

Assignment: Data Analysis

Task 1: Data Aggregation Using `aggregate` and `dplyr`

Question 1:

Run the following lines and study how they work. Then state what they do and output for us.

```
# Creating the dataframe
df1 <- data.frame(Name =
  c('James', 'Paul', 'Richards', 'Marico', 'Samantha', 'Ravi', 'Raghu',
  'Richards', 'George', 'Ema', 'Samantha', 'Catherine'),
  State = c('Alaska', 'California', 'Texas', 'North
  Carolina', 'California', 'Texas', 'Alaska', 'Texas', 'North
  Carolina', 'Alaska', 'California', 'Texas'),
  Sales = c(14, 24, 31, 12, 13, 7, 9, 31, 18, 16, 18, 14))

# Aggregating using aggregate
agg_result <- aggregate(df1$Sales, by = list(df1$State), FUN = sum)
print(agg_result)

##           Group.1   x
## 1          Alaska  39
## 2      California  55
## 3 North Carolina  30
## 4          Texas  83

# Aggregating using dplyr
library(dplyr)

## Warning: package 'dplyr' was built under R version 4.5.2

##
## 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

dplyr_result <- df1 %>%
  group_by(State) %>%
  summarise(sum_sales = sum(Sales))
print(dplyr_result)
```

```

## # A tibble: 4 × 2
##   State      sum_sales
##   <chr>        <dbl>
## 1 Alaska       39
## 2 California   55
## 3 North Carolina 30
## 4 Texas        83

```

Explanation:

- The above code creates a data frame df1 containing Name, State, and Sales.
- Then it calculates the sum of sales grouped by State using both aggregate() and dplyr package.

Output: - The output will show the sum of Sales for each state using both methods (aggregate() and dplyr).

Task 2: World Cup Matches Data Analysis

Question 2a:

Use R to read the WorldCupMatches.csv from the DATA folder on Google Drive. Then perform the following:

(a) *Find the size of the data frame. How many rows, how many columns?*

```

# Read the WorldCupMatches dataset
wc_data <- read.csv("C:\\\\Users\\\\Lenovo\\\\Downloads\\\\WorldCupMatches.csv")

# Find the size of the data frame
data_size <- dim(wc_data)
data_size

## [1] 852  20

```

Explanation:

- The dim() function returns the number of rows and columns in the data frame.

Output: - The number of rows and columns in the WorldCupMatches dataset.

(b) *Use summary function to report the statistical summary of your data.*

```

# Get the statistical summary of the dataset
summary(wc_data)

##          Year           Datetime            Stage           Stadium
##  Min.   :1930   Length:852             Length:852         Length:852
##  1st Qu.:1970   Class :character    Class :character   Class :character
##  Median :1990   Mode  :character    Mode  :character   Mode  :character
##  Mean   :1985

```

```

## 3rd Qu.:2002
## Max. :2014
##
##      City          Home.Team.Name   Home.Team.Goals  Away.Team.Goals
## Length:852          Length:852       Min. : 0.000  Min. :0.000
## Class :character    Class :character  1st Qu.: 1.000  1st Qu.:0.000
## Mode  :character    Mode  :character  Median : 2.000  Median :1.000
##                               Mean  : 1.811  Mean  :1.022
##                               3rd Qu.: 3.000  3rd Qu.:2.000
##                               Max. :10.000 Max. :7.000
##
##      Away.Team.Name   Win.conditions   Attendance
## Half.time.Home.Goals
## Length:852          Length:852       Min. : 2000  Min. :0.0000
## Class :character    Class :character  1st Qu.: 30000 1st Qu.:0.0000
## Mode  :character    Mode  :character  Median : 41580  Median :0.0000
##                               Mean  : 45165  Mean  :0.7089
##                               3rd Qu.: 61375  3rd Qu.:1.0000
##                               Max. :173850  Max. :6.0000
##                               NA's  :2
##      Half.time.Away.Goals  Referee        Assistant.1  Assistant.2
## Min. :0.0000          Length:852       Length:852       Length:852
## 1st Qu.:0.0000          Class :character  Class :character  Class
## :character
## Median :0.0000          Mode  :character  Mode  :character  Mode
## :character
## Mean   :0.4284
## 3rd Qu.:1.0000
## Max.  :5.0000
##
##      RoundID          MatchID   Home.Team.Initials
## Away.Team.Initials
## Min.  : 201  Min.  :     25  Length:852       Length:852
## 1st Qu.: 262  1st Qu.: 1189  Class :character  Class
## :character
## Median : 337  Median : 2191  Mode  :character  Mode
## :character
## Mean   :10661773  Mean   : 61346868
## 3rd Qu.: 249722  3rd Qu.: 43950059
## Max.  :97410600  Max.  :300186515
##

```

Output: - A summary of each column, including mean, min, max, and other statistics.

(c) *Find how many unique locations Olympics were held at.*

```

# Find unique Locations
unique_locations <- length(unique(wc_data$Location))
unique_locations

## [1] 0

```

Output: - The number of unique locations where World Cup matches were held.

(d) Find the average attendance.

```
# Find average attendance
avg_attendance <- mean(wc_data$Attendance, na.rm = TRUE)
avg_attendance

## [1] 45164.8
```

Output: - The average attendance at the World Cup matches.

(e) For each Home Team, what is the total number of goals scored?

```
# Aggregating goals scored by Home Team using dplyr
home_goals <- wc_data %>%
  group_by(`Home.Team.Name`) %>%
  summarise(total_goals = sum(`Home.Team.Goals`, na.rm = TRUE))
print(home_goals)

## # A tibble: 78 × 2
##   Home.Team.Name total_goals
##   <chr>           <int>
## 1 Algeria          5
## 2 Angola            0
## 3 Argentina        111
## 4 Australia         7
## 5 Austria           31
## 6 Belgium           27
## 7 Bolivia            1
## 8 Brazil           180
## 9 Bulgaria          11
## 10 Cameroon          11
## # ... with 68 more rows
```

Output: - The total number of goals scored by each Home Team.

(f) What is the average number of attendees for each year?

```
# Average attendance for each year
avg_attendance_year <- wc_data %>%
  group_by(Year) %>%
  summarise(avg_attendance = mean(Attendance, na.rm = TRUE))
avg_attendance_year

## # A tibble: 20 × 2
##   Year avg_attendance
##   <int>      <dbl>
## 1 1930       32808.
## 2 1934       21353.
## 3 1938       20872.
## 4 1950       47511.
## 5 1954       29562.
## 6 1958       23423.
```

```

## 7 1962      27912.
## 8 1966      48848.
## 9 1970      50124.
## 10 1974     49099.
## 11 1978     40679.
## 12 1982     40572.
## 13 1986     46039.
## 14 1990     48389.
## 15 1994     68991.
## 16 1998     43517.
## 17 2002     42269.
## 18 2006     52491.
## 19 2010     49670.
## 20 2014     55375.

```

Output: - The average attendance for each year, along with any trends or patterns.

Task 3: Metabolites Data Analysis

Question 3:

Use R to read the metabolites.csv from the DATA folder on Google Drive. Then perform the following:

(a) *Find how many Alzheimer's patients there are in the data set.*

```

# Read the metabolites dataset
metabolites_data <- read.csv("C:\\\\Users\\\\Lenovo\\\\Downloads\\\\metabolite.csv")

# Find number of Alzheimer's patients
alzheimers_patients <- sum(metabolites_data$Diagnosis == 'Alzheimer', na.rm = TRUE)
alzheimers_patients

## [1] 0

```

Explanation:

- Counts the number of Alzheimer's patients in the dataset by checking the Diagnosis column.

Output: - The count of Alzheimer's patients in the dataset.

(b) *Determine the number of missing values for each column.*

```

# Find missing values for each column
missing_values <- colSums(is.na(metabolites_data))
missing_values

```

| ## | Label | Phe | Pro |
|-----------|------------|------------|-----------|
| Ser | 0 | 0 | 0 |
| ## | Thr | ADMA | alpha.AAA |
| c4.OH.Pro | 0 | 0 | 0 |
| ## | Carnosine | Creatinine | DOPA |
| Dopamine | 1 | 0 | 0 |
| ## | Histamine | Kynurenine | Met.SO |
| Nitro.Tyr | 0 | 0 | 1 |
| ## | PEA | Putrescine | Sarcosine |
| Serotonin | 69 | 0 | 0 |
| ## | Spermidine | Spermine | t4.OH.Pro |
| Taurine | 0 | 60 | 0 |
| ## | SDMA | C0 | C10 |
| C10.1 | 0 | 0 | 0 |
| ## | C10.2 | C12 | C12.DC |
| C12.1 | 0 | 0 | 1 |
| ## | C14 | C14.1 | C14.1.OH |
| C14.2 | 0 | 0 | 1 |
| ## | C14.2.OH | C16 | C16.OH |
| C16.1 | 2 | 0 | 1 |
| ## | C16.1.OH | C16.2 | C16.2.OH |
| C18 | 2 | 2 | 1 |
| ## | C18.1 | C18.1.OH | C18.2 |
| C2 | 0 | 7 | 0 |
| ## | C3 | C3.OH | C3.1 |
| C4 | | | |

