

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
## # i 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
0			
##	Thr	ADMA	alpha.AAA
c4.OH.Pro			
##	0	0	0
20			
##	Carnosine	Creatinine	DOPA
Dopamine			
##	1	0	0
20			
##	Histamine	Kynurenine	Met.S0
Nitro.Tyr			
##	0	0	1
62			
##	PEA	Putrescine	Sarcosine
Serotonin			
##	69	0	0
0			
##	Spermidine	Spermine	t4.OH.Pro
Taurine			
##	0	60	0
2			
##	SDMA	C0	C10
C10.1			
##	0	0	0
0			
##	C10.2	C12	C12.DC
C12.1			
##	0	0	1
0			
##	C14	C14.1	C14.1.OH
C14.2			
##	0	0	1
0			
##	C14.2.OH	C16	C16.OH
C16.1			
##	2	0	1
0			
##	C16.1.OH	C16.2	C16.2.OH
C18			
##	2	2	1
0			
##	C18.1	C18.1.OH	C18.2
C2			
##	0	7	0
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.aa.C30.0		
##	0	0	0
0			
##	PC.aa.C30.1	PC.aa.C30.2	PC.aa.C32.1
	PC.aa.C32.2		
##	10	0	0
0			
##	PC.aa.C34.0	PC.aa.C34.1	PC.aa.C34.2
	PC.aa.C34.3		
##	0	0	0
0			
##	PC.aa.C36.0	PC.aa.C36.1	PC.aa.C36.2
	PC.aa.C36.3		
##	0	0	0
0			
##	PC.aa.C36.4	PC.aa.C36.5	PC.aa.C38.0
	PC.aa.C38.1		
##	0	0	0
52			
##	PC.aa.C38.2	PC.aa.C38.3	PC.aa.C38.4
	PC.aa.C38.5		
##	19	0	0
0			
##	PC.aa.C38.6	PC.aa.C40.1	PC.aa.C40.2
	PC.aa.C40.3		
##	0	0	0
0			
##	PC.aa.C40.4	PC.aa.C40.5	PC.aa.C40.6
	PC.aa.C42.0		
##	0	0	0
0			
##	PC.aa.C42.1	PC.aa.C42.2	PC.aa.C42.3
	PC.aa.C42.4		
##	0	1	0
0			
##	PC.aa.C42.5	PC.aa.C44.3	PC.aa.C44.4
	PC.aa.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
##      EDTAc_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)

```

##	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	NA	0.675
------	-----------	------	-----	-----	-----	------	-------	----	-------

80.1

## 5	Alzheimer	79.8	126	115	199	1.24	1.360	NA	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	Kynurenine	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

## 1	17.8	0.147	0.188	NA	24.0	125	1.13	18.2
------	------	-------	-------	----	------	-----	------	------

0.059

## 4	18.7	0.255	0.353	NA	23.1	159	1.34	23.5
------	------	-------	-------	----	------	-----	------	------

0.071

## 5	22.5	0.390	0.473	NA	26.9	149	1.24	13.6
------	------	-------	-------	----	------	-----	------	------

0.139

## 8	23.3	0.215	0.276	NA	10.7	133	1.04	13.3
------	------	-------	-------	----	------	-----	------	------

0.051

## 9	22.1	0.166	0.327	NA	16.0	215	1.24	15.8
------	------	-------	-------	----	------	-----	------	------

0.061

## 10	18.5	0.320	0.236	NA	27.9	140	1.84	15.6
-------	------	-------	-------	----	------	-----	------	------

0.068

##	C10.1	C10.2	C12	C12.DC	C12.1	C14	C14.1	C14.1.OH	C14.2	C14.2.OH
----	-------	-------	-----	--------	-------	-----	-------	----------	-------	----------

C16

## 1	0.312	0.038	0.030	0.042	0.290	0.023	0.019	0.008	0.008	0.006
------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.046

## 4	0.317	0.040	0.045	0.048	0.275	0.026	0.028	0.010	0.013	0.011
------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.074

## 5	0.472	0.074	0.056	0.079	0.394	0.034	0.043	0.016	0.025	0.017
------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.062

## 8	0.217	0.030	0.041	0.035	0.174	0.024	0.017	0.007	0.006	0.007
------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.060

## 9	0.258	0.036	0.037	0.038	0.228	0.022	0.018	0.007	0.007	0.007
------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.054

## 10	0.319	0.041	0.039	0.037	0.284	0.026	0.022	0.008	0.013	0.007
-------	-------	-------	-------	-------	-------	-------	-------	-------	-------	-------

0.076

##	C16.OH	C16.1	C16.1.OH	C16.2	C16.2.OH	C18	C18.1	C18.1.OH	C18.2	C2
----	--------	-------	----------	-------	----------	-----	-------	----------	-------	----

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 0.092
## 4	0.685	3.33	18.6	13.30	0.053 0.079
## 5	0.513	1.78	13.8	5.03	NA 0.102
## 8	0.605	2.32	12.1	11.50	NA 0.093
## 9	0.486	2.05	10.4	11.40	NA 0.082
## 10	0.578	2.28	13.8	8.10	NA 0.111
## 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 21.7
## 4	106.0	93.6	1.590	0.190	2.57 20.9
## 5	83.4	35.9	0.709	0.135	1.83 20.5
## 8	83.6	60.6	1.580	0.251	2.16 18.4
## 9	89.6	55.1	1.390	0.221	1.62 18.3
## 10	80.7	82.6	1.640	0.244	1.75 14.6
## 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 1.230
## 4	48.8	41.2	122.0	1.76	0.070 1.160
## 5	28.5	21.9	98.1	1.70	0.048 1.100
## 8	34.4	41.5	110.0	2.03	0.106 1.150
## 9	32.0	41.8	102.0	1.77	0.081 0.902
## 10	38.1	31.2	138.0	1.80	0.092 0.904
## 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 0.074
## 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.aa.C30.0	PC.aa.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.aa.C30.2	PC.aa.C32.1	PC.aa.C32.2	PC.aa.C34.0	PC.aa.C34.1	PC.aa.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.aa.C34.3	PC.aa.C36.0	PC.aa.C36.1	PC.aa.C36.2	PC.aa.C36.3	PC.aa.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.aa.C36.5	PC.aa.C38.0	PC.aa.C38.1	PC.aa.C38.2	PC.aa.C38.3	PC.aa.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.aa.C38.5	PC.aa.C38.6	PC.aa.C40.1	PC.aa.C40.2	PC.aa.C40.3	PC.aa.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.aa.C40.5	PC.aa.C40.6	PC.aa.C42.0	PC.aa.C42.1	PC.aa.C42.2	PC.aa.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	EDTAc_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
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              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 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.Pro)] <- 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 Kynurenine Met.S0 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

```

