

Multiple Regression Diagnostics and Simultaneous Multiple Regression Tutorial

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Multiple Regression Diagnostics Demonstration

Applied Prompt:

A local resident recently joined their a newly developed charitable giving program and is interested in finding out more about prosocial behavior among group members. Without knowing much about the literature, they developed a short survey to give to a nearby school district to find out.

****Research Question**:** What influences the amount of money given to a program for community outreach.

Ultimately, they must report their findings to their program coordinator and a statistician, who is also on the board.

Variables in Data:

ID — identifying number for each observation

Belief — belief that charitable giving has a positive effect on a scale of 1-10

Need — rating of perceived amount of need required in the community on a scale of 1-10

Interest — rating of level of interest in the project on a scale of 1-10

Happy — rating of happiness felt when making donations on a scale of 1-10

Amount — amount given from 0 - 10 dollars

For all variables:

A value of 99 implies that a response was missing for that variable and case

Load in Libraries

```
library(mice)
library(MVN)
library(lmtest)
library(apaTables)
library(psych)
library(jmv)
library(tidyverse)
```

Load in Data

```
#Load in the data frame into an object `dat`

dat <- read.csv("RegDiagnostics2.csv")

#View(dat)
```

Change Missing Data Labels

```
#Remove all `99` values from data frame and replace with `NA`s
# MAKE SURE YOU DO NOT HAVE AN ID SET TO 99 OR IT WILL REMOVE THIS CASE NUMBER!!

dat[dat=="99"] <- NA
```

Exploratory Data Analysis

Descriptive Statistics

```
# Descriptive statistics (Default: No Removal of Missing Data)
describe(dat)
```

```
##          vars    n   mean    sd median trimmed   mad   min   max range  skew
## ID          1 106 152.50 30.74  152.5  152.50 39.29 100.0 205.0 105.0  0.00
## Amount      2 105   8.03  0.66   8.2   8.10  0.44   4.5   9.1   4.6 -2.74
## Need        3 104   5.83  1.77   5.9   5.90  2.08   1.7   9.5   7.8 -0.32
## Interest    4 104   6.14  0.75   6.2   6.16  0.89   4.4   7.5   3.1 -0.25
## Happy       5 106   8.84  0.85   9.1   8.92  0.59   3.4  10.0   6.6 -2.67
## Belief      6 104   7.53  0.38   7.5   7.53  0.30   6.1   8.3   2.2 -0.56
##          kurtosis   se
## ID          -1.23 2.99
## Amount      11.27 0.06
## Need        -0.71 0.17
## Interest    -0.65 0.07
## Happy       13.81 0.08
## Belief      1.63 0.04
```

