

#Problem 2

#Calculate the Manhattan distance between any two vectors, vec1 and vec2

#Assign vectors, fill with any two vectors #vec1 <- c(1,2,3,4) #vec2 <- c(2,3,4,5)

```
Manhattan_dist <- function(vec1, vec2){ dist <- sum(abs(vec1 - vec2)) return (dist) }
```

```
Manhattan_dist(vec1,vec2)
```

#Calculate the Euclidian distance between any two vectors, vec1 and vec2

#Assign vectors, fill with any two vectors #vec1 <- c(1,2,3,4) #vec2 <- c(2,3,4,5)

```
Euclidian_dist <- function(vec1, vec2){ dist <- sqrt(sum((vec1 - vec2)^2)) return (dist) }
```

```
Euclidian_dist(vec1,vec2)
```

#Problem #4

```
library(ggplot2)
```

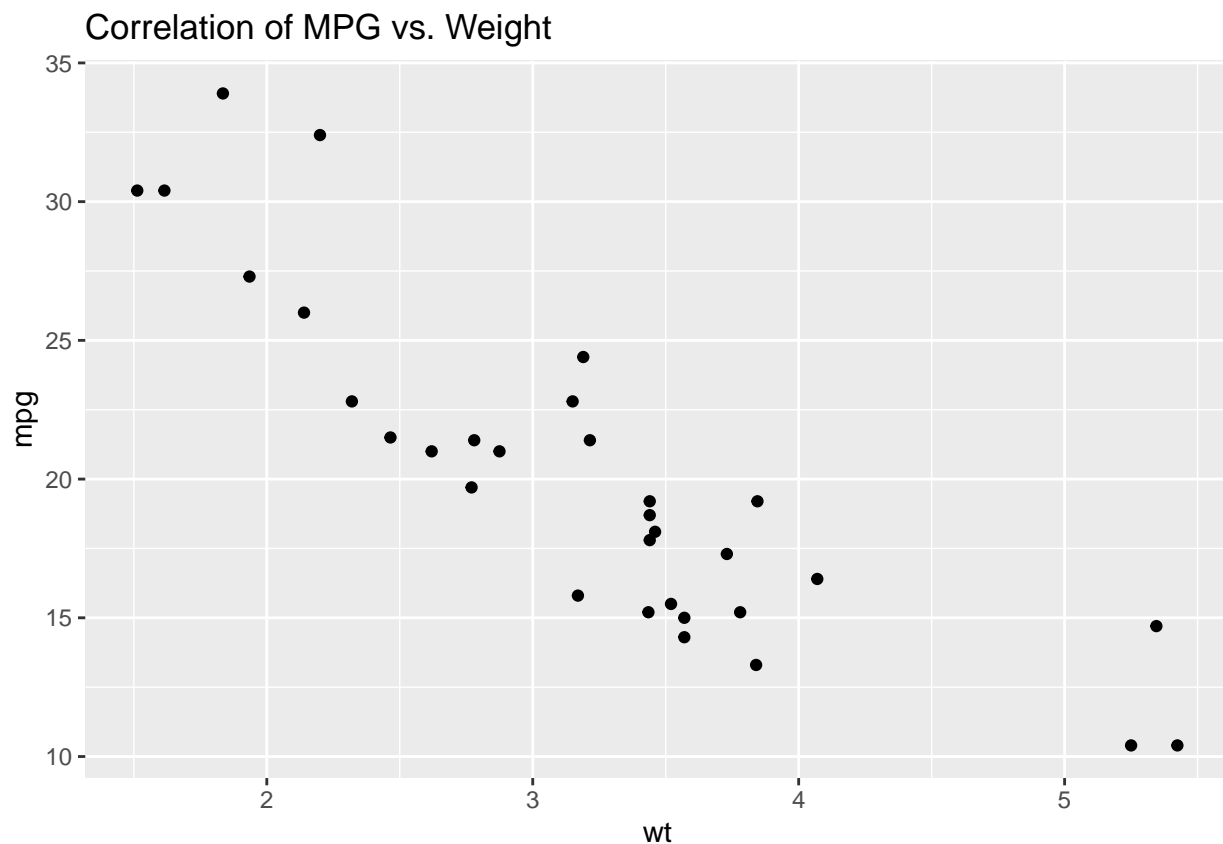
```
data("mtcars")
```

#Calculate correlation between mpg and wt

```
cor_mpg_wt <- cor(mtcars$mpg, mtcars$wt)
```