| | | | |
|----------------|-----------------|----------------|----------------|
| ## | 0 | 8 | 2 |
| 0 | | | |
| ## | C3.DC..C4.OH. | C4.1 | C5 |
| C5.M.DC | | | |
| ## | 0 | 0 | 0 |
| 1 | | | |
| ## | C5.OH..C3.DC.M. | C5.1 | C5.1.DC |
| C6..C4.1.DC. | | | |
| ## | 0 | 5 | 2 |
| 0 | | | |
| ## | C5.DC..C6.OH. | C6.1 | C7.DC |
| C8 | | | |
| ## | 4 | 2 | 1 |
| 0 | | | |
| ## | C9 | lysoPC.a.C14.0 | lysoPC.a.C16.0 |
| lysoPC.a.C16.1 | | | |
| ## | 1 | 0 | 0 |
| 0 | | | |
| ## | lysoPC.a.C17.0 | lysoPC.a.C18.0 | lysoPC.a.C18.1 |
| lysoPC.a.C18.2 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | lysoPC.a.C20.3 | lysoPC.a.C20.4 | lysoPC.a.C24.0 |
| lysoPC.a.C26.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | lysoPC.a.C26.1 | lysoPC.a.C28.0 | lysoPC.a.C28.1 |
| PC.aa.C24.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C26.0 | PC.aa.C28.1 | PC.aa.C30.0 |
| PC.aa.C32.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C32.1 | PC.aa.C32.2 | PC.aa.C32.3 |
| PC.aa.C34.1 | | | |
| ## | 0 | 47 | 0 |
| 0 | | | |
| ## | PC.aa.C34.2 | PC.aa.C34.3 | PC.aa.C34.4 |
| PC.aa.C36.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C36.1 | PC.aa.C36.2 | PC.aa.C36.3 |
| PC.aa.C36.4 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C36.5 | PC.aa.C36.6 | PC.aa.C38.0 |
| PC.aa.C38.3 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |

| | | | |
|-------------|-------------|-------------|-------------|
| ## | PC.aa.C38.4 | PC.aa.C38.5 | PC.aa.C38.6 |
| PC.aa.C40.1 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C40.2 | PC.aa.C40.3 | PC.aa.C40.4 |
| PC.aa.C40.5 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C40.6 | PC.aa.C42.0 | PC.aa.C42.1 |
| PC.aa.C42.2 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.aa.C42.4 | PC.aa.C42.5 | PC.aa.C42.6 |
| PC.ae.C30.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C30.1 | PC.ae.C30.2 | PC.ae.C32.1 |
| PC.ae.C32.2 | | | |
| ## | 10 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C34.0 | PC.ae.C34.1 | PC.ae.C34.2 |
| PC.ae.C34.3 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C36.0 | PC.ae.C36.1 | PC.ae.C36.2 |
| PC.ae.C36.3 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C36.4 | PC.ae.C36.5 | PC.ae.C38.0 |
| PC.ae.C38.1 | | | |
| ## | 0 | 0 | 0 |
| 52 | | | |
| ## | PC.ae.C38.2 | PC.ae.C38.3 | PC.ae.C38.4 |
| PC.ae.C38.5 | | | |
| ## | 19 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C38.6 | PC.ae.C40.1 | PC.ae.C40.2 |
| PC.ae.C40.3 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C40.4 | PC.ae.C40.5 | PC.ae.C40.6 |
| PC.ae.C42.0 | | | |
| ## | 0 | 0 | 0 |
| 0 | | | |
| ## | PC.ae.C42.1 | PC.ae.C42.2 | PC.ae.C42.3 |
| PC.ae.C42.4 | | | |
| ## | 0 | 1 | 0 |
| 0 | | | |
| ## | PC.ae.C42.5 | PC.ae.C44.3 | PC.ae.C44.4 |
| PC.ae.C44.5 | | | |

```

##          0          0          0
0
##      PC.ae.C44.6    SM..OH..C14.1    SM..OH..C16.1
SM..OH..C22.1
##          0          0          0
0
##      SM..OH..C22.2    SM..OH..C24.1    SM.C16.0
SM.C16.1
##          0          0          0
0
##      SM.C18.0    SM.C18.1    SM.C20.2
SM.C24.0
##          0          0          0
0
##      SM.C24.1    SM.C26.0    SM.C26.1
H1_1
##          0          0          0
0
##          H1        Urea_N    L.Arginine_N
L.Leucine_N
##          0          1          1
1
##      EDTAca_N X2.Hydroxybutyrate X3.Hydroxybutyrate
Acetate
##          1          1          1
1
##      Acetoacetate    Acetone    Betaine
Carnitine
##          1          1          1
1
##      Choline    Creatine    Dimethyl.sulfone
Ethanol
##          1          1          1
2
##      Formate    Glucose    Glycerol
Hypoxanthine
##          2          1          1
1
##      Isobutyrate    Isopropanol    Lactate
Malonate
##          1          1          1
1

```

Output: - The number of missing values for each column.

(c) Remove the rows which have missing values for the Dopamine column.

```

# Remove rows where Dopamine column has missing values
metabolites_no_na_dopamine <-
metabolites_data[ !is.na(metabolites_data$Dopamine), ]
head(metabolites_no_na_dopamine)

```



```

C3
## 1 0.008 0.009 0.007 0.005 0.013 0.013 0.024 0.003 0.016 1.97
0.354
## 4 0.011 0.015 0.008 0.006 0.009 0.020 0.035 0.004 0.033 2.10
0.278
## 5 NA 0.024 0.014 0.012 0.025 0.031 0.034 0.012 0.017 5.62
0.436
## 8 0.006 0.010 0.005 0.004 0.008 0.020 0.025 0.004 0.019 1.66
0.258
## 9 0.005 0.012 0.005 0.005 0.009 0.014 0.026 0.003 0.016 2.21
0.233
## 10 0.009 0.012 0.007 0.006 0.014 0.024 0.031 0.004 0.028 2.58
0.364
## C3.OH C3.1 C4 C3.DC..C4.OH. C4.1 C5 C5.M.DC C5.OH..C3.DC.M.
C5.1
## 1 0.008 0.015 0.082 0.045 0.025 0.094 0.023 0.026
0.030
## 4 0.010 0.017 0.110 0.077 0.031 0.145 0.034 0.041
0.035
## 5 0.029 0.035 0.106 0.099 0.069 0.141 0.094 0.058
0.073
## 8 0.008 0.012 0.082 0.047 0.021 0.107 0.023 0.023
0.021
## 9 0.008 0.014 0.088 0.029 0.024 0.127 0.024 0.024
0.025
## 10 0.010 0.015 0.100 0.038 0.025 0.292 0.024 0.023
NA
## C5.1.DC C6..C4.1.DC. C5.DC..C6.OH. C6.1 C7.DC C8 C9
lysoPC.a.C14.0
## 1 0.020 0.022 0.014 0.018 0.011 0.062 0.016
2.23
## 4 0.016 0.029 0.016 0.027 0.017 0.091 0.018
2.19
## 5 0.049 0.052 0.040 0.040 0.036 0.192 0.041
1.88
## 8 0.017 0.036 0.011 NA 0.009 0.062 0.011
2.13
## 9 0.016 0.026 0.018 0.015 0.013 0.064 0.014
2.10
## 10 0.014 0.044 0.017 0.018 0.014 0.067 0.013
2.06
## lysoPC.a.C16.0 lysoPC.a.C16.1 lysoPC.a.C17.0 lysoPC.a.C18.0
lysoPC.a.C18.1
## 1 37.9 2.66 0.446 9.00
8.58
## 4 32.8 2.39 0.323 7.21
7.22
## 5 24.5 1.27 0.382 6.66
5.39
## 8 33.7 3.09 0.455 6.96

```

| | | | | | |
|-------------------|----------------|----------------|----------------|-------------|-------------|
| 7.31 | | | | | |
| ## 9 | 36.0 | 3.46 | 0.435 | 7.27 | |
| 8.11 | | | | | |
| ## 10 | 32.7 | 2.48 | 0.368 | 6.49 | |
| 7.86 | | | | | |
| ## lysoPC.a.C18.2 | lysoPC.a.C20.3 | lysoPC.a.C20.4 | lysoPC.a.C24.0 | | |
| lysoPC.a.C26.0 | | | | | |
| ## 1 | 7.27 | 1.83 | 8.25 | 0.079 | |
| 0.113 | | | | | |
| ## 4 | 7.62 | 1.64 | 6.75 | 0.066 | |
| 0.086 | | | | | |
| ## 5 | 3.60 | 0.97 | 6.26 | 0.084 | |
| 0.118 | | | | | |
| ## 8 | 7.53 | 2.35 | 8.73 | 0.061 | |
| 0.083 | | | | | |
| ## 9 | 6.75 | 2.08 | 7.82 | 0.068 | |
| 0.083 | | | | | |
| ## 10 | 10.10 | 1.86 | 12.40 | 0.061 | |
| 0.086 | | | | | |
| ## lysoPC.a.C26.1 | lysoPC.a.C28.0 | lysoPC.a.C28.1 | PC.aa.C24.0 | PC.aa.C26.0 | |
| ## 1 | 0.053 | 0.108 | 0.072 | 0.082 | 0.438 |
| ## 4 | 0.045 | 0.076 | 0.076 | 0.076 | 0.486 |
| ## 5 | 0.053 | 0.092 | 0.072 | 0.069 | 0.401 |
| ## 8 | 0.047 | 0.124 | 0.078 | 0.082 | 0.424 |
| ## 9 | 0.037 | 0.096 | 0.061 | 0.064 | 0.430 |
| ## 10 | 0.052 | 0.088 | 0.074 | 0.065 | 0.453 |
| ## PC.aa.C28.1 | PC.aa.C30.0 | PC.aa.C32.0 | PC.aa.C32.1 | PC.aa.C32.2 | PC.aa.C32.3 |
| ## 1 | 0.571 | 2.35 | 11.4 | 9.22 | NA |
| ## 4 | 0.685 | 3.33 | 18.6 | 13.30 | 0.053 |
| ## 5 | 0.513 | 1.78 | 13.8 | 5.03 | NA |
| ## 8 | 0.605 | 2.32 | 12.1 | 11.50 | NA |
| ## 9 | 0.486 | 2.05 | 10.4 | 11.40 | NA |
| ## 10 | 0.578 | 2.28 | 13.8 | 8.10 | NA |
| ## PC.aa.C34.1 | PC.aa.C34.2 | PC.aa.C34.3 | PC.aa.C34.4 | PC.aa.C36.0 | PC.aa.C36.1 |
| ## 1 | 109.0 | 71.0 | 1.430 | 0.200 | 2.38 |
| ## 4 | 106.0 | 93.6 | 1.590 | 0.190 | 2.57 |
| ## 5 | 83.4 | 35.9 | 0.709 | 0.135 | 1.83 |
| ## 8 | 83.6 | 60.6 | 1.580 | 0.251 | 2.16 |
| ## 9 | 89.6 | 55.1 | 1.390 | 0.221 | 1.62 |
| ## 10 | 80.7 | 82.6 | 1.640 | 0.244 | 1.75 |
| ## PC.aa.C36.2 | PC.aa.C36.3 | PC.aa.C36.4 | PC.aa.C36.5 | PC.aa.C36.6 | PC.aa.C38.0 |
| ## 1 | 42.4 | 42.7 | 120.0 | 1.86 | 0.084 |
| ## 4 | 48.8 | 41.2 | 122.0 | 1.76 | 0.070 |
| ## 5 | 28.5 | 21.9 | 98.1 | 1.70 | 0.048 |
| ## 8 | 34.4 | 41.5 | 110.0 | 2.03 | 0.106 |
| ## 9 | 32.0 | 41.8 | 102.0 | 1.77 | 0.081 |
| ## 10 | 38.1 | 31.2 | 138.0 | 1.80 | 0.092 |
| ## PC.aa.C38.3 | PC.aa.C38.4 | PC.aa.C38.5 | PC.aa.C38.6 | PC.aa.C40.1 | PC.aa.C40.2 |
| ## 1 | 32.1 | 95.1 | 16.8 | 41.6 | 0.195 |
| ## 4 | 28.7 | 92.7 | 14.3 | 29.9 | 0.220 |
| | | | | | 0.097 |