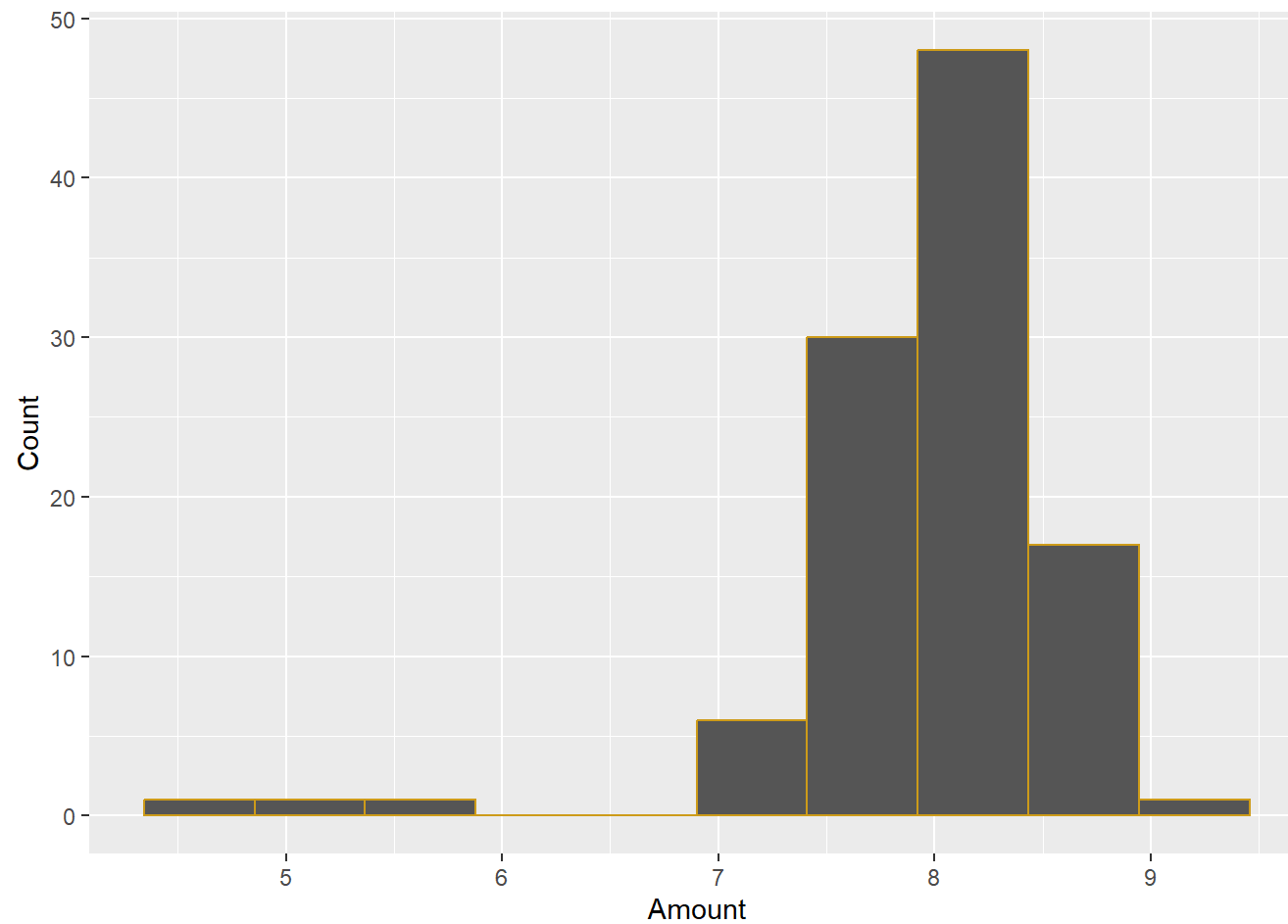
```
# Descriptive statistics (Removal of Missing Data)
describe(dat,
          na.rm=FALSE)
```

```
##          vars    n   mean    sd median trimmed   mad   min   max range  skew
## ID          1 100 153.46 30.51  153.5  153.62 38.55 100.0 205.0 105.0 -0.04
## Amount      2 100   8.06  0.60   8.2   8.11  0.44   4.5   9.1   4.6 -2.66
## Need        3 100   5.78  1.78   5.9   5.85  2.08   1.7   9.5   7.8 -0.28
## Interest    4 100   6.12  0.75   6.2   6.14  0.89   4.4   7.5   3.1 -0.25
## Happy       5 100   8.86  0.85   9.1   8.96  0.59   3.4  10.0   6.6 -2.84
## Belief      6 100   7.54  0.38   7.5   7.54  0.30   6.1   8.3   2.2 -0.61
##          kurtosis   se
## ID          -1.21 3.05
## Amount      12.74 0.06
## Need        -0.75 0.18
## Interest    -0.68 0.07
## Happy       14.99 0.08
## Belief      1.78 0.04
```