#Plot correlation onto scatter plot

```
ggplot(mtcars, aes(x = wt, y = mpg)) +  
  geom_point() +  
  labs(title = "Correlation of MPG vs. Weight")
```



#Problem #5

```
setwd("C:/Users/kathryn.poniatowski/OneDrive - Mass Transport Authority, Flint/UofM/CSC 587/Poniatowski")
```

#Import test data into dataframe

```
metabolite <- read.csv("metabolite.csv")
```

#Calculate missing percentage in columns

```
missing_percentage <- colMeans(is.na(metabolite))
```

#Drop columns with over 75% missing

```
metabolite_columns <- missing_percentage <= 0.75
```

#Assign columns to amended dataframe

```
metabolite_amended <- metabolite[, metabolite_columns]
```

Replace missing values in remaining columns with the median of the column

```
metabolite_amended <- metabolite_amended
```

```
for (col in colnames(metabolite_amended)) {
```

Calculate the median excluding NA values

```
median_value <- median(metabolite_amended[[col]], na.rm = TRUE)
```

Replace NA values with the median

```
metabolite_amended[[col]][is.na(metabolite_amended[[col]])] <- median_value
```

```
}
```

```
head(metabolite_amended)
```

```
##      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
## 2 Alzheimer 93.4 138 142 217 1.05      0.929      0.189      1.350      48.8
## 3 Alzheimer 68.6 161 158 208 1.00      0.620      0.198      0.998      30.4
## 4 Alzheimer 94.1 129 162 201 1.10      0.795      0.198      0.675      80.1
## 5 Alzheimer 79.8 126 115 199 1.24      1.360      0.198      1.280      60.5
## 6 Alzheimer 82.5 167 173 333 1.35      1.150      0.198      1.010      24.0
##      DOPA Dopamine Histamine Kynurenine Met.SO Putrescine Sarcosine Serotonin
## 1 0.265 0.233      0.225      5.21 0.526      0.068      17.8      0.147
## 2 0.252 0.231      0.211      5.44 0.387      0.087      20.2      0.231
## 3 0.268 0.231      0.217      5.20 0.651      0.260      14.4      0.196
## 4 0.264 0.234      0.209      5.80 0.389      0.110      18.7      0.255
## 5 0.271 0.231      0.210      4.46 0.466      0.118      22.5      0.390
## 6 0.275 0.231      0.212      7.01 0.417      0.262      30.8      0.140
##      Spermidine t4.OH.Pro Taurine SDMA  C0  C10 C10.1 C10.2  C12 C12.DC C12.1
## 1      0.188      24.0      125 1.13 18.2 0.059 0.312 0.038 0.030 0.042 0.290
## 2      0.233      29.3      120 1.65 17.0 0.051 0.288 0.039 0.038 0.038 0.265
## 3      0.384      20.9      139 1.57 12.6 0.083 0.357 0.054 0.032 0.048 0.302
## 4      0.353      23.1      159 1.34 23.5 0.071 0.317 0.040 0.045 0.048 0.275
## 5      0.473      26.9      149 1.24 13.6 0.139 0.472 0.074 0.056 0.079 0.394
## 6      0.856      26.0      379 1.44 26.7 0.058 0.238 0.042 0.039 0.035 0.196
##      C14 C14.1 C14.1.OH C14.2 C14.2.OH  C16 C16.OH C16.1 C16.1.OH C16.2
## 1 0.023 0.019      0.008 0.008      0.006 0.046      0.008 0.009      0.007 0.005
## 2 0.026 0.017      0.008 0.009      0.009 0.070      0.009 0.013      0.006 0.006
## 3 0.021 0.031      0.010 0.010      0.009 0.076      0.011 0.019      0.010 0.005
## 4 0.026 0.028      0.010 0.013      0.011 0.074      0.011 0.015      0.008 0.006
```

