU	NONCOVID_14_75y_female_ICU		
## 1	0.31	0.16	
## 2	0.00	0.00	
## 3	0.04	0.08	
## 4	0.01	0.00	
## 5	0.32	0.05	
## 6	0.02	0.02	

```
##   NONCOVID_15_83y_unknown_ICU NONCOVID_16_40y_female_ICU
## 1                        0.59                        0.34
## 2                        0.00                        0.00
## 3                        0.03                        0.07
## 4                        0.04                        0.00
## 5                        0.00                        0.13
## 6                        0.19                        0.00
##   NONCOVID_17_84y_female_ICU NONCOVID_18_88y_male_ICU
## 1                        0.37                        0.33
## 2                        0.00                        0.00
## 3                        0.07                        0.06
## 4                        0.01                        0.00
## 5                        0.18                        0.00
## 6                        0.00                        0.00
##   NONCOVID_19_66y_female_ICU NONCOVID_20_62y_female_ICU
## 1                        0.25                        0.20
## 2                        0.00                        0.00
## 3                        0.11                        0.01
## 4                        0.00                        0.02
## 5                        0.04                        0.00
## 6                        0.03                        0.07
##   NONCOVID_21_71y_male_NonICU NONCOVID_22_63y_male_NonICU
## 1                        0.40                        0.30
## 2                        0.00                        0.00
## 3                        0.04                        0.02
## 4                        0.02                        0.02
## 5                        0.00                        0.00
## 6                        0.00                        0.00
##   NONCOVID_23_42y_female_NonICU NONCOVID_24_32y_female_NonICU
## 1                        0.70                        0.75
## 2                        0.00                        0.00
## 3                        0.02                        0.27
## 4                        0.01                        0.00
## 5                        0.00                        0.06
## 6                        0.00                        0.00
##   NONCOVID_25_62y_male_NonICU NONCOVID_26_36y_male_ICU
## 1                        2.80                        0.22
## 2                        0.00                        0.00
## 3                        0.04                        0.28
## 4                        0.00                        0.00
## 5                        0.00                        0.00
## 6                        0.00                        0.00
```

Formatting and cleaning data

```
# goal is to merge both dataframes - we can merge them based on participant ID

# transposing the genes df and turning it back to a df so it can have 'participant_id' as rows (the same as columns)
genes_transposed <- as.data.frame(t(genes))

# assigning row x with gene name to be the column headers
colnames(genes_transposed) <- as.character(genes_transposed[1, ])

# getting rid of the x row because it has redundant info (same values as column names)
```



```

genes_transposed <- genes_transposed[-1, ]

# setting the row name to participant id so we can use this column to merge with metadata
# used this source to find the rownames_to_column function https://forum.posit.co/t/rstudio-rownames-to
genes_transposed <- rownames_to_column(genes_transposed, var = "participant_id")

# reshaping the gene so that genes and expression can be their own columns using pivot_longer
# used this source to learn how to implement pivot_longer: https://tidyr.tidyverse.org/reference/pivot_
genes_long <- genes_transposed %>%
  pivot_longer(
    cols = -participant_id, #accessing all columns with the exception of participant_id because it is
    names_to = "gene", #setting name column to gene
    values_to = "expression" #setting values column to expression
  )

# merging long gene data and series metadata to become one dataframe
all_data <- merge(genes_long, series, by = "participant_id") #merging it on 'participant_id' column

# taking a look at the new comprehensive dataframe
head(all_data)

```

```

##           participant_id      gene expression geo_accession
## 1 COVID_01_39y_male_NonICU    A1CF         0.00    GSM4753021
## 2 COVID_01_39y_male_NonICU    A1BG         0.49    GSM4753021
## 3 COVID_01_39y_male_NonICU    AADAC         0.00    GSM4753021
## 4 COVID_01_39y_male_NonICU AADACL2         0.00    GSM4753021
## 5 COVID_01_39y_male_NonICU AADACL3         0.00    GSM4753021
## 6 COVID_01_39y_male_NonICU AADACL4         0.00    GSM4753021
##           status X.Sample_submission_date last_update_date type
## 1 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
## 2 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
## 3 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
## 4 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
## 5 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
## 6 Public on Aug 29 2020                Aug 28 2020      Aug 29 2020 SRA
##   channel_count      source_name_ch1 organism_ch1
## 1             1 Leukocytes from whole blood Homo sapiens
## 2             1 Leukocytes from whole blood Homo sapiens
## 3             1 Leukocytes from whole blood Homo sapiens
## 4             1 Leukocytes from whole blood Homo sapiens
## 5             1 Leukocytes from whole blood Homo sapiens
## 6             1 Leukocytes from whole blood Homo sapiens
##   disease_status age  sex icu_status apacheii charlson_score
## 1 disease state: COVID-19 39 male      no          15          0
## 2 disease state: COVID-19 39 male      no          15          0
## 3 disease state: COVID-19 39 male      no          15          0
## 4 disease state: COVID-19 39 male      no          15          0
## 5 disease state: COVID-19 39 male      no          15          0
## 6 disease state: COVID-19 39 male      no          15          0
##   mechanical_ventilation ventilator.free_days
## 1                      yes                   0
## 2                      yes                   0
## 3                      yes                   0
## 4                      yes                   0