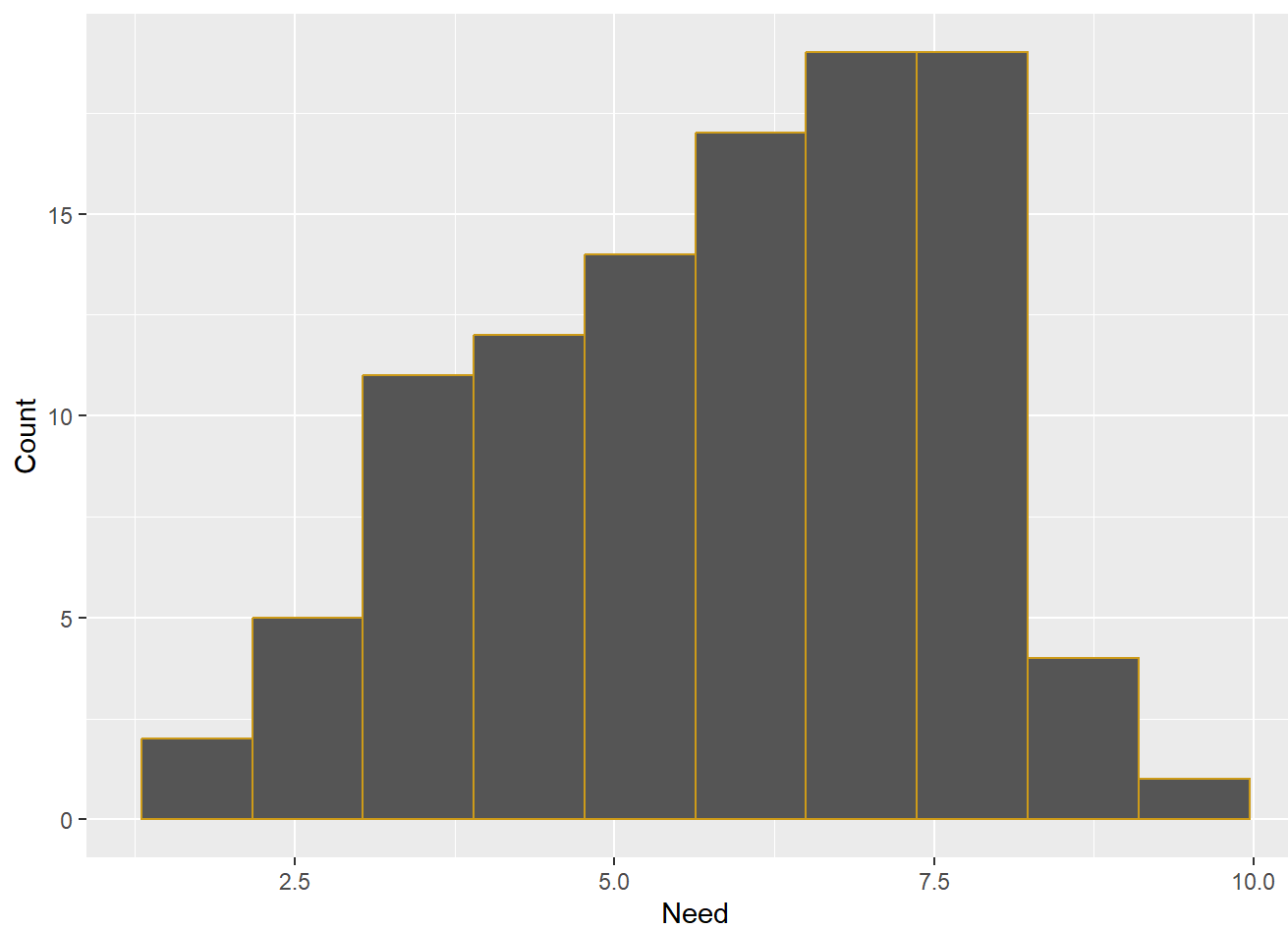
Generate Univariate and Bivariate Visualizations

```
# Univariate Histograms of Continuous Variables
```