| | | | | | | |
|----------------|-------------|-------------|-------------|-------------|-------------|-------|
| ## 5 | 23.3 | 101.0 | 13.8 | 36.2 | 0.165 | 0.044 |
| ## 8 | 31.3 | 81.7 | 14.5 | 42.8 | 0.225 | 0.099 |
| ## 9 | 31.1 | 75.9 | 13.6 | 37.6 | 0.181 | 0.100 |
| ## 10 | 22.2 | 98.2 | 12.7 | 31.5 | 0.189 | 0.130 |
| ## PC.aa.C40.3 | PC.aa.C40.4 | PC.aa.C40.5 | PC.aa.C40.6 | PC.aa.C42.0 | PC.aa.C42.1 | |
| ## 1 | 0.491 | 3.48 | 5.66 | 21.8 | 0.364 | 0.226 |
| ## 4 | 0.433 | 3.59 | 5.06 | 14.0 | 0.427 | 0.223 |
| ## 5 | 0.525 | 3.37 | 5.29 | 22.5 | 0.125 | 0.095 |
| ## 8 | 0.384 | 4.02 | 5.49 | 20.6 | 0.365 | 0.193 |
| ## 9 | 0.298 | 2.88 | 4.88 | 17.5 | 0.396 | 0.216 |
| ## 10 | 0.315 | 2.87 | 3.32 | 12.6 | 0.301 | 0.186 |
| ## PC.aa.C42.2 | PC.aa.C42.4 | PC.aa.C42.5 | PC.aa.C42.6 | PC.ae.C30.0 | PC.ae.C30.1 | |
| ## 1 | 0.108 | 0.272 | 0.272 | 0.291 | 0.173 | 0.027 |
| ## 4 | 0.119 | 0.268 | 0.267 | 0.254 | 0.223 | 0.049 |
| ## 5 | 0.083 | 0.206 | 0.205 | 0.280 | 0.095 | 0.082 |
| ## 8 | 0.089 | 0.266 | 0.237 | 0.288 | 0.212 | 0.028 |
| ## 9 | 0.086 | 0.215 | 0.206 | 0.223 | 0.179 | 0.012 |
| ## 10 | 0.092 | 0.228 | 0.199 | 0.224 | 0.164 | 0.017 |
| ## PC.ae.C30.2 | PC.ae.C32.1 | PC.ae.C32.2 | PC.ae.C34.0 | PC.ae.C34.1 | PC.ae.C34.2 | |
| ## 1 | 0.022 | 1.65 | 0.371 | 0.880 | 3.66 | 2.48 |
| ## 4 | 0.023 | 2.47 | 0.459 | 0.964 | 4.06 | 3.09 |
| ## 5 | 0.023 | 1.72 | 0.316 | 1.060 | 3.28 | 1.70 |
| ## 8 | 0.033 | 1.68 | 0.335 | 1.140 | 3.51 | 2.28 |
| ## 9 | 0.022 | 1.53 | 0.291 | 0.683 | 3.20 | 2.03 |
| ## 10 | 0.024 | 1.70 | 0.341 | 0.800 | 3.56 | 2.52 |
| ## PC.ae.C34.3 | PC.ae.C36.0 | PC.ae.C36.1 | PC.ae.C36.2 | PC.ae.C36.3 | PC.ae.C36.4 | |
| ## 1 | 0.813 | 0.498 | 5.64 | 1.90 | 1.17 | 6.96 |
| ## 4 | 1.020 | 0.552 | 4.75 | 2.01 | 1.35 | 8.36 |
| ## 5 | 0.722 | 0.553 | 5.95 | 1.47 | 0.76 | 4.78 |
| ## 8 | 0.868 | 0.479 | 5.65 | 1.97 | 1.25 | 7.15 |
| ## 9 | 0.663 | 0.384 | 4.79 | 1.54 | 1.12 | 6.43 |
| ## 10 | 0.983 | 0.498 | 4.85 | 1.93 | 1.04 | 7.08 |
| ## PC.ae.C36.5 | PC.ae.C38.0 | PC.ae.C38.1 | PC.ae.C38.2 | PC.ae.C38.3 | PC.ae.C38.4 | |
| ## 1 | 4.79 | 0.474 | 0.287 | 0.538 | 2.66 | 6.33 |
| ## 4 | 5.97 | 0.397 | 0.022 | 0.144 | 1.97 | 5.99 |
| ## 5 | 4.00 | 0.430 | 0.271 | 0.246 | 1.80 | 5.45 |
| ## 8 | 4.04 | 0.550 | 0.078 | 0.559 | 2.90 | 5.73 |
| ## 9 | 3.32 | 0.442 | NA | 0.224 | 2.53 | 5.03 |
| ## 10 | 4.52 | 0.458 | NA | 0.053 | 1.99 | 5.79 |
| ## PC.ae.C38.5 | PC.ae.C38.6 | PC.ae.C40.1 | PC.ae.C40.2 | PC.ae.C40.3 | PC.ae.C40.4 | |
| ## 1 | 5.51 | 1.95 | 0.574 | 0.575 | 0.940 | 1.76 |
| ## 4 | 5.63 | 1.97 | 0.425 | 0.540 | 0.742 | 1.45 |
| ## 5 | 4.34 | 1.51 | 0.430 | 0.432 | 0.632 | 1.10 |
| ## 8 | 4.53 | 1.71 | 0.584 | 0.499 | 0.991 | 1.84 |
| ## 9 | 3.82 | 1.28 | 0.472 | 0.555 | 0.737 | 1.35 |
| ## 10 | 4.84 | 1.58 | 0.670 | 0.572 | 0.757 | 1.16 |
| ## PC.ae.C40.5 | PC.ae.C40.6 | PC.ae.C42.0 | PC.ae.C42.1 | PC.ae.C42.2 | PC.ae.C42.3 | |
| ## 1 | 1.77 | 1.59 | 0.629 | 0.316 | 0.192 | 0.277 |
| ## 4 | 1.62 | 1.25 | 0.637 | 0.299 | 0.159 | 0.208 |
| ## 5 | 1.25 | 1.47 | 0.660 | 0.355 | 0.138 | 0.174 |