## 5	0.034	0.043	0.016	0.025	0.017	0.062	0.007	0.024	0.014	0.012
## 6	0.029	0.023	0.009	0.010	0.007	0.081	0.006	0.012	0.005	0.007
##	C16.2.OH	C18	C18.1	C18.1.OH	C18.2	C2	C3	C3.OH	C3.1	C4
## 1	0.013	0.013	0.024	0.003	0.016	1.97	0.354	0.008	0.015	0.082
## 2	0.012	0.014	0.025	0.003	0.028	1.95	0.184	0.009	0.013	0.108
## 3	0.013	0.016	0.025	0.004	0.018	1.70	0.371	0.011	0.012	0.057
## 4	0.009	0.020	0.035	0.004	0.033	2.10	0.278	0.010	0.017	0.110
## 5	0.025	0.031	0.034	0.012	0.017	5.62	0.436	0.029	0.035	0.106
## 6	0.015	0.017	0.035	0.004	0.029	3.49	0.461	0.008	0.014	0.123
##	C3.DC..C4.OH.	C4.1	C5	C5.M.DC	C5.OH..C3.DC.M.	C5.1	C5.1.DC	C6..C4.1.DC.		
## 1	0.045	0.025	0.094	0.023		0.026	0.030	0.020		0.022
## 2	0.080	0.025	0.077	0.032		0.026	0.024	0.021		0.030
## 3	0.035	0.039	0.096	0.045		0.024	0.037	0.018		0.022
## 4	0.077	0.031	0.145	0.034		0.041	0.035	0.016		0.029
## 5	0.099	0.069	0.141	0.094		0.058	0.073	0.049		0.052
## 6	0.068	0.026	0.090	0.019		0.037	0.022	0.016		0.063
##	C5.DC..C6.OH.	C6.1	C7.DC	C8	C9	lysoPC.a.C14.0	lysoPC.a.C16.0			
## 1	0.014	0.018	0.011	0.062	0.016		2.23			37.9
## 2	0.018	0.015	0.010	0.058	0.014		1.97			22.1
## 3	0.029	0.031	0.021	0.090	0.017		2.12			33.7
## 4	0.016	0.027	0.017	0.091	0.018		2.19			32.8
## 5	0.040	0.040	0.036	0.192	0.041		1.88			24.5
## 6	0.016	0.019	0.014	0.073	0.014		2.11			29.1
##	lysoPC.a.C16.1	lysoPC.a.C17.0	lysoPC.a.C18.0	lysoPC.a.C18.1	lysoPC.a.C18.2					
## 1	2.66		0.446		9.00		8.58			7.27
## 2	1.31		0.270		5.35		3.94			4.42
## 3	2.53		0.399		7.51		7.73			8.02
## 4	2.39		0.323		7.21		7.22			7.62
## 5	1.27		0.382		6.66		5.39			3.60
## 6	2.09		0.348		5.84		6.30			8.10
##	lysoPC.a.C20.3	lysoPC.a.C20.4	lysoPC.a.C24.0	lysoPC.a.C26.0	lysoPC.a.C26.1					
## 1	1.830		8.25		0.079		0.113			0.053
## 2	0.958		4.60		0.059		0.066			0.042
## 3	2.050		9.84		0.075		0.126			0.049
## 4	1.640		6.75		0.066		0.086			0.045
## 5	0.970		6.26		0.084		0.118			0.053
## 6	1.970		7.04		0.083		0.112			0.050
##	lysoPC.a.C28.0	lysoPC.a.C28.1	PC.aa.C24.0	PC.aa.C26.0	PC.aa.C28.1	PC.aa.C30.0				
## 1	0.108		0.072		0.082		0.438		0.571	2.35
## 2	0.076		0.058		0.065		0.409		0.521	1.99
## 3	0.078		0.092		0.099		0.458		0.605	2.69
## 4	0.076		0.076		0.076		0.486		0.685	3.33
## 5	0.092		0.072		0.069		0.401		0.513	1.78
## 6	0.099		0.083		0.073		0.450		0.620	2.61
##	PC.aa.C32.0	PC.aa.C32.1	PC.aa.C32.2	PC.aa.C32.3	PC.aa.C34.1	PC.aa.C34.2				
## 1	11.4		9.22		0.117		0.092		109.0	71.0
## 2	12.7		5.40		0.117		0.067		64.2	60.5
## 3	16.6		11.60		0.117		0.105		108.0	83.1
## 4	18.6		13.30		0.053		0.079		106.0	93.6
## 5	13.8		5.03		0.117		0.102		83.4	35.9
## 6	14.7		8.98		0.117		0.107		90.2	85.6
##	PC.aa.C34.3	PC.aa.C34.4	PC.aa.C36.0	PC.aa.C36.1	PC.aa.C36.2	PC.aa.C36.3				
## 1	1.430		0.200		2.38		21.7		42.4	42.7
## 2	0.879		0.127		2.05		14.3		35.6	24.3