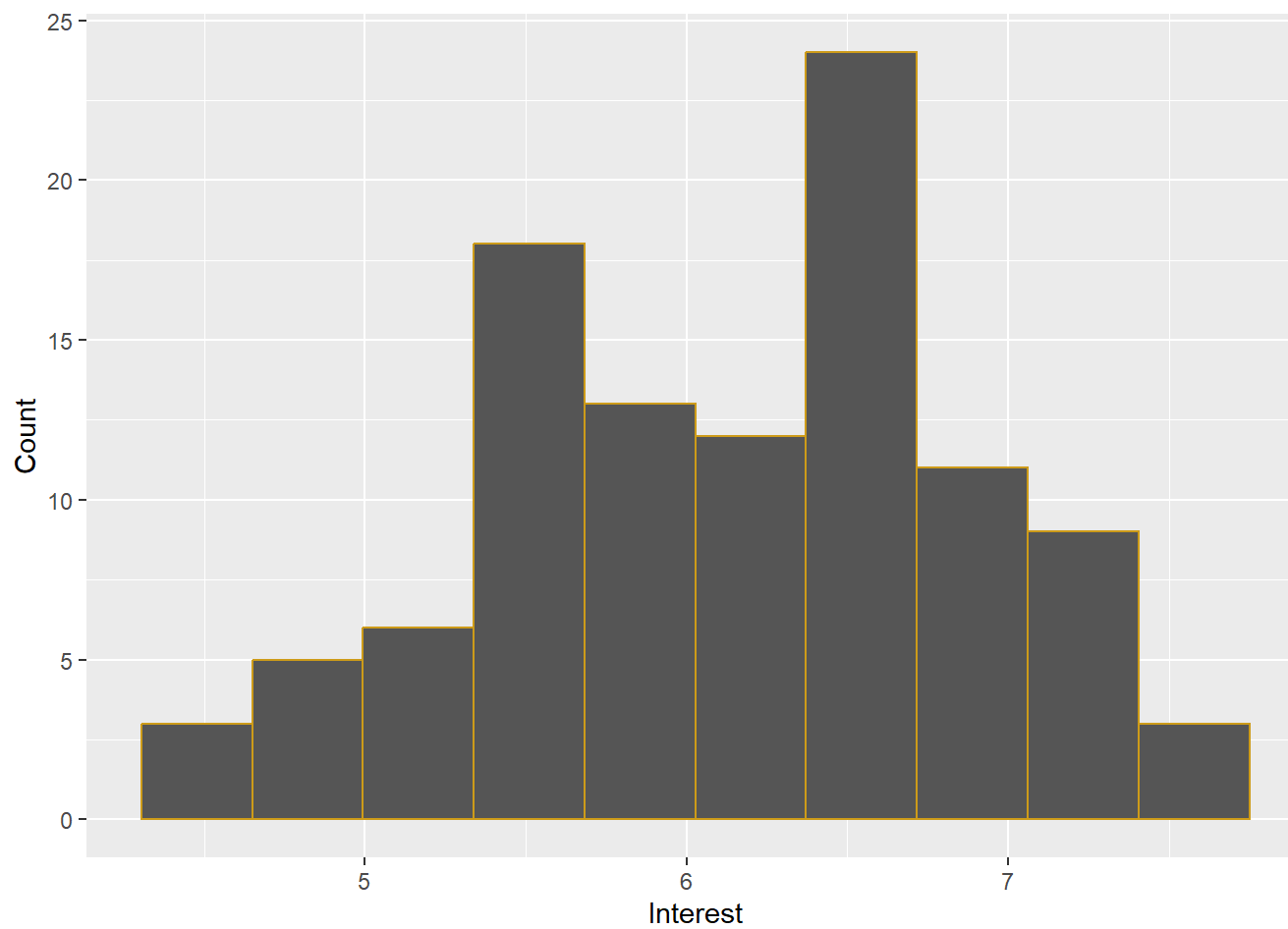
```
ggplot(data = dat,  
       mapping = aes(x = Amount)) +  
  geom_histogram(bins = 10, color = "goldenrod3") +  
  labs(y = "Count", x = "Amount")
```