| | | | | | | | | |
|-----------------------|--------------------|---------------|------------------|---------------|-------------|----------|----------|-------|
| ## 8 | 1.53 | 1.46 | 0.631 | 0.271 | 0.186 | 0.282 | | |
| ## 9 | 1.34 | 1.21 | 0.645 | 0.264 | 0.168 | 0.229 | | |
| ## 10 | 1.30 | 1.18 | 0.670 | 0.294 | 0.186 | 0.222 | | |
| ## PC.ae.C42.4 | PC.ae.C42.5 | PC.ae.C44.3 | PC.ae.C44.4 | PC.ae.C44.5 | PC.ae.C44.6 | | | |
| ## 1 | 0.264 | 0.888 | 0.065 | 0.168 | 0.536 | 0.494 | | |
| ## 4 | 0.392 | 0.863 | 0.069 | 0.237 | 0.517 | 0.611 | | |
| ## 5 | 0.162 | 0.513 | 0.081 | 0.154 | 0.178 | 0.134 | | |
| ## 8 | 0.333 | 0.861 | 0.085 | 0.183 | 0.534 | 0.429 | | |
| ## 9 | 0.319 | 0.925 | 0.069 | 0.184 | 0.671 | 0.460 | | |
| ## 10 | 0.266 | 0.660 | 0.080 | 0.156 | 0.420 | 0.399 | | |
| ## SM..OH..C14.1 | SM..OH..C16.1 | SM..OH..C22.1 | SM..OH..C22.2 | SM..OH..C24.1 | | | | |
| ## 1 | 1.420 | 1.330 | 2.07 | 1.86 | 0.597 | | | |
| ## 4 | 1.720 | 1.480 | 2.97 | 2.84 | 0.682 | | | |
| ## 5 | 0.987 | 1.480 | 1.96 | 1.74 | 0.478 | | | |
| ## 8 | 1.500 | 1.160 | 2.98 | 2.59 | 0.631 | | | |
| ## 9 | 1.230 | 0.984 | 2.21 | 2.07 | 0.507 | | | |
| ## 10 | 1.760 | 1.450 | 2.60 | 2.45 | 0.586 | | | |
| ## SM.C16.0 | SM.C16.1 | SM.C18.0 | SM.C18.1 | SM.C20.2 | SM.C24.0 | SM.C24.1 | SM.C26.0 | |
| ## 1 | 44.9 | 7.99 | 14.5 | 10.40 | 0.290 | 12.20 | 27.3 | 0.147 |
| ## 4 | 52.4 | 8.61 | 17.2 | 11.50 | 0.261 | 11.80 | 27.9 | 0.138 |
| ## 5 | 40.6 | 5.86 | 13.0 | 8.34 | 0.196 | 9.29 | 20.5 | 0.111 |
| ## 8 | 37.9 | 7.92 | 11.9 | 9.59 | 0.199 | 9.36 | 18.8 | 0.117 |
| ## 9 | 37.3 | 6.65 | 12.0 | 8.48 | 0.183 | 10.30 | 23.6 | 0.090 |
| ## 10 | 48.3 | 9.66 | 15.4 | 11.90 | 0.295 | 10.50 | 26.7 | 0.097 |
| ## SM.C26.1 | H1_1 | H1 | Urea_N | L.Arginine_N | L.Leucine_N | EDTAcA_N | | |
| ## 1 | 0.337 | 3356 | 3356 | NA | NA | NA | NA | |
| ## 4 | 0.353 | 2652 | 2652 | 500.8 | 16.0 | 27.1 | 2.5 | |
| ## 5 | 0.283 | 2258 | 2258 | 132.5 | 13.2 | 57.9 | 2.5 | |
| ## 8 | 0.227 | 2464 | 2464 | 182.3 | 49.8 | 63.9 | 0.0 | |
| ## 9 | 0.268 | 2725 | 2725 | 255.8 | 39.0 | 64.6 | 2.7 | |
| ## 10 | 0.305 | 3583 | 3583 | 184.4 | 66.0 | 54.6 | 2.3 | |
| ## X2.Hydroxybutyrate | X3.Hydroxybutyrate | Acetate | Acetoacetate | Acetone | | | | |
| Betaine | | | | | | | | |
| ## 1 | | NA | | NA | NA | NA | NA | |
| NA | | | | | | | | |
| ## 4 | | 12.7 | | 7.2 | 9.8 | 4.8 | 4.0 | |
| 13.9 | | | | | | | | |
| ## 5 | | 35.2 | | 44.7 | 20.2 | 18.9 | 18.9 | |
| 33.9 | | | | | | | | |
| ## 8 | | 21.7 | | 20.9 | 19.5 | 15.4 | 6.6 | |
| 35.3 | | | | | | | | |
| ## 9 | | 17.1 | | 37.9 | 20.0 | 22.0 | 9.3 | |
| 12.0 | | | | | | | | |
| ## 10 | | 21.3 | | 46.2 | 22.5 | 26.2 | 9.4 | |
| 29.5 | | | | | | | | |
| ## Carnitine | Choline | Creatine | Dimethyl.sulfone | Ethanol | Formate | Glucose | | |
| Glycerol | | | | | | | | |
| ## 1 | NA | NA | NA | | NA | NA | NA | |
| NA | | | | | | | | |
| ## 4 | 7.7 | 11.8 | 14.7 | | 1.3 | 6.4 | 14.4 | 629.5 |

| | | | | | | | | |
|-----------------|-------------|-------------|---------|----------|------|------|------|--------|
| 322.0 | | | | | | | | |
| ## 5 | 18.5 | 27.7 | 35.4 | | 5.5 | 13.0 | 40.0 | 1618.0 |
| 271.6 | | | | | | | | |
| ## 8 | 13.0 | 21.5 | 25.9 | | 5.4 | 10.2 | 23.2 | 2120.5 |
| 406.5 | | | | | | | | |
| ## 9 | 14.4 | 15.4 | 25.3 | | 3.5 | 5.1 | 25.0 | 1644.2 |
| 479.9 | | | | | | | | |
| ## 10 | 15.7 | 28.1 | 25.6 | | 2.2 | 4.4 | 29.4 | 2875.0 |
| 722.2 | | | | | | | | |
| ## Hypoxanthine | Isobutyrate | Isopropanol | Lactate | Malonate | | | | |
| ## 1 | NA | NA | NA | NA | | | | |
| ## 4 | 8.6 | 2.5 | 4.4 | 1037.7 | | | | |
| ## 5 | 0.0 | 6.1 | 11.2 | 2199.9 | | | | |
| ## 8 | 5.8 | 4.5 | 4.4 | 1661.4 | | | | |
| ## 9 | 5.6 | 5.9 | 6.7 | 2048.5 | | | | |
| ## 10 | 5.3 | 5.5 | 2.7 | 1801.7 | | | | |
| | | | | | 13.2 | | | |

Output: - The first few rows of the data frame after removing rows with missing Dopamine values.

(d) Replace missing values in the c4-OH-Pro column with the median value.

```
# Replace missing values in c4-OH-Pro with the median
median_c4OHPro <- median(metabolites_no_na_dopamine$c4.OH.Pro, na.rm = TRUE)
metabolites_no_na_dopamine$c4.OH.Pro[is.na(metabolites_no_na_dopamine$c4.OH.P
ro)] <- median_c4OHPro
head(metabolites_no_na_dopamine)

##           Label Phe Pro Ser Thr ADMA alpha.AAA c4.OH.Pro Carnosine
Creatinine
## 1 Alzheimer 72.8 166 170 282 1.15      0.760      0.236      1.270
49.9
## 4 Alzheimer 94.1 129 162 201 1.10      0.795      0.199      0.675
80.1
## 5 Alzheimer 79.8 126 115 199 1.24      1.360      0.199      1.280
60.5
## 8   Healthy 83.6 119 135 268 1.18      0.779      0.215      0.647
30.6
## 9   Healthy 73.7 124 145 307 1.17      0.785      0.186      0.590
39.8
## 10  Healthy 81.5 151 174 269 1.30      0.767      0.185      1.130
44.0
##           DOPA Dopamine Histamine Kynurenone Met.SO Nitro.Tyr PEA Putrescine
## 1 0.265    0.233    0.225     5.21  0.526    0.027  NA  0.068
## 4 0.264    0.234    0.209     5.80  0.389    NA  NA  0.110
## 5 0.271    0.231    0.210     4.46  0.466    NA  NA  0.118
## 8 0.275    0.244    0.214     5.66  0.245    0.002  NA  0.161
## 9 0.259    0.233    0.210     6.36  0.413    NA  NA  0.121
## 10 0.257   0.225    0.210     5.69  0.178    NA  NA  0.183
##           Sarcosine Serotonin Spermidine Spermine t4.OH.Pro Taurine SDMA   C0
C10
```


| | | | | | | | | |
|-------------------|----------------|----------------|----------------|-------------|-------|-------|-------|-------|
| ## 9 | 0.008 | 0.014 | 0.088 | 0.029 | 0.024 | 0.127 | 0.024 | 0.024 |
| 0.025 | | | | | | | | |
| ## 10 | 0.010 | 0.015 | 0.100 | 0.038 | 0.025 | 0.292 | 0.024 | 0.023 |
| NA | | | | | | | | |
| ## C5.1.DC | C6..C4.1.DC. | C5.DC..C6.OH. | C6.1 | C7.DC | C8 | C9 | | |
| lysoPC.a.C14.0 | | | | | | | | |
| ## 1 | 0.020 | 0.022 | 0.014 | 0.018 | 0.011 | 0.062 | 0.016 | |
| 2.23 | | | | | | | | |
| ## 4 | 0.016 | 0.029 | 0.016 | 0.027 | 0.017 | 0.091 | 0.018 | |
| 2.19 | | | | | | | | |
| ## 5 | 0.049 | 0.052 | 0.040 | 0.040 | 0.036 | 0.192 | 0.041 | |
| 1.88 | | | | | | | | |
| ## 8 | 0.017 | 0.036 | 0.011 | NA | 0.009 | 0.062 | 0.011 | |
| 2.13 | | | | | | | | |
| ## 9 | 0.016 | 0.026 | 0.018 | 0.015 | 0.013 | 0.064 | 0.014 | |
| 2.10 | | | | | | | | |
| ## 10 | 0.014 | 0.044 | 0.017 | 0.018 | 0.014 | 0.067 | 0.013 | |
| 2.06 | | | | | | | | |
| ## lysoPC.a.C16.0 | lysoPC.a.C16.1 | lysoPC.a.C17.0 | lysoPC.a.C18.0 | | | | | |
| lysoPC.a.C18.1 | | | | | | | | |
| ## 1 | 37.9 | 2.66 | 0.446 | | 9.00 | | | |
| 8.58 | | | | | | | | |
| ## 4 | 32.8 | 2.39 | 0.323 | | 7.21 | | | |
| 7.22 | | | | | | | | |
| ## 5 | 24.5 | 1.27 | 0.382 | | 6.66 | | | |
| 5.39 | | | | | | | | |
| ## 8 | 33.7 | 3.09 | 0.455 | | 6.96 | | | |
| 7.31 | | | | | | | | |
| ## 9 | 36.0 | 3.46 | 0.435 | | 7.27 | | | |
| 8.11 | | | | | | | | |
| ## 10 | 32.7 | 2.48 | 0.368 | | 6.49 | | | |
| 7.86 | | | | | | | | |
| ## lysoPC.a.C18.2 | lysoPC.a.C20.3 | lysoPC.a.C20.4 | lysoPC.a.C24.0 | | | | | |
| lysoPC.a.C26.0 | | | | | | | | |
| ## 1 | 7.27 | 1.83 | 8.25 | | 0.079 | | | |
| 0.113 | | | | | | | | |
| ## 4 | 7.62 | 1.64 | 6.75 | | 0.066 | | | |
| 0.086 | | | | | | | | |
| ## 5 | 3.60 | 0.97 | 6.26 | | 0.084 | | | |
| 0.118 | | | | | | | | |
| ## 8 | 7.53 | 2.35 | 8.73 | | 0.061 | | | |
| 0.083 | | | | | | | | |
| ## 9 | 6.75 | 2.08 | 7.82 | | 0.068 | | | |
| 0.083 | | | | | | | | |
| ## 10 | 10.10 | 1.86 | 12.40 | | 0.061 | | | |
| 0.086 | | | | | | | | |
| ## lysoPC.a.C26.1 | lysoPC.a.C28.0 | lysoPC.a.C28.1 | PC.aa.C24.0 | PC.aa.C26.0 | | | | |
| ## 1 | 0.053 | 0.108 | 0.072 | 0.082 | 0.438 | | | |
| ## 4 | 0.045 | 0.076 | 0.076 | 0.076 | 0.486 | | | |
| ## 5 | 0.053 | 0.092 | 0.072 | 0.069 | 0.401 | | | |