## 3	1.930	0.210	2.30	19.9	44.9	43.9
## 4	1.590	0.190	2.57	20.9	48.8	41.2
## 5	0.709	0.135	1.83	20.5	28.5	21.9
## 6	1.790	0.213	2.48	15.5	43.2	46.0
##	PC.aa.C36.4	PC.aa.C36.5	PC.aa.C36.6	PC.aa.C38.0	PC.aa.C38.3	PC.aa.C38.4
## 1	120.0	1.86	0.084	1.230	32.1	95.1
## 2	83.7	1.05	0.046	0.946	21.9	78.9
## 3	146.0	2.09	0.057	1.210	34.5	107.0
## 4	122.0	1.76	0.070	1.160	28.7	92.7
## 5	98.1	1.70	0.048	1.100	23.3	101.0
## 6	114.0	3.47	0.103	1.390	28.9	78.0
##	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	16.80	41.6	0.195	0.074	0.491	3.48
## 2	9.91	25.1	0.211	0.057	0.358	3.39
## 3	17.50	36.6	0.212	0.118	0.395	3.56
## 4	14.30	29.9	0.220	0.097	0.433	3.59
## 5	13.80	36.2	0.165	0.044	0.525	3.37
## 6	13.10	48.4	0.205	0.120	0.346	2.63
##	PC.aa.C40.5	PC.aa.C40.6	PC.aa.C42.0	PC.aa.C42.1	PC.aa.C42.2	PC.aa.C42.4
## 1	5.66	21.8	0.364	0.226	0.108	0.272
## 2	4.08	14.2	0.419	0.216	0.109	0.336
## 3	5.34	16.7	0.476	0.281	0.118	0.300
## 4	5.06	14.0	0.427	0.223	0.119	0.268
## 5	5.29	22.5	0.125	0.095	0.083	0.206
## 6	3.25	18.9	0.451	0.233	0.135	0.228
##	PC.aa.C42.5	PC.aa.C42.6	PC.aa.C30.0	PC.aa.C30.1	PC.aa.C30.2	PC.aa.C32.1
## 1	0.272	0.291	0.173	0.027	0.022	1.65
## 2	0.317	0.248	0.147	0.024	0.020	2.01
## 3	0.206	0.267	0.209	0.046	0.030	2.40
## 4	0.267	0.254	0.223	0.049	0.023	2.47
## 5	0.205	0.280	0.095	0.082	0.023	1.72
## 6	0.254	0.271	0.221	0.039	0.029	2.01
##	PC.aa.C32.2	PC.aa.C34.0	PC.aa.C34.1	PC.aa.C34.2	PC.aa.C34.3	PC.aa.C36.0
## 1	0.371	0.880	3.66	2.48	0.813	0.498
## 2	0.360	0.763	2.68	2.32	0.905	0.398
## 3	0.477	0.938	4.04	2.95	1.030	0.554
## 4	0.459	0.964	4.06	3.09	1.020	0.552
## 5	0.316	1.060	3.28	1.70	0.722	0.553
## 6	0.397	0.920	3.26	2.58	1.000	0.443
##	PC.aa.C36.1	PC.aa.C36.2	PC.aa.C36.3	PC.aa.C36.4	PC.aa.C36.5	PC.aa.C38.0
## 1	5.64	1.90	1.170	6.96	4.79	0.474
## 2	3.89	1.54	0.873	6.40	5.36	0.325
## 3	5.95	2.29	1.240	9.05	6.63	0.478
## 4	4.75	2.01	1.350	8.36	5.97	0.397
## 5	5.95	1.47	0.760	4.78	4.00	0.430
## 6	4.95	2.05	1.170	7.04	4.47	0.590
##	PC.aa.C38.2	PC.aa.C38.3	PC.aa.C38.4	