```

```
## 5          yes          0
## 6          yes          0
##  hospital.free_days_post_45_day_followup ferritin.ng.ml. crp.mg.l.
## 1              0              946      73.1
## 2              0              946      73.1
## 3              0              946      73.1
## 4              0              946      73.1
## 5              0              946      73.1
## 6              0              946      73.1
##  ddimer.mg.l_feu. procalcitonin.ng.ml.. lactate.mmol.l. fibrinogen sofa
## 1          1.3          36          0.9          513      8
## 2          1.3          36          0.9          513      8
## 3          1.3          36          0.9          513      8
## 4          1.3          36          0.9          513      8
## 5          1.3          36          0.9          513      8
## 6          1.3          36          0.9          513      8
```

Choosing focuses of interests: genes, continuous covariates and categorical covariates

gene: AASDHPPT continuous covariate: ferritin levels categorical covariate: disease status, gender

```
#subsetting the dataframe so that it only contains columns that I want to perform further analysis on

#getting rid of all unknown values in the data
unique(all_data$ferritin.ng.ml.)
unique(all_data$Age)
#used the above lines to look at unique values, and saw that unknown is formatted as " unknown", so will
all_data[all_data == " unknown"] <- NA

# Drop rows with any NA values
all_data <- na.omit(all_data)

#using select to subset https://www.educative.io/answers/what-is-the-select-function-in-r
covid_data <- all_data %>% select(participant_id, gene,expression, ferritin.ng.ml., sex, disease_status)

#taking a look at the new subsetted data frame
head(covid_data)
```

```
##      participant_id      gene expression ferritin.ng.ml.    sex
## 1 COVID_01_39y_male_NonICU A1CF      0.00          946  male
## 2 COVID_01_39y_male_NonICU A1BG      0.49          946  male
## 3 COVID_01_39y_male_NonICU AADAC      0.00          946  male
## 4 COVID_01_39y_male_NonICU AADACL2    0.00          946  male
## 5 COVID_01_39y_male_NonICU AADACL3    0.00          946  male
## 6 COVID_01_39y_male_NonICU AADACL4    0.00          946  male
##      disease_status
## 1 disease state: COVID-19
## 2 disease state: COVID-19
## 3 disease state: COVID-19
## 4 disease state: COVID-19
## 5 disease state: COVID-19
## 6 disease state: COVID-19
```