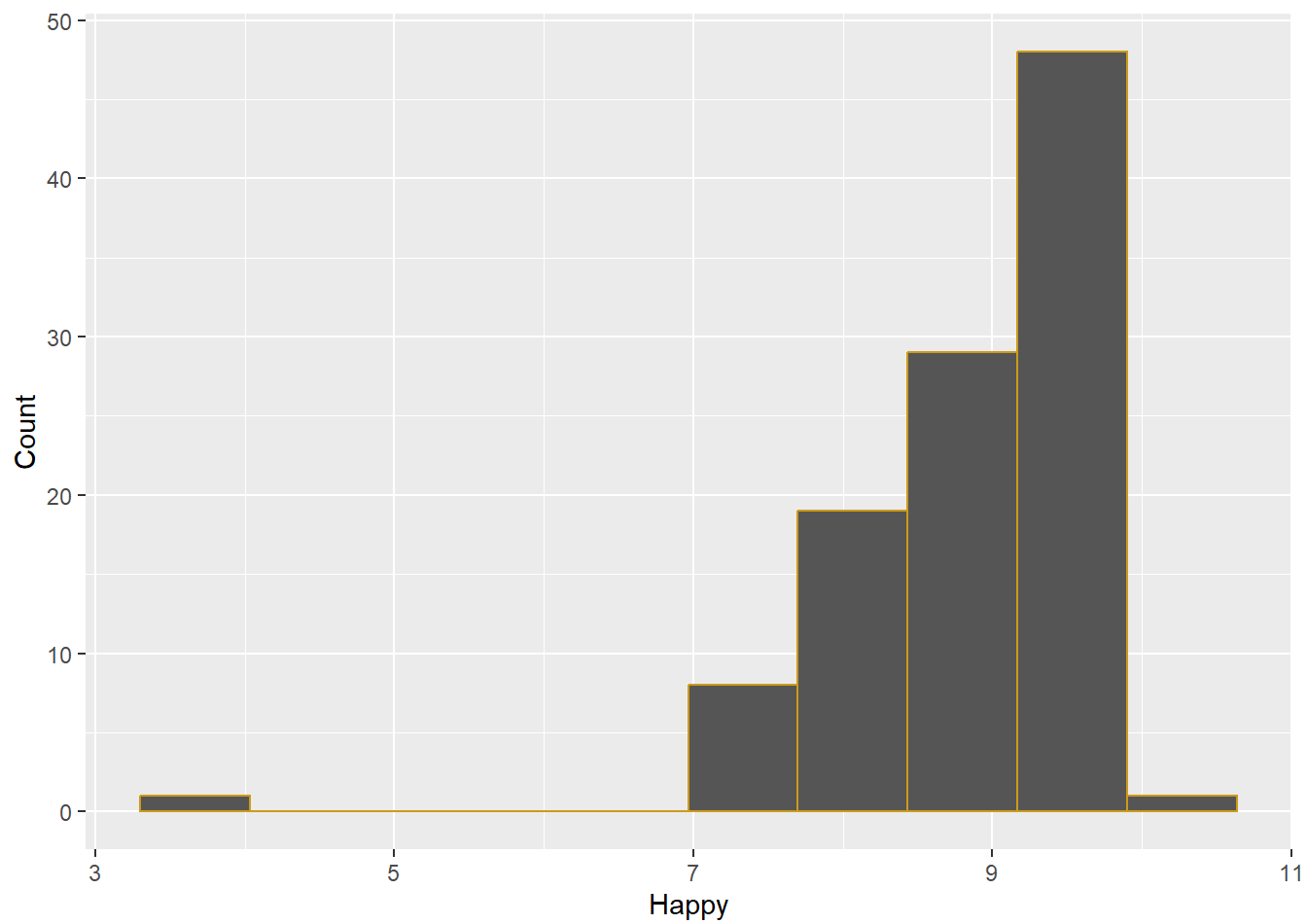
```
ggplot(data = dat,  
       mapping = aes(x = Need)) +  
  geom_histogram(bins = 10, color = "goldenrod3") +  
  labs(y = "Count", x = "Need")
```