PC.aa.C38.5	PC.aa.C38.6	PC.aa.C40.1
## 1	0.538	2.66	6.33	5.51	1.95	0.574
## 2	0.127	1.80	5.37	4.49	1.63	0.281
## 3	0.154	2.87	7.06	5.64	1.98	0.759
## 4	0.144	1.97	5.99	5.63	1.97	0.425
## 5	0.246	1.80	5.45	4.34	1.51	0.430
## 6	0.312	2.46	5.55	4.60	1.80	0.481
##	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	0.575	0.940	1.76	1.77	1.59	0.629			
## 2	0.491	0.702	1.43	1.55	1.20	0.616			
## 3	0.654	0.817	1.51	1.64	1.49	0.686			
## 4	0.540	0.742	1.45	1.62	1.25	0.637			
## 5	0.432	0.632	1.10	1.25	1.47	0.660			
## 6	0.598	0.826	1.25	1.38	1.61	0.669			
##	PC.ae.C42.1	PC.ae.C42.2	PC.ae.C42.3	PC.ae.C42.4	PC.ae.C42.5	PC.ae.C44.3			
## 1	0.316	0.192	0.277	0.264	0.888	0.065			
## 2	0.260	0.157	0.200	0.311	0.840	0.071			
## 3	0.356	0.241	0.288	0.319	0.957	0.065			
## 4	0.299	0.159	0.208	0.392	0.863	0.069			
## 5	0.355	0.138	0.174	0.162	0.513	0.081			
## 6	0.265	0.195	0.253	0.316	0.814	0.085			
##	PC.ae.C44.4	PC.ae.C44.5	PC.ae.C44.6	SM..OH..C14.1	SM..OH..C16.1	SM..OH..C22.1			
## 1	0.168	0.536	0.494	1.420	1.33	2.07			
## 2	0.220	0.470	0.515	1.390	1.25	2.47			
## 3	0.228	0.565	0.603	1.840	1.58	2.69			
## 4	0.237	0.517	0.611	1.720	1.48	2.97			
## 5	0.154	0.178	0.134	0.987	1.48	1.96			
## 6	0.232	0.554	0.539	1.320	1.12	2.51			
##	SM..OH..C22.2	SM..OH..C24.1	SM.C16.0	SM.C16.1	SM.C18.0	SM.C18.1	SM.C20.2		
## 1	1.86	0.597	44.9	7.99	14.5	10.40	0.290		
## 2	2.20	0.640	42.1	6.88	12.7	8.52	0.211		
## 3	2.63	0.665	44.8	8.91	14.6	11.60	0.304		
## 4	2.84	0.682	52.4	8.61	17.2	11.50	0.261		
## 5	1.74	0.478	40.6	5.86	13.0	8.34	0.196		
## 6	2.16	0.640	42.6	8.49	13.0	10.60	0.270		
##	SM.C24.0	SM.C24.1	SM.C26.0	SM.C26.1	H1_1	H1	Urea_N	L.Arginine_N	L.Leucine_N
## 1	12.20	27.3	0.147	0.337	3356	3356	185.05	45.1	55.75
## 2	10.40	25.6	0.130	0.317	2509	2509	201.90	22.5	35.30
## 3	11.50	28.8	0.163	0.364	2661	2661	193.30	21.0	25.40
## 4	11.80	27.9	0.138	0.353	2652	2652	500.80	16.0	27.10
## 5	9.29	20.5	0.111	0.283	2258	2258	132.50	13.2	57.90
## 6	9.58	23.7	0.135	0.316	3031	3031	193.30	32.2	26.50
##	EDTAca_N	