## 1 0.059	17.8	0.147	0.188	NA	24.0	125	1.13	18.2		
## 4 0.071	18.7	0.255	0.353	NA	23.1	159	1.34	23.5		
## 5 0.139	22.5	0.390	0.473	NA	26.9	149	1.24	13.6		
## 8 0.051	23.3	0.215	0.276	NA	10.7	133	1.04	13.3		
## 9 0.061	22.1	0.166	0.327	NA	16.0	215	1.24	15.8		
## 10 0.068	18.5	0.320	0.236	NA	27.9	140	1.84	15.6		
## C16	C10.1	C10.2	C12	C12.DC	C12.1	C14	C14.1	C14.1.OH	C14.2	C14.2.OH
## 1 0.046	0.312	0.038	0.030	0.042	0.290	0.023	0.019	0.008	0.008	0.006
## 4 0.074	0.317	0.040	0.045	0.048	0.275	0.026	0.028	0.010	0.013	0.011
## 5 0.062	0.472	0.074	0.056	0.079	0.394	0.034	0.043	0.016	0.025	0.017
## 8 0.060	0.217	0.030	0.041	0.035	0.174	0.024	0.017	0.007	0.006	0.007
## 9 0.054	0.258	0.036	0.037	0.038	0.228	0.022	0.018	0.007	0.007	0.007
## 10 0.076	0.319	0.041	0.039	0.037	0.284	0.026	0.022	0.008	0.013	0.007
## C3	C16.OH	C16.1	C16.1.OH	C16.2	C16.2.OH	C18	C18.1	C18.1.OH	C18.2	C2
## 1 0.354	0.008	0.009	0.007	0.005	0.013	0.013	0.024	0.003	0.016	1.97
## 4 0.278	0.011	0.015	0.008	0.006	0.009	0.020	0.035	0.004	0.033	2.10
## 5 0.436	NA	0.024	0.014	0.012	0.025	0.031	0.034	0.012	0.017	5.62
## 8 0.258	0.006	0.010	0.005	0.004	0.008	0.020	0.025	0.004	0.019	1.66
## 9 0.233	0.005	0.012	0.005	0.005	0.009	0.014	0.026	0.003	0.016	2.21
## 10 0.364	0.009	0.012	0.007	0.006	0.014	0.024	0.031	0.004	0.028	2.58
## C5.1	C3.OH	C3.1	C4	C3.DC..C4.OH.	C4.1	C5	C5.M.DC	C5.OH..C3.DC.M.		
## 1 0.030	0.008	0.015	0.082		0.045	0.025	0.094	0.023		0.026
## 4 0.035	0.010	0.017	0.110		0.077	0.031	0.145	0.034		0.041
## 5 0.073	0.029	0.035	0.106		0.099	0.069	0.141	0.094		0.058
## 8 0.021	0.008	0.012	0.082		0.047	0.021	0.107	0.023		0.023

## 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	0.092
## 4	0.685	3.33	18.6	13.30	0.053	0.079
## 5	0.513	1.78	13.8	5.03	NA	0.102
## 8	0.605	2.32	12.1	11.50	NA	0.093
## 9	0.486	2.05	10.4	11.40	NA	0.082
## 10	0.578	2.28	13.8	8.10	NA	0.111
##	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	21.7
## 4	106.0	93.6	1.590	0.190	2.57	20.9
## 5	83.4	35.9	0.709	0.135	1.83	20.5
## 8	83.6	60.6	1.580	0.251	2.16	18.4
## 9	89.6	55.1	1.390	0.221	1.62	18.3
## 10	80.7	82.6	1.640	0.244	1.75	14.6
##	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	1.230
## 4	48.8	41.2	122.0	1.76	0.070	1.160
## 5	28.5	21.9	98.1	1.70	0.048	1.100
## 8	34.4	41.5	110.0	2.03	0.106	1.150
## 9	32.0	41.8	102.0	1.77	0.081	0.902
## 10	38.1	31.2	138.0	1.80	0.092	0.904
##	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	0.074
## 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.aa.C30.0	PC.aa.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.aa.C30.2	PC.aa.C32.1	PC.aa.C32.2	PC.aa.C34.0	PC.aa.C34.1	PC.aa.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 EDTAc_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
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 Kynurenine 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	9.22