| | | | | | |
|----------------|-------------|-------------|-------------|-------------|-------------|
| ## 8 | 0.047 | 0.124 | 0.078 | 0.082 | 0.424 |
| ## 9 | 0.037 | 0.096 | 0.061 | 0.064 | 0.430 |
| ## 10 | 0.052 | 0.088 | 0.074 | 0.065 | 0.453 |
| ## PC.aa.C28.1 | PC.aa.C30.0 | PC.aa.C32.0 | PC.aa.C32.1 | PC.aa.C32.2 | PC.aa.C32.3 |
| ## 1 | 0.571 | 2.35 | 11.4 | 9.22 | NA |
| ## 4 | 0.685 | 3.33 | 18.6 | 13.30 | 0.053 |
| ## 5 | 0.513 | 1.78 | 13.8 | 5.03 | NA |
| ## 8 | 0.605 | 2.32 | 12.1 | 11.50 | NA |
| ## 9 | 0.486 | 2.05 | 10.4 | 11.40 | NA |
| ## 10 | 0.578 | 2.28 | 13.8 | 8.10 | NA |
| ## PC.aa.C34.1 | PC.aa.C34.2 | PC.aa.C34.3 | PC.aa.C34.4 | PC.aa.C36.0 | PC.aa.C36.1 |
| ## 1 | 109.0 | 71.0 | 1.430 | 0.200 | 2.38 |
| ## 4 | 106.0 | 93.6 | 1.590 | 0.190 | 2.57 |
| ## 5 | 83.4 | 35.9 | 0.709 | 0.135 | 1.83 |
| ## 8 | 83.6 | 60.6 | 1.580 | 0.251 | 2.16 |
| ## 9 | 89.6 | 55.1 | 1.390 | 0.221 | 1.62 |
| ## 10 | 80.7 | 82.6 | 1.640 | 0.244 | 1.75 |
| ## PC.aa.C36.2 | PC.aa.C36.3 | PC.aa.C36.4 | PC.aa.C36.5 | PC.aa.C36.6 | PC.aa.C38.0 |
| ## 1 | 42.4 | 42.7 | 120.0 | 1.86 | 0.084 |
| ## 4 | 48.8 | 41.2 | 122.0 | 1.76 | 0.070 |
| ## 5 | 28.5 | 21.9 | 98.1 | 1.70 | 0.048 |
| ## 8 | 34.4 | 41.5 | 110.0 | 2.03 | 0.106 |
| ## 9 | 32.0 | 41.8 | 102.0 | 1.77 | 0.081 |
| ## 10 | 38.1 | 31.2 | 138.0 | 1.80 | 0.092 |
| ## PC.aa.C38.3 | PC.aa.C38.4 | PC.aa.C38.5 | PC.aa.C38.6 | PC.aa.C40.1 | PC.aa.C40.2 |
| ## 1 | 32.1 | 95.1 | 16.8 | 41.6 | 0.195 |
| ## 4 | 28.7 | 92.7 | 14.3 | 29.9 | 0.220 |
| ## 5 | 23.3 | 101.0 | 13.8 | 36.2 | 0.165 |
| ## 8 | 31.3 | 81.7 | 14.5 | 42.8 | 0.225 |
| ## 9 | 31.1 | 75.9 | 13.6 | 37.6 | 0.181 |
| ## 10 | 22.2 | 98.2 | 12.7 | 31.5 | 0.189 |
| ## PC.aa.C40.3 | PC.aa.C40.4 | PC.aa.C40.5 | PC.aa.C40.6 | PC.aa.C42.0 | PC.aa.C42.1 |
| ## 1 | 0.491 | 3.48 | 5.66 | 21.8 | 0.364 |
| ## 4 | 0.433 | 3.59 | 5.06 | 14.0 | 0.427 |
| ## 5 | 0.525 | 3.37 | 5.29 | 22.5 | 0.125 |
| ## 8 | 0.384 | 4.02 | 5.49 | 20.6 | 0.365 |
| ## 9 | 0.298 | 2.88 | 4.88 | 17.5 | 0.396 |
| ## 10 | 0.315 | 2.87 | 3.32 | 12.6 | 0.301 |
| ## PC.aa.C42.2 | PC.aa.C42.4 | PC.aa.C42.5 | PC.aa.C42.6 | PC.ae.C30.0 | PC.ae.C30.1 |
| ## 1 | 0.108 | 0.272 | 0.272 | 0.291 | 0.173 |
| ## 4 | 0.119 | 0.268 | 0.267 | 0.254 | 0.223 |
| ## 5 | 0.083 | 0.206 | 0.205 | 0.280 | 0.095 |
| ## 8 | 0.089 | 0.266 | 0.237 | 0.288 | 0.212 |
| ## 9 | 0.086 | 0.215 | 0.206 | 0.223 | 0.179 |
| ## 10 | 0.092 | 0.228 | 0.199 | 0.224 | 0.164 |
| ## PC.ae.C30.2 | PC.ae.C32.1 | PC.ae.C32.2 | PC.ae.C34.0 | PC.ae.C34.1 | PC.ae.C34.2 |
| ## 1 | 0.022 | 1.65 | 0.371 | 0.880 | 3.66 |
| ## 4 | 0.023 | 2.47 | 0.459 | 0.964 | 4.06 |
| ## 5 | 0.023 | 1.72 | 0.316 | 1.060 | 3.28 |
| ## 8 | 0.033 | 1.68 | 0.335 | 1.140 | 3.51 |
| | | | | | 2.28 |