X2.Hydroxybutyrate	X3.Hydroxybutyrate	Acetate	Acetoacetate	Acetone			
## 1	2.9	19.80	44.1	20.2	21.4	10.15			
## 2	2.0	12.40	8.5	13.2	5.7	5.10			
## 3	1.8	11.33	11.7	5.8	9.3	5.60			
## 4	2.5	12.70	7.2	9.8	4.8	4.00			
## 5	2.5	35.20	44.7	20.2	18.9	18.90			
## 6	0.0	17.20	16.0	23.6	7.8	5.50			
##	Betaine	Carnitine	Choline	Creatine	Dimethyl.sulfone	Ethanol	Formate	Glucose	
## 1	32.25	13.1	22.15	26.7	3.55	7.2	28.9	2239.35	
## 2	22.00	8.7	14.20	14.5	4.70	16.6	24.6	1489.70	
## 3	19.10	15.3	14.50	17.8	2.10	8.1	27.4	1343.90	
## 4	13.90	7.7	11.80	14.7	1.30	6.4	14.4	629.50	
## 5	33.90	18.5	27.70	35.4	5.50	13.0	40.0	1618.00	
## 6	16.90	16.7	25.90	18.6	3.40	5.0	35.5	1791.80	
##	Glycerol	Hypoxanthine	Isobutyrate	Isopropanol	Lactate	Malonate			
## 1	449.1	7.35	4.6	3.3	1768.7	11.35			
## 2	324.6	6.30	3.6	1.9	1171.6	10.40			
## 3	201.3	6.00	2.5	2.5	1938.1	13.10			
## 4	322.0	8.60	2.5	4.4	1037.7	7.60			
## 5	271.6	0.00	6.1	11.2	2199.9	11.70			

```
## 6      274.2      8.80      2.3      2.4 1486.7      11.80
```

```
library(ggplot2)
library(stats)

# Exclude non-numeric columns from the dataframe
numeric_data <- metabolite_amended[sapply(metabolite_amended, is.numeric)]

# Perform PCA on the numeric data (excluding the Label column if it exists)
pca <- prcomp(numeric_data, scale. = TRUE)

# Get the first two principal components (PC1, PC2)
pca_data <- as.data.frame(pca$x[, 1:2])

# Add the 'Label' column from the original dataframe to the PCA data
pca_data$Label <- metabolite_amended$Label

# Plotting the scatter plot of PC1 vs PC2, colored by 'Label'
ggplot(pca_data, aes(x = PC1, y = PC2, color = Label)) +
  geom_point() +
  labs(title = "Metabolite PCA Scatter Plot") +
  scale_color_manual(values = c("Group1" = "blue", "Group2" = "red"))
```

```
## Warning: No shared levels found between `names(values)` of the manual scale and the
## data's colour values.
## No shared levels found between `names(values)` of the manual scale and the
## data's colour values.
```

Metabolite PCA Scatter Plot