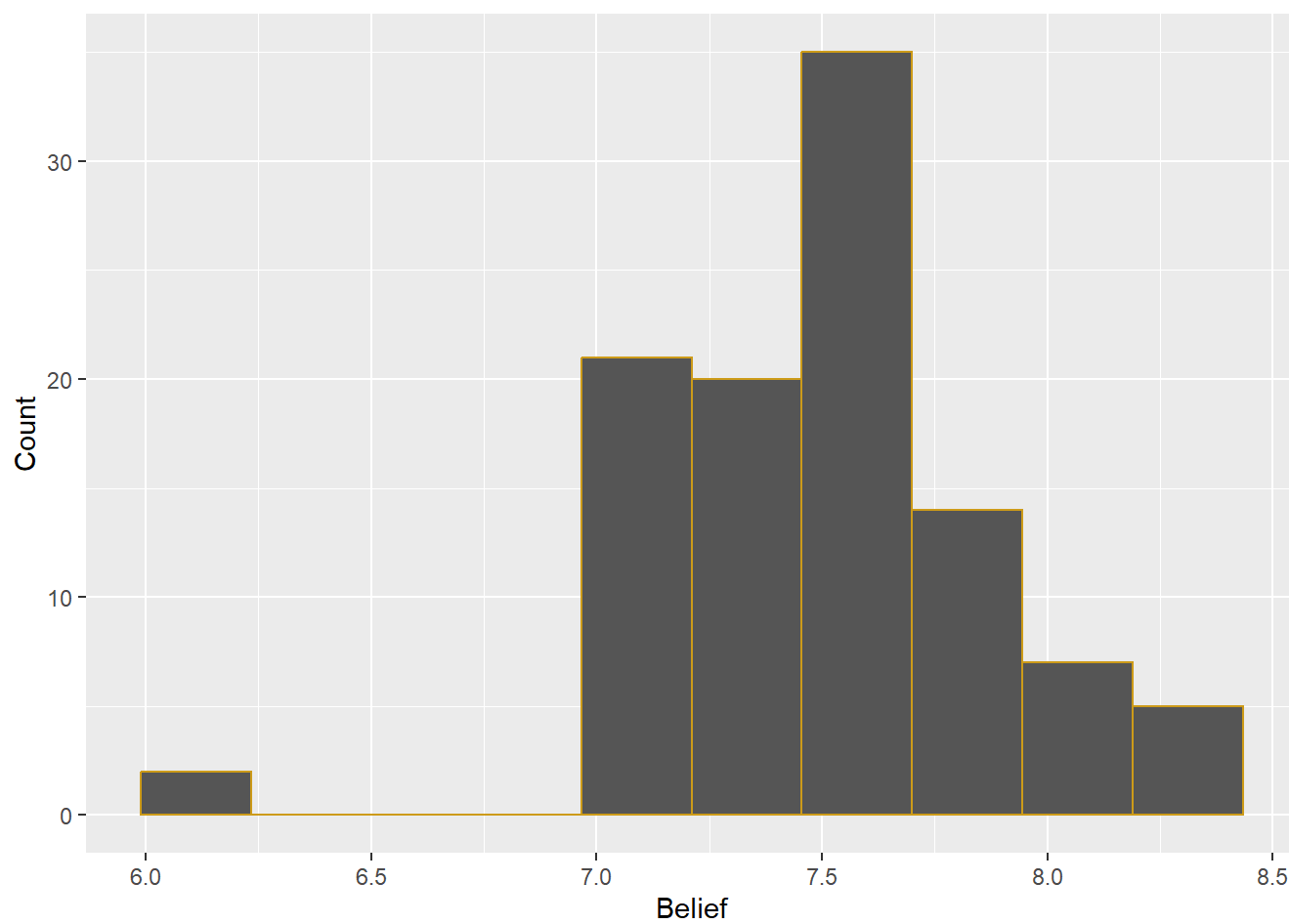
```
ggplot(data = dat,  
       mapping = aes(x = Interest)) +  
  geom_histogram(bins = 10, color = "goldenrod3") +  
  labs(y = "Count", x = "Interest")
```



```
ggplot(data = dat,  
       mapping = aes(x = Happy)) +  
  geom_histogram(bins = 10, color = "goldenrod3") +  
  labs(y = "Count", x = "Happy")
```