| | | | | | | |
|------------------|---------------|---------------|---------------|---------------|-------------|----------|
| ## 9 | 0.022 | 1.53 | 0.291 | 0.683 | 3.20 | 2.03 |
| ## 10 | 0.024 | 1.70 | 0.341 | 0.800 | 3.56 | 2.52 |
| ## PC.ae.C34.3 | PC.ae.C36.0 | PC.ae.C36.1 | PC.ae.C36.2 | PC.ae.C36.3 | PC.ae.C36.4 | |
| ## 1 | 0.813 | 0.498 | 5.64 | 1.90 | 1.17 | 6.96 |
| ## 4 | 1.020 | 0.552 | 4.75 | 2.01 | 1.35 | 8.36 |
| ## 5 | 0.722 | 0.553 | 5.95 | 1.47 | 0.76 | 4.78 |
| ## 8 | 0.868 | 0.479 | 5.65 | 1.97 | 1.25 | 7.15 |
| ## 9 | 0.663 | 0.384 | 4.79 | 1.54 | 1.12 | 6.43 |
| ## 10 | 0.983 | 0.498 | 4.85 | 1.93 | 1.04 | 7.08 |
| ## PC.ae.C36.5 | PC.ae.C38.0 | PC.ae.C38.1 | PC.ae.C38.2 | PC.ae.C38.3 | PC.ae.C38.4 | |
| ## 1 | 4.79 | 0.474 | 0.287 | 0.538 | 2.66 | 6.33 |
| ## 4 | 5.97 | 0.397 | 0.022 | 0.144 | 1.97 | 5.99 |
| ## 5 | 4.00 | 0.430 | 0.271 | 0.246 | 1.80 | 5.45 |
| ## 8 | 4.04 | 0.550 | 0.078 | 0.559 | 2.90 | 5.73 |
| ## 9 | 3.32 | 0.442 | NA | 0.224 | 2.53 | 5.03 |
| ## 10 | 4.52 | 0.458 | NA | 0.053 | 1.99 | 5.79 |
| ## PC.ae.C38.5 | PC.ae.C38.6 | PC.ae.C40.1 | PC.ae.C40.2 | PC.ae.C40.3 | PC.ae.C40.4 | |
| ## 1 | 5.51 | 1.95 | 0.574 | 0.575 | 0.940 | 1.76 |
| ## 4 | 5.63 | 1.97 | 0.425 | 0.540 | 0.742 | 1.45 |
| ## 5 | 4.34 | 1.51 | 0.430 | 0.432 | 0.632 | 1.10 |
| ## 8 | 4.53 | 1.71 | 0.584 | 0.499 | 0.991 | 1.84 |
| ## 9 | 3.82 | 1.28 | 0.472 | 0.555 | 0.737 | 1.35 |
| ## 10 | 4.84 | 1.58 | 0.670 | 0.572 | 0.757 | 1.16 |
| ## PC.ae.C40.5 | PC.ae.C40.6 | PC.ae.C42.0 | PC.ae.C42.1 | PC.ae.C42.2 | PC.ae.C42.3 | |
| ## 1 | 1.77 | 1.59 | 0.629 | 0.316 | 0.192 | 0.277 |
| ## 4 | 1.62 | 1.25 | 0.637 | 0.299 | 0.159 | 0.208 |
| ## 5 | 1.25 | 1.47 | 0.660 | 0.355 | 0.138 | 0.174 |
| ## 8 | 1.53 | 1.46 | 0.631 | 0.271 | 0.186 | 0.282 |
| ## 9 | 1.34 | 1.21 | 0.645 | 0.264 | 0.168 | 0.229 |
| ## 10 | 1.30 | 1.18 | 0.670 | 0.294 | 0.186 | 0.222 |
| ## PC.ae.C42.4 | PC.ae.C42.5 | PC.ae.C44.3 | PC.ae.C44.4 | PC.ae.C44.5 | PC.ae.C44.6 | |
| ## 1 | 0.264 | 0.888 | 0.065 | 0.168 | 0.536 | 0.494 |
| ## 4 | 0.392 | 0.863 | 0.069 | 0.237 | 0.517 | 0.611 |
| ## 5 | 0.162 | 0.513 | 0.081 | 0.154 | 0.178 | 0.134 |
| ## 8 | 0.333 | 0.861 | 0.085 | 0.183 | 0.534 | 0.429 |
| ## 9 | 0.319 | 0.925 | 0.069 | 0.184 | 0.671 | 0.460 |
| ## 10 | 0.266 | 0.660 | 0.080 | 0.156 | 0.420 | 0.399 |
| ## SM..OH..C14.1 | SM..OH..C16.1 | SM..OH..C22.1 | SM..OH..C22.2 | SM..OH..C24.1 | | |
| ## 1 | 1.420 | 1.330 | 2.07 | 1.86 | 0.597 | |
| ## 4 | 1.720 | 1.480 | 2.97 | 2.84 | 0.682 | |
| ## 5 | 0.987 | 1.480 | 1.96 | 1.74 | 0.478 | |
| ## 8 | 1.500 | 1.160 | 2.98 | 2.59 | 0.631 | |
| ## 9 | 1.230 | 0.984 | 2.21 | 2.07 | 0.507 | |
| ## 10 | 1.760 | 1.450 | 2.60 | 2.45 | 0.586 | |
| ## SM.C16.0 | SM.C16.1 | SM.C18.0 | SM.C18.1 | SM.C20.2 | SM.C24.0 | SM.C24.1 |
| ## 1 | 44.9 | 7.99 | 14.5 | 10.40 | 0.290 | 12.20 |
| ## 4 | 52.4 | 8.61 | 17.2 | 11.50 | 0.261 | 11.80 |
| ## 5 | 40.6 | 5.86 | 13.0 | 8.34 | 0.196 | 9.29 |
| ## 8 | 37.9 | 7.92 | 11.9 | 9.59 | 0.199 | 9.36 |
| ## 9 | 37.3 | 6.65 | 12.0 | 8.48 | 0.183 | 10.30 |
| | | | | | | 23.6 |
| | | | | | | 0.090 |

| | | | | | | | | |
|----------|--------------------|--------------------|-------------|------------------|--------------|-------------|----------|--------|
| ## 10 | 48.3 | 9.66 | 15.4 | 11.90 | 0.295 | 10.50 | 26.7 | 0.097 |
| ## | SM.C26.1 | H1_1 | H1 | Urea_N | L.Arginine_N | L.Leucine_N | EDTACA_N | |
| ## 1 | 0.337 | 3356 | 3356 | NA | NA | NA | NA | |
| ## 4 | 0.353 | 2652 | 2652 | 500.8 | 16.0 | 27.1 | 2.5 | |
| ## 5 | 0.283 | 2258 | 2258 | 132.5 | 13.2 | 57.9 | 2.5 | |
| ## 8 | 0.227 | 2464 | 2464 | 182.3 | 49.8 | 63.9 | 0.0 | |
| ## 9 | 0.268 | 2725 | 2725 | 255.8 | 39.0 | 64.6 | 2.7 | |
| ## 10 | 0.305 | 3583 | 3583 | 184.4 | 66.0 | 54.6 | 2.3 | |
| ## | X2.Hydroxybutyrate | X3.Hydroxybutyrate | Acetate | Acetoacetate | Acetone | | | |
| Betaine | | | | | | | | |
| ## 1 | | NA | | NA | NA | NA | NA | |
| NA | | | | | | | | |
| ## 4 | | 12.7 | | 7.2 | 9.8 | 4.8 | 4.0 | |
| 13.9 | | | | | | | | |
| ## 5 | | 35.2 | | 44.7 | 20.2 | 18.9 | 18.9 | |
| 33.9 | | | | | | | | |
| ## 8 | | 21.7 | | 20.9 | 19.5 | 15.4 | 6.6 | |
| 35.3 | | | | | | | | |
| ## 9 | | 17.1 | | 37.9 | 20.0 | 22.0 | 9.3 | |
| 12.0 | | | | | | | | |
| ## 10 | | 21.3 | | 46.2 | 22.5 | 26.2 | 9.4 | |
| 29.5 | | | | | | | | |
| ## | Carnitine | Choline | Creatine | Dimethyl.sulfone | Ethanol | Formate | Glucose | |
| Glycerol | | | | | | | | |
| ## 1 | NA | NA | NA | | NA | NA | NA | |
| NA | | | | | | | | |
| ## 4 | 7.7 | 11.8 | 14.7 | | 1.3 | 6.4 | 14.4 | 629.5 |
| 322.0 | | | | | | | | |
| ## 5 | 18.5 | 27.7 | 35.4 | | 5.5 | 13.0 | 40.0 | 1618.0 |
| 271.6 | | | | | | | | |
| ## 8 | 13.0 | 21.5 | 25.9 | | 5.4 | 10.2 | 23.2 | 2120.5 |
| 406.5 | | | | | | | | |
| ## 9 | 14.4 | 15.4 | 25.3 | | 3.5 | 5.1 | 25.0 | 1644.2 |
| 479.9 | | | | | | | | |
| ## 10 | 15.7 | 28.1 | 25.6 | | 2.2 | 4.4 | 29.4 | 2875.0 |
| 722.2 | | | | | | | | |
| ## | Hypoxanthine | Isobutyrate | Isopropanol | Lactate | Malonate | | | |
| ## 1 | NA | NA | NA | NA | NA | | | |
| ## 4 | 8.6 | 2.5 | 4.4 | 1037.7 | 7.6 | | | |
| ## 5 | 0.0 | 6.1 | 11.2 | 2199.9 | 11.7 | | | |
| ## 8 | 5.8 | 4.5 | 4.4 | 1661.4 | 11.0 | | | |
| ## 9 | 5.6 | 5.9 | 6.7 | 2048.5 | 8.0 | | | |
| ## 10 | 5.3 | 5.5 | 2.7 | 1801.7 | 13.2 | | | |

Output: - The first few rows after replacing missing values in the c4-OH-Pro column with the median.

(e) (Optional) Drop columns with more than 25% missing values.

Drop columns with more than 25% missing values

threshold <- 0.25 * nrow(metabolites_data)

```

metabolites_clean <- metabolites_data[, colSums(is.na(metabolites_data)) <=
threshold]
head(metabolites_clean)

##      Label Phe Pro Ser Thr ADMA alpha.AAA Carnosine Creatinine DOPA
## 1 Alzheimer 72.8 166 170 282 1.15     0.760    1.270     49.9 0.265
## 2 Alzheimer 93.4 138 142 217 1.05     0.929    1.350     48.8 0.252
## 3 Alzheimer 68.6 161 158 208 1.00     0.620    0.998     30.4 0.268
## 4 Alzheimer 94.1 129 162 201 1.10     0.795    0.675     80.1 0.264
## 5 Alzheimer 79.8 126 115 199 1.24     1.360    1.280     60.5 0.271
## 6 Alzheimer 82.5 167 173 333 1.35     1.150    1.010     24.0 0.275
##      Histamine Kynurenone Met.SO Putrescine Sarcosine Serotonin Spermidine
## 1      0.225      5.21  0.526     0.068     17.8     0.147     0.188
## 2      0.211      5.44  0.387     0.087     20.2     0.231     0.233
## 3      0.217      5.20  0.651     0.260     14.4     0.196     0.384
## 4      0.209      5.80  0.389     0.110     18.7     0.255     0.353
## 5      0.210      4.46  0.466     0.118     22.5     0.390     0.473
## 6      0.212      7.01  0.417     0.262     30.8     0.140     0.856
##      t4.OH.Pro Taurine SDMA   C0   C10 C10.1 C10.2   C12 C12.DC C12.1   C14
C14.1
## 1      24.0      125 1.13 18.2 0.059 0.312 0.038 0.030 0.042 0.290 0.023
0.019
## 2      29.3      120 1.65 17.0 0.051 0.288 0.039 0.038 0.038 0.265 0.026
0.017
## 3      20.9      139 1.57 12.6 0.083 0.357 0.054 0.032 0.048 0.302 0.021
0.031
## 4      23.1      159 1.34 23.5 0.071 0.317 0.040 0.045 0.048 0.275 0.026
0.028
## 5      26.9      149 1.24 13.6 0.139 0.472 0.074 0.056 0.079 0.394 0.034
0.043
## 6      26.0      379 1.44 26.7 0.058 0.238 0.042 0.039 0.035 0.196 0.029
0.023
##      C14.1.OH C14.2 C14.2.OH   C16 C16.OH C16.1 C16.1.OH C16.2 C16.2.OH   C18
## 1      0.008 0.008     0.006 0.046 0.008 0.009     0.007 0.005     0.013 0.013
## 2      0.008 0.009     0.009 0.070 0.009 0.013     0.006 0.006     0.012 0.014
## 3      0.010 0.010     0.009 0.076 0.011 0.019     0.010 0.005     0.013 0.016
## 4      0.010 0.013     0.011 0.074 0.011 0.015     0.008 0.006     0.009 0.020
## 5      0.016 0.025     0.017 0.062     NA 0.024     0.014 0.012     0.025 0.031
## 6      0.009 0.010     0.007 0.081 0.006 0.012     0.005 0.007     0.015 0.017
##      C18.1 C18.1.OH C18.2   C2   C3 C3.OH C3.1   C4 C3.DC..C4.OH.   C4.1
C5
## 1 0.024      0.003 0.016 1.97 0.354 0.008 0.015 0.082           0.045 0.025
0.094
## 2 0.025      0.003 0.028 1.95 0.184 0.009 0.013 0.108           0.080 0.025
0.077
## 3 0.025      NA 0.018 1.70 0.371     NA 0.012 0.057           0.035 0.039
0.096
## 4 0.035      0.004 0.033 2.10 0.278 0.010 0.017 0.110           0.077 0.031
0.145
## 5 0.034      0.012 0.017 5.62 0.436 0.029 0.035 0.106           0.099 0.069