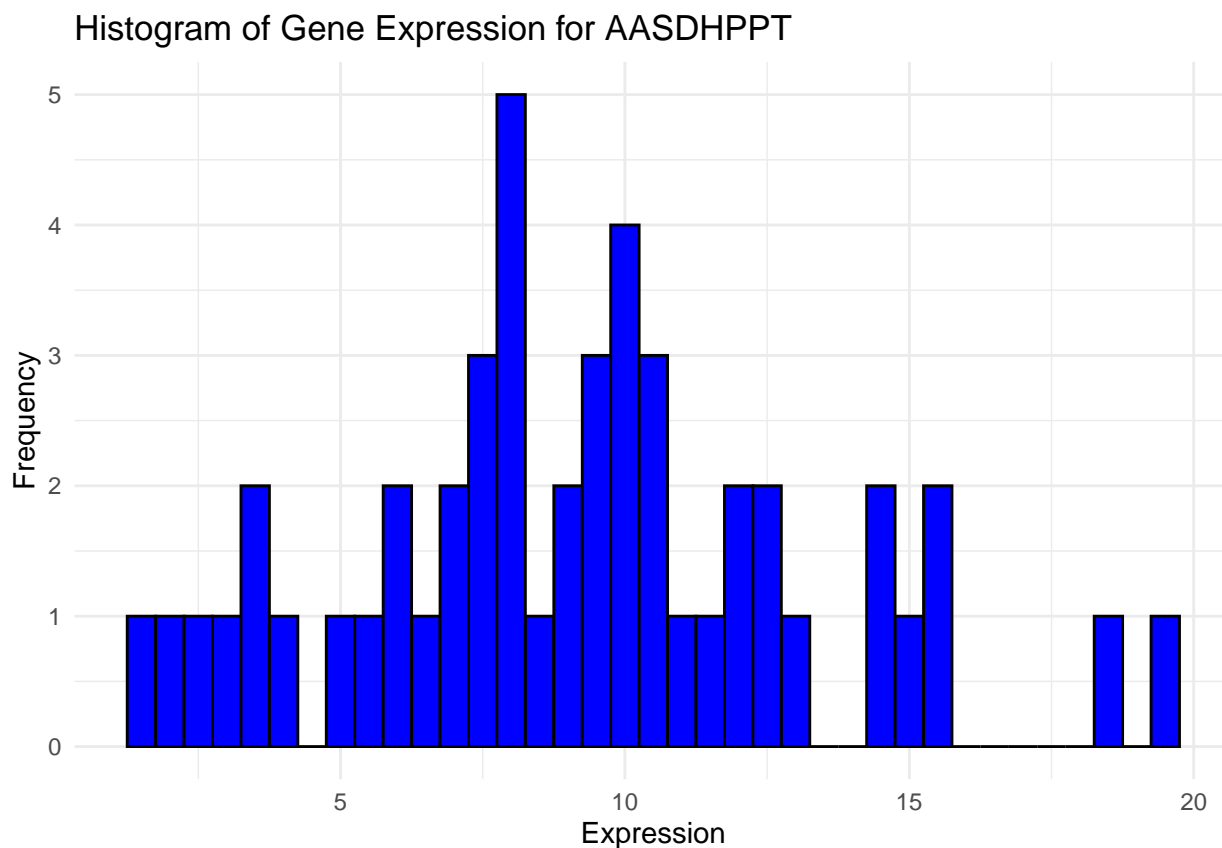
Plotting gene expression (histogram)

```
library(tidyverse)

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v forcats 1.0.0      v purrr 1.0.2
## v lubridate 1.9.2    v stringr 1.5.0
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()      masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors

AASDHPPT_data <- covid_data %>%
  filter(gene == "AASDHPPT") %>%
  select(expression) %>%
  mutate(expression = as.numeric(expression)) # Ensure expression is numeric

# Plot histogram
ggplot(AASDHPPT_data, aes(x = as.numeric(expression))) +
  geom_histogram(binwidth = 0.5, color = "black", fill = "blue") +
  labs(title = "Histogram of Gene Expression for AASDHPPT",
       x = "Expression",
       y = "Frequency") +
  theme_minimal() #choosing preferred theme
```