## 2	0.065	0.409	0.521	1.99	12.7	5.40
## 3	0.099	0.458	0.605	2.69	16.6	11.60
## 4	0.076	0.486	0.685	3.33	18.6	13.30
## 5	0.069	0.401	0.513	1.78	13.8	5.03
## 6	0.073	0.450	0.620	2.61	14.7	8.98
##	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	2.38
## 2	0.067	64.2	60.5	0.879	0.127	2.05
## 3	0.105	108.0	83.1	1.930	0.210	2.30
## 4	0.079	106.0	93.6	1.590	0.190	2.57
## 5	0.102	83.4	35.9	0.709	0.135	1.83
## 6	0.107	90.2	85.6	1.790	0.213	2.48
##	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	0.084
## 2	14.3	35.6	24.3	83.7	1.05	0.046
## 3	19.9	44.9	43.9	146.0	2.09	0.057
## 4	20.9	48.8	41.2	122.0	1.76	0.070
## 5	20.5	28.5	21.9	98.1	1.70	0.048
## 6	15.5	43.2	46.0	114.0	3.47	0.103
##	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	0.195
## 2	0.946	21.9	78.9	9.91	25.1	0.211
## 3	1.210	34.5	107.0	17.50	36.6	0.212
## 4	1.160	28.7	92.7	14.30	29.9	0.220
## 5	1.100	23.3	101.0	13.80	36.2	0.165
## 6	1.390	28.9	78.0	13.10	48.4	0.205
##	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	0.364
## 2	0.057	0.358	3.39	4.08	14.2	0.419
## 3	0.118	0.395	3.56	5.34	16.7	0.476
## 4	0.097	0.433	3.59	5.06	14.0	0.427
## 5	0.044	0.525	3.37	5.29	22.5	0.125
## 6	0.120	0.346	2.63	3.25	18.9	0.451
##	PC.aa.C42.1	PC.aa.C42.2	PC.aa.C42.4	PC.aa.C42.5	PC.aa.C42.6	PC.aa.C30.0
## 1	0.226	0.108	0.272	0.272	0.291	0.173
## 2	0.216	0.109	0.336	0.317	0.248	0.147
## 3	0.281	0.118	0.300	0.206	0.267	0.209
## 4	0.223	0.119	0.268	0.267	0.254	0.223
## 5	0.095	0.083	0.206	0.205	0.280	0.095
## 6	0.233	0.135	0.228	0.254	0.271	0.221
##	PC.aa.C30.1	PC.aa.C30.2	PC.aa.C32.1	PC.aa.C32.2	PC.aa.C34.0	PC.aa.C34.1
## 1	0.027	0.022	1.65	0.371	0.880	3.66
## 2	0.024	0.020	2.01	0.360	0.763	2.68
## 3	0.046	0.030	2.40	0.477	0.938	4.04
## 4	0.049	0.023	2.47	0.459	0.964	4.06
## 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		
## 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		8.7	
14.2									
## 3		11.7	5.8	9.3	5.6	19.1		15.3	
14.5									
## 4		7.2	9.8	4.8	4.0	13.9		7.7	
11.8									
## 5		44.7	20.2	18.9	18.9	33.9		18.5	
27.7									
## 6		16.0	23.6	7.8	5.5	16.9		16.7	
25.9									
##	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		6.3
## 3	17.8		2.1	8.1	27.4	1343.9	201.3		6.0
## 4	14.7		1.3	6.4	14.4	629.5	322.0		8.6
## 5	35.4		5.5	13.0	40.0	1618.0	271.6		0.0
## 6	18.6		3.4	5.0	35.5	1791.8	274.2		8.8
##	Isobutyrate	Isopropanol	Lactate	Malonate					
## 1	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.