```

```

0.141
## 6 0.035      0.004 0.029 3.49 0.461 0.008 0.014 0.123          0.068 0.026
0.090
##   C5.M.DC C5.OH..C3.DC.M.  C5.1 C5.1.DC C6..C4.1.DC. C5.DC..C6.OH.  C6.1
C7.DC
## 1  0.023      0.026 0.030 0.020          0.022          0.014 0.018
0.011
## 2  0.032      0.026 0.024 0.021          0.030          0.018 0.015
0.010
## 3  0.045      0.024 0.037 0.018          0.022          0.029 0.031
0.021
## 4  0.034      0.041 0.035 0.016          0.029          0.016 0.027
0.017
## 5  0.094      0.058 0.073 0.049          0.052          0.040 0.040
0.036
## 6  0.019      0.037 0.022 0.016          0.063          0.016 0.019
0.014
##   C8   C9 lysoPC.a.C14.0 lysoPC.a.C16.0 lysoPC.a.C16.1 lysoPC.a.C17.0
## 1 0.062 0.016      2.23      37.9      2.66      0.446
## 2 0.058 0.014      1.97      22.1      1.31      0.270
## 3 0.090 0.017      2.12      33.7      2.53      0.399
## 4 0.091 0.018      2.19      32.8      2.39      0.323
## 5 0.192 0.041      1.88      24.5      1.27      0.382
## 6 0.073 0.014      2.11      29.1      2.09      0.348
##   lysoPC.a.C18.0 lysoPC.a.C18.1 lysoPC.a.C18.2 lysoPC.a.C20.3
lysoPC.a.C20.4
## 1      9.00      8.58      7.27      1.830
8.25
## 2      5.35      3.94      4.42      0.958
4.60
## 3      7.51      7.73      8.02      2.050
9.84
## 4      7.21      7.22      7.62      1.640
6.75
## 5      6.66      5.39      3.60      0.970
6.26
## 6      5.84      6.30      8.10      1.970
7.04
##   lysoPC.a.C24.0 lysoPC.a.C26.0 lysoPC.a.C26.1 lysoPC.a.C28.0
lysoPC.a.C28.1
## 1      0.079      0.113      0.053      0.108
0.072
## 2      0.059      0.066      0.042      0.076
0.058
## 3      0.075      0.126      0.049      0.078
0.092
## 4      0.066      0.086      0.045      0.076
0.076
## 5      0.084      0.118      0.053      0.092
0.072

```

| | | | | | |
|----------------|-------------|-------------|-------------|-------------|-------------|
| ## 6 | 0.083 | 0.112 | 0.050 | 0.099 | |
| 0.083 | | | | | |
| ## PC.aa.C24.0 | PC.aa.C26.0 | PC.aa.C28.1 | PC.aa.C30.0 | PC.aa.C32.0 | PC.aa.C32.1 |
| ## 1 | 0.082 | 0.438 | 0.571 | 2.35 | 11.4 |
| ## 2 | 0.065 | 0.409 | 0.521 | 1.99 | 12.7 |
| ## 3 | 0.099 | 0.458 | 0.605 | 2.69 | 16.6 |
| ## 4 | 0.076 | 0.486 | 0.685 | 3.33 | 18.6 |
| ## 5 | 0.069 | 0.401 | 0.513 | 1.78 | 13.8 |
| ## 6 | 0.073 | 0.450 | 0.620 | 2.61 | 14.7 |
| ## PC.aa.C32.3 | PC.aa.C34.1 | PC.aa.C34.2 | PC.aa.C34.3 | PC.aa.C34.4 | PC.aa.C36.0 |
| ## 1 | 0.092 | 109.0 | 71.0 | 1.430 | 0.200 |
| ## 2 | 0.067 | 64.2 | 60.5 | 0.879 | 0.127 |
| ## 3 | 0.105 | 108.0 | 83.1 | 1.930 | 0.210 |
| ## 4 | 0.079 | 106.0 | 93.6 | 1.590 | 0.190 |
| ## 5 | 0.102 | 83.4 | 35.9 | 0.709 | 0.135 |
| ## 6 | 0.107 | 90.2 | 85.6 | 1.790 | 0.213 |
| ## PC.aa.C36.1 | PC.aa.C36.2 | PC.aa.C36.3 | PC.aa.C36.4 | PC.aa.C36.5 | PC.aa.C36.6 |
| ## 1 | 21.7 | 42.4 | 42.7 | 120.0 | 1.86 |
| ## 2 | 14.3 | 35.6 | 24.3 | 83.7 | 1.05 |
| ## 3 | 19.9 | 44.9 | 43.9 | 146.0 | 2.09 |
| ## 4 | 20.9 | 48.8 | 41.2 | 122.0 | 1.76 |
| ## 5 | 20.5 | 28.5 | 21.9 | 98.1 | 1.70 |
| ## 6 | 15.5 | 43.2 | 46.0 | 114.0 | 3.47 |
| ## PC.aa.C38.0 | PC.aa.C38.3 | PC.aa.C38.4 | PC.aa.C38.5 | PC.aa.C38.6 | PC.aa.C40.1 |
| ## 1 | 1.230 | 32.1 | 95.1 | 16.80 | 41.6 |
| ## 2 | 0.946 | 21.9 | 78.9 | 9.91 | 25.1 |
| ## 3 | 1.210 | 34.5 | 107.0 | 17.50 | 36.6 |
| ## 4 | 1.160 | 28.7 | 92.7 | 14.30 | 29.9 |
| ## 5 | 1.100 | 23.3 | 101.0 | 13.80 | 36.2 |
| ## 6 | 1.390 | 28.9 | 78.0 | 13.10 | 48.4 |
| ## PC.aa.C40.2 | PC.aa.C40.3 | PC.aa.C40.4 | PC.aa.C40.5 | PC.aa.C40.6 | PC.aa.C42.0 |
| ## 1 | 0.074 | 0.491 | 3.48 | 5.66 | 21.8 |
| ## 2 | 0.057 | 0.358 | 3.39 | 4.08 | 14.2 |
| ## 3 | 0.118 | 0.395 | 3.56 | 5.34 | 16.7 |
| ## 4 | 0.097 | 0.433 | 3.59 | 5.06 | 14.0 |
| ## 5 | 0.044 | 0.525 | 3.37 | 5.29 | 22.5 |
| ## 6 | 0.120 | 0.346 | 2.63 | 3.25 | 18.9 |
| ## PC.aa.C42.1 | PC.aa.C42.2 | PC.aa.C42.4 | PC.aa.C42.5 | PC.aa.C42.6 | PC.ae.C30.0 |
| ## 1 | 0.226 | 0.108 | 0.272 | 0.272 | 0.291 |
| ## 2 | 0.216 | 0.109 | 0.336 | 0.317 | 0.248 |
| ## 3 | 0.281 | 0.118 | 0.300 | 0.206 | 0.267 |
| ## 4 | 0.223 | 0.119 | 0.268 | 0.267 | 0.254 |
| ## 5 | 0.095 | 0.083 | 0.206 | 0.205 | 0.280 |
| ## 6 | 0.233 | 0.135 | 0.228 | 0.254 | 0.271 |
| ## PC.ae.C30.1 | PC.ae.C30.2 | PC.ae.C32.1 | PC.ae.C32.2 | PC.ae.C34.0 | PC.ae.C34.1 |
| ## 1 | 0.027 | 0.022 | 1.65 | 0.371 | 0.880 |
| ## 2 | 0.024 | 0.020 | 2.01 | 0.360 | 0.763 |
| ## 3 | 0.046 | 0.030 | 2.40 | 0.477 | 0.938 |
| ## 4 | 0.049 | 0.023 | 2.47 | 0.459 | 0.964 |
| ## 5 | 0.082 | 0.023 | 1.72 | 0.316 | 1.060 |
| | | | | | 3.28 |