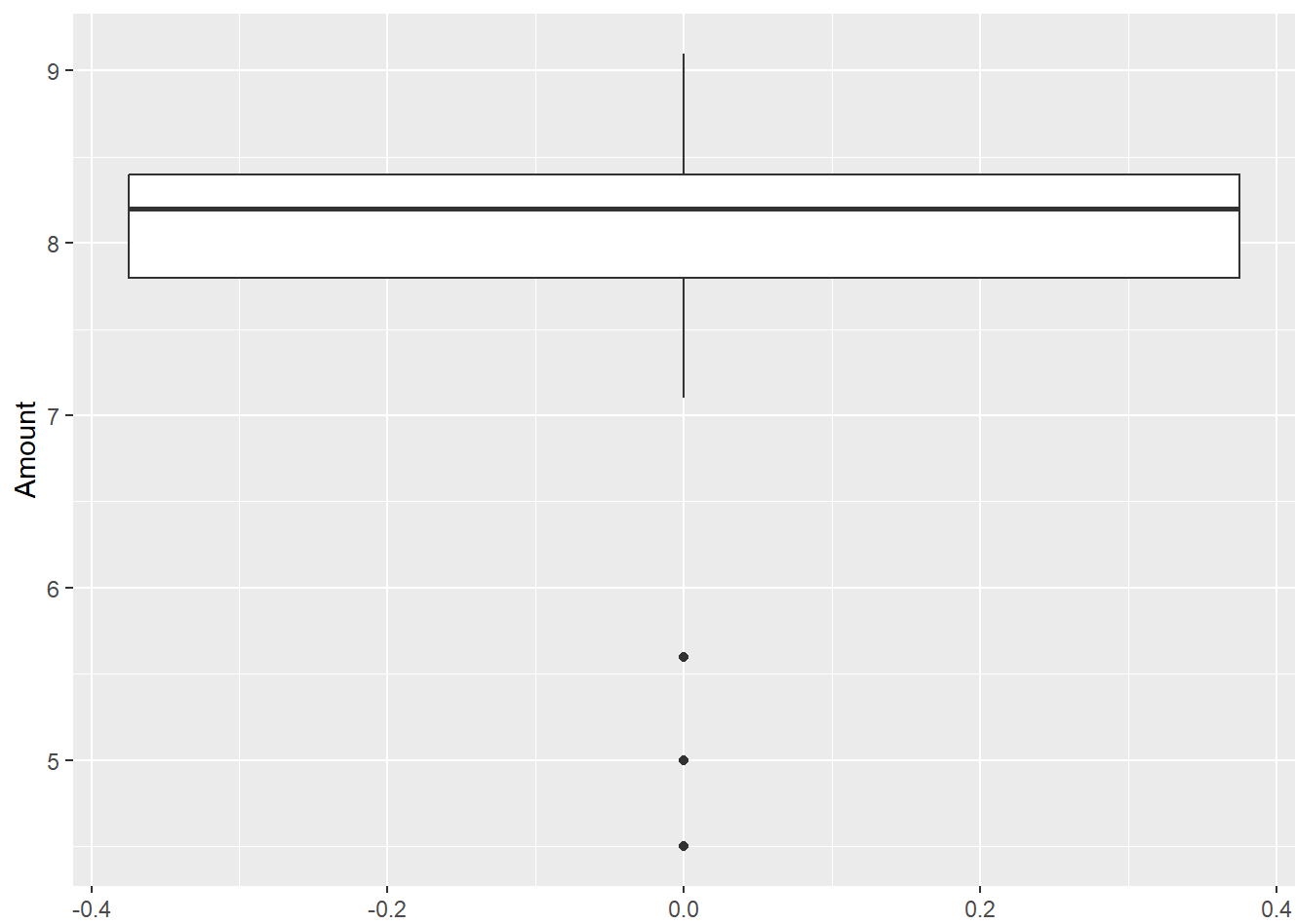


```
ggplot(data = dat,  
       mapping = aes(x = Belief)) +  
  geom_histogram(bins = 10, color = "goldenrod3") +  
  labs(y = "Count", x = "Belief")
```