Plotting gene expression vs. ferritin (scatterplot)

```
# referred to these sites when plotting
# https://r-graph-gallery.com/interactive-charts.html
# https://r-graph-gallery.com/scatterplot.html
# http://www.sthda.com/english/wiki/ggplot2-scatter-plots-quick-start-guide-r-software-and-data-visuali

# using plotly to add interactive element to the graph
#install.packages("plotly")
library(plotly)

##
## Attaching package: 'plotly'

## The following object is masked from 'package:ggplot2':
##
##     last_plot

## The following object is masked from 'package:stats':
##
##     filter

## The following object is masked from 'package:graphics':
##
##     layout

# subsetting the data so that it only contains rows where gene = AASDHPPT and also the sex variable so
data <- covid_data %>%
  filter(gene == "AASDHPPT")

# certain columns in the dataset with numeric values are stored in character types, setting them to num
data$expression <- as.numeric(data$expression)
data$ferritin.ng.ml. <- as.numeric(data$ferritin.ng.ml. )

scatter_plot <- ggplot(data, aes(x = expression, y = ferritin.ng.ml.)) +
  geom_point(color = "purple",size = 3) + #increasing size of the data points on the graph
  geom_smooth(method = "lm", se = FALSE, color = "#33FFF7",linetype = "dashed") +
  labs(title = "AASDHPPT Expression and Ferritin Levels Appear Slightly Negatively Correlated",
        subtitle = "Exploring AASDHPPT Expression vs. Ferritin Levels in Human Subjects",
        x= "AASDHPPT Expression" ,
        y = "Ferritin (ng/mL)" ) +
  theme_bw() # changing to preferred theme +

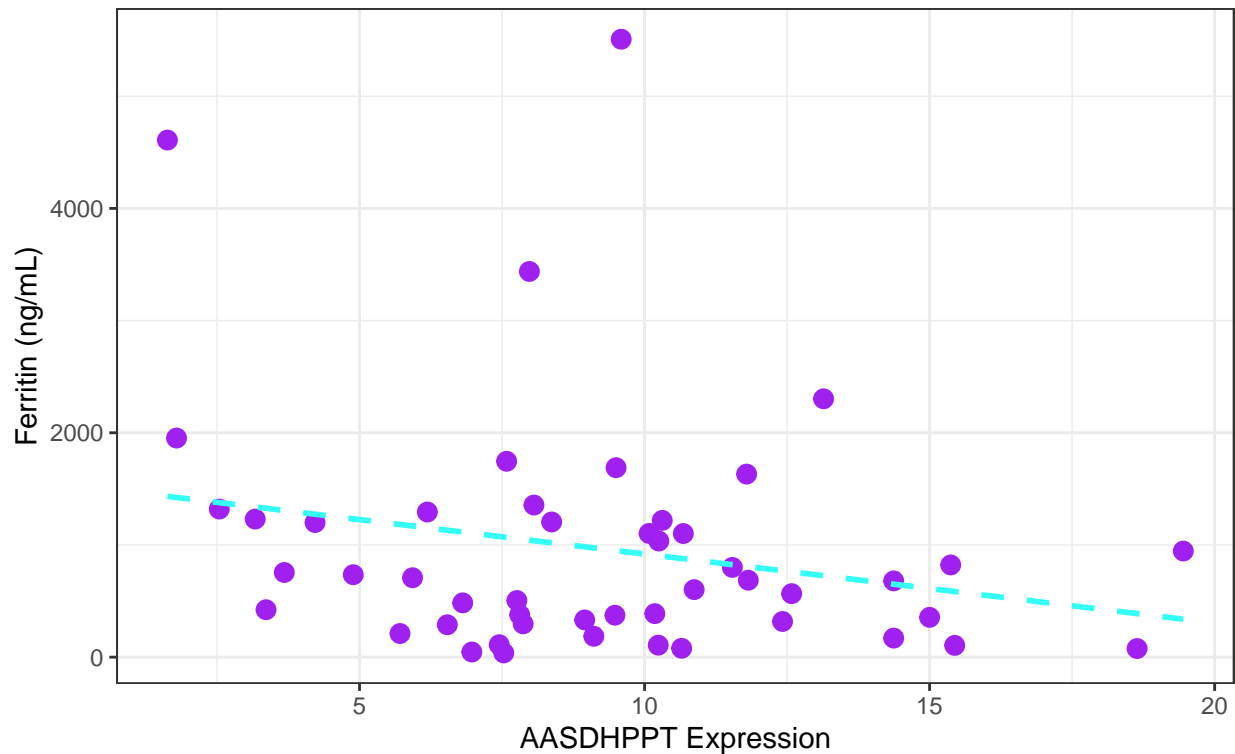
interactive_scatter <- ggplotly(scatter_plot)

## `geom_smooth()` using formula = 'y ~ x'
# displaying regular graph and interactive
scatter_plot

## `geom_smooth()` using formula = 'y ~ x'
```

AASDHPPT Expression and Ferritin Levels Appear Slightly Negatively Correlated

Exploring AASDHPPT Expression vs. Ferritin Levels in Human Subjects



interactive_scatter

Plotting gene expression vs.disease status by sex (boxplot)

```
# creating boxplot
ggplot(data, aes(x = disease_status, y = as.numeric(expression), fill = sex)) +
  geom_boxplot() +
  scale_x_discrete(labels = c("disease state: COVID-19" = "COVID-19 Positive", "disease state: non-COVID-19" = "COVID-19 Negative")) +
  labs(title = "Observing patterns of AASDHPPT Expression by Gender and Disease Status", x = "Disease Status", y = "AASDHPPT Expression") +
  theme_bw() # changing to preferred theme
```