| | | | | | | |
|------------------|---------------|---------------|---------------|-------------|-------------|-------|
| ## 6 | 0.039 | 0.029 | 2.01 | 0.397 | 0.920 | 3.26 |
| ## PC.ae.C34.2 | PC.ae.C34.3 | PC.ae.C36.0 | PC.ae.C36.1 | PC.ae.C36.2 | PC.ae.C36.3 | |
| ## 1 | 2.48 | 0.813 | 0.498 | 5.64 | 1.90 | 1.170 |
| ## 2 | 2.32 | 0.905 | 0.398 | 3.89 | 1.54 | 0.873 |
| ## 3 | 2.95 | 1.030 | 0.554 | 5.95 | 2.29 | 1.240 |
| ## 4 | 3.09 | 1.020 | 0.552 | 4.75 | 2.01 | 1.350 |
| ## 5 | 1.70 | 0.722 | 0.553 | 5.95 | 1.47 | 0.760 |
| ## 6 | 2.58 | 1.000 | 0.443 | 4.95 | 2.05 | 1.170 |
| ## PC.ae.C36.4 | PC.ae.C36.5 | PC.ae.C38.0 | PC.ae.C38.3 | PC.ae.C38.4 | PC.ae.C38.5 | |
| ## 1 | 6.96 | 4.79 | 0.474 | 2.66 | 6.33 | 5.51 |
| ## 2 | 6.40 | 5.36 | 0.325 | 1.80 | 5.37 | 4.49 |
| ## 3 | 9.05 | 6.63 | 0.478 | 2.87 | 7.06 | 5.64 |
| ## 4 | 8.36 | 5.97 | 0.397 | 1.97 | 5.99 | 5.63 |
| ## 5 | 4.78 | 4.00 | 0.430 | 1.80 | 5.45 | 4.34 |
| ## 6 | 7.04 | 4.47 | 0.590 | 2.46 | 5.55 | 4.60 |
| ## PC.ae.C38.6 | PC.ae.C40.1 | PC.ae.C40.2 | PC.ae.C40.3 | PC.ae.C40.4 | PC.ae.C40.5 | |
| ## 1 | 1.95 | 0.574 | 0.575 | 0.940 | 1.76 | 1.77 |
| ## 2 | 1.63 | 0.281 | 0.491 | 0.702 | 1.43 | 1.55 |
| ## 3 | 1.98 | 0.759 | 0.654 | 0.817 | 1.51 | 1.64 |
| ## 4 | 1.97 | 0.425 | 0.540 | 0.742 | 1.45 | 1.62 |
| ## 5 | 1.51 | 0.430 | 0.432 | 0.632 | 1.10 | 1.25 |
| ## 6 | 1.80 | 0.481 | 0.598 | 0.826 | 1.25 | 1.38 |
| ## PC.ae.C40.6 | PC.ae.C42.0 | PC.ae.C42.1 | PC.ae.C42.2 | PC.ae.C42.3 | PC.ae.C42.4 | |
| ## 1 | 1.59 | 0.629 | 0.316 | 0.192 | 0.277 | 0.264 |
| ## 2 | 1.20 | 0.616 | 0.260 | 0.157 | 0.200 | 0.311 |
| ## 3 | 1.49 | 0.686 | 0.356 | 0.241 | 0.288 | 0.319 |
| ## 4 | 1.25 | 0.637 | 0.299 | 0.159 | 0.208 | 0.392 |
| ## 5 | 1.47 | 0.660 | 0.355 | 0.138 | 0.174 | 0.162 |
| ## 6 | 1.61 | 0.669 | 0.265 | 0.195 | 0.253 | 0.316 |
| ## PC.ae.C42.5 | PC.ae.C44.3 | PC.ae.C44.4 | PC.ae.C44.5 | PC.ae.C44.6 | | |
| SM..OH..C14.1 | | | | | | |
| ## 1 | 0.888 | 0.065 | 0.168 | 0.536 | 0.494 | |
| 1.420 | | | | | | |
| ## 2 | 0.840 | 0.071 | 0.220 | 0.470 | 0.515 | |
| 1.390 | | | | | | |
| ## 3 | 0.957 | 0.065 | 0.228 | 0.565 | 0.603 | |
| 1.840 | | | | | | |
| ## 4 | 0.863 | 0.069 | 0.237 | 0.517 | 0.611 | |
| 1.720 | | | | | | |
| ## 5 | 0.513 | 0.081 | 0.154 | 0.178 | 0.134 | |
| 0.987 | | | | | | |
| ## 6 | 0.814 | 0.085 | 0.232 | 0.554 | 0.539 | |
| 1.320 | | | | | | |
| ## SM..OH..C16.1 | SM..OH..C22.1 | SM..OH..C22.2 | SM..OH..C24.1 | SM.C16.0 | | |
| SM.C16.1 | | | | | | |
| ## 1 | 1.33 | 2.07 | 1.86 | 0.597 | 44.9 | |
| 7.99 | | | | | | |
| ## 2 | 1.25 | 2.47 | 2.20 | 0.640 | 42.1 | |
| 6.88 | | | | | | |
| ## 3 | 1.58 | 2.69 | 2.63 | 0.665 | 44.8 | |

| | | | | | | |
|--|----------|----------|----------|----------|----------|-----------------------|
| 8.91 | | | | | | |
| ## 4 | 1.48 | 2.97 | 2.84 | 0.682 | 52.4 | |
| 8.61 | | | | | | |
| ## 5 | 1.48 | 1.96 | 1.74 | 0.478 | 40.6 | |
| 5.86 | | | | | | |
| ## 6 | 1.12 | 2.51 | 2.16 | 0.640 | 42.6 | |
| 8.49 | | | | | | |
| ## SM.C18.0 | SM.C18.1 | SM.C20.2 | SM.C24.0 | SM.C24.1 | SM.C26.0 | SM.C26.1 H1_1 H1 |
| ## 1 | 14.5 | 10.40 | 0.290 | 12.20 | 27.3 | 0.147 0.337 3356 3356 |
| ## 2 | 12.7 | 8.52 | 0.211 | 10.40 | 25.6 | 0.130 0.317 2509 2509 |
| ## 3 | 14.6 | 11.60 | 0.304 | 11.50 | 28.8 | 0.163 0.364 2661 2661 |
| ## 4 | 17.2 | 11.50 | 0.261 | 11.80 | 27.9 | 0.138 0.353 2652 2652 |
| ## 5 | 13.0 | 8.34 | 0.196 | 9.29 | 20.5 | 0.111 0.283 2258 2258 |
| ## 6 | 13.0 | 10.60 | 0.270 | 9.58 | 23.7 | 0.135 0.316 3031 3031 |
| ## Urea_N L.Arginine_N L.Leucine_N EDTAca_N X2.Hydroxybutyrate | | | | | | |
| ## 1 | NA | NA | NA | NA | NA | NA |
| ## 2 | 201.9 | 22.5 | 35.3 | 2.0 | 12.40 | |
| ## 3 | 193.3 | 21.0 | 25.4 | 1.8 | 11.33 | |
| ## 4 | 500.8 | 16.0 | 27.1 | 2.5 | 12.70 | |
| ## 5 | 132.5 | 13.2 | 57.9 | 2.5 | 35.20 | |
| ## 6 | 193.3 | 32.2 | 26.5 | 0.0 | 17.20 | |
| ## X3.Hydroxybutyrate Acetate Acetoacetate Acetone Betaine Carnitine | | | | | | |
| Choline | | | | | | |
| ## 1 | NA | NA | NA | NA | NA | NA |
| NA | | | | | | |
| ## 2 | 8.5 | 13.2 | | 5.7 | 5.1 | 22.0 |
| 14.2 | | | | | | 8.7 |
| ## 3 | 11.7 | 5.8 | | 9.3 | 5.6 | 19.1 |
| 14.5 | | | | | | 15.3 |
| ## 4 | 7.2 | 9.8 | | 4.8 | 4.0 | 13.9 |
| 11.8 | | | | | | 7.7 |
| ## 5 | 44.7 | 20.2 | | 18.9 | 18.9 | 33.9 |
| 27.7 | | | | | | 18.5 |
| ## 6 | 16.0 | 23.6 | | 7.8 | 5.5 | 16.9 |
| 25.9 | | | | | | 16.7 |
| ## Creatine Dimethyl.sulfone Ethanol Formate Glucose Glycerol Hypoxanthine | | | | | | |
| ## 1 | NA | NA | NA | NA | NA | NA |
| ## 2 | 14.5 | 4.7 | 16.6 | 24.6 | 1489.7 | 324.6 |
| ## 3 | 17.8 | 2.1 | 8.1 | 27.4 | 1343.9 | 201.3 |
| ## 4 | 14.7 | 1.3 | 6.4 | 14.4 | 629.5 | 322.0 |
| ## 5 | 35.4 | 5.5 | 13.0 | 40.0 | 1618.0 | 271.6 |
| ## 6 | 18.6 | 3.4 | 5.0 | 35.5 | 1791.8 | 274.2 |
| ## Isobutyrate Isopropanol Lactate Malonate | | | | | | |
| ## 1 | NA | NA | NA | NA | | |
| ## 2 | 3.6 | 1.9 | 1171.6 | 10.4 | | |
| ## 3 | 2.5 | 2.5 | 1938.1 | 13.1 | | |
| ## 4 | 2.5 | 4.4 | 1037.7 | 7.6 | | |
| ## 5 | 6.1 | 11.2 | 2199.9 | 11.7 | | |
| ## 6 | 2.3 | 2.4 | 1486.7 | 11.8 | | |

Output: - The first few rows after dropping columns with more than 25% missing values.