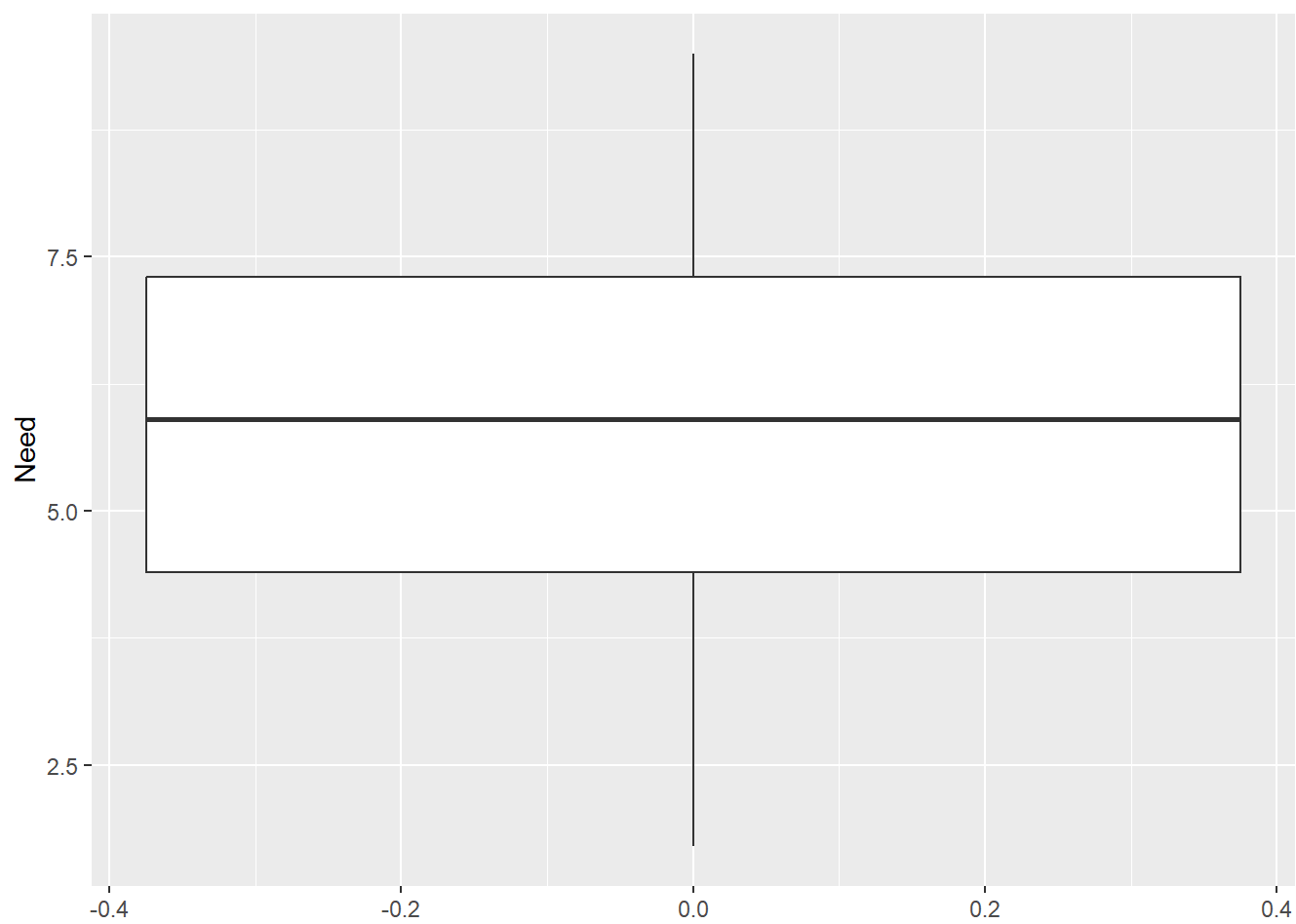


Univariate Boxplots of Continuous Variables

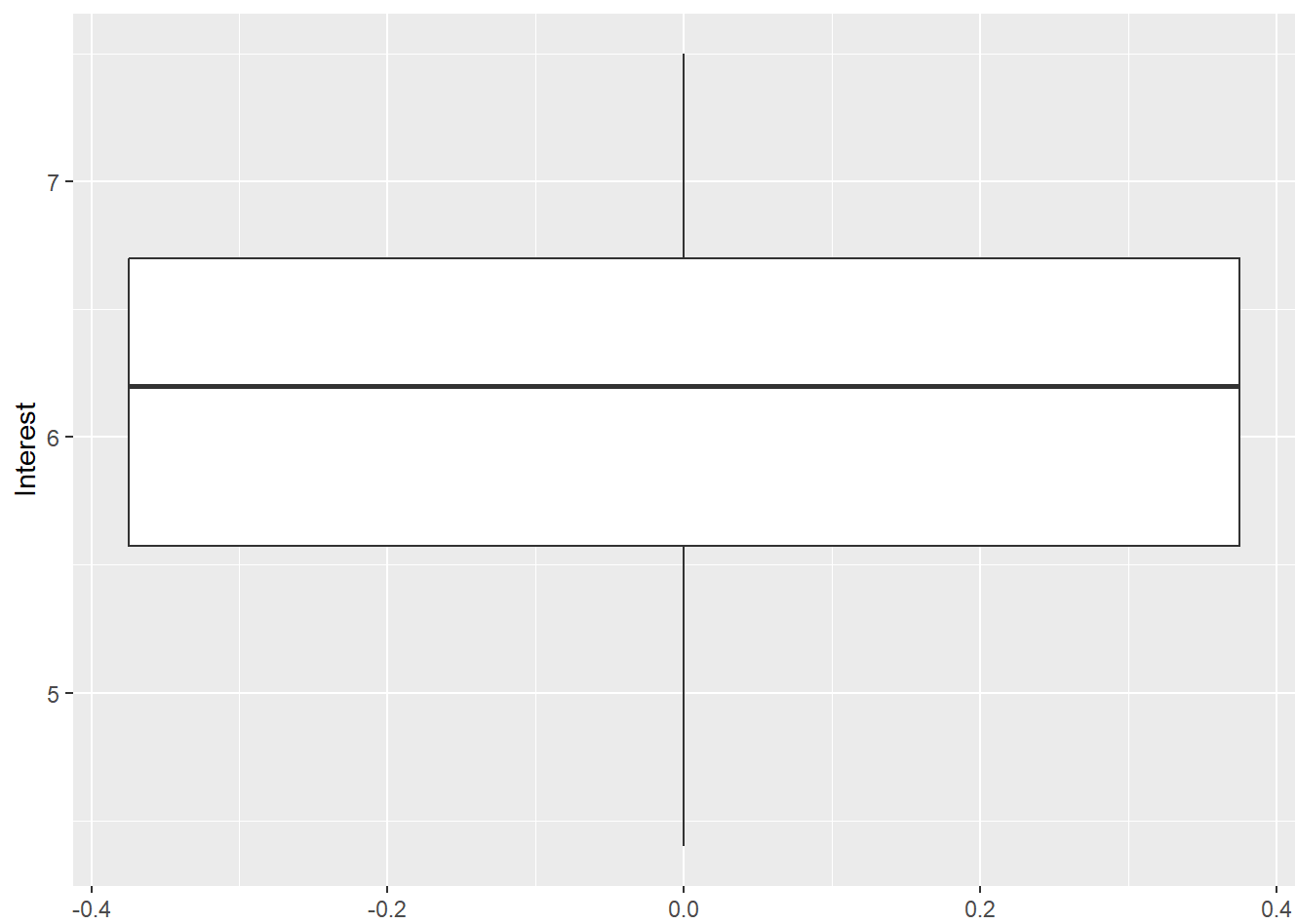
```
ggplot(data = dat,  
       mapping = aes(y = Amount)) +  
  geom_boxplot() +  
  labs(y = "Amount")
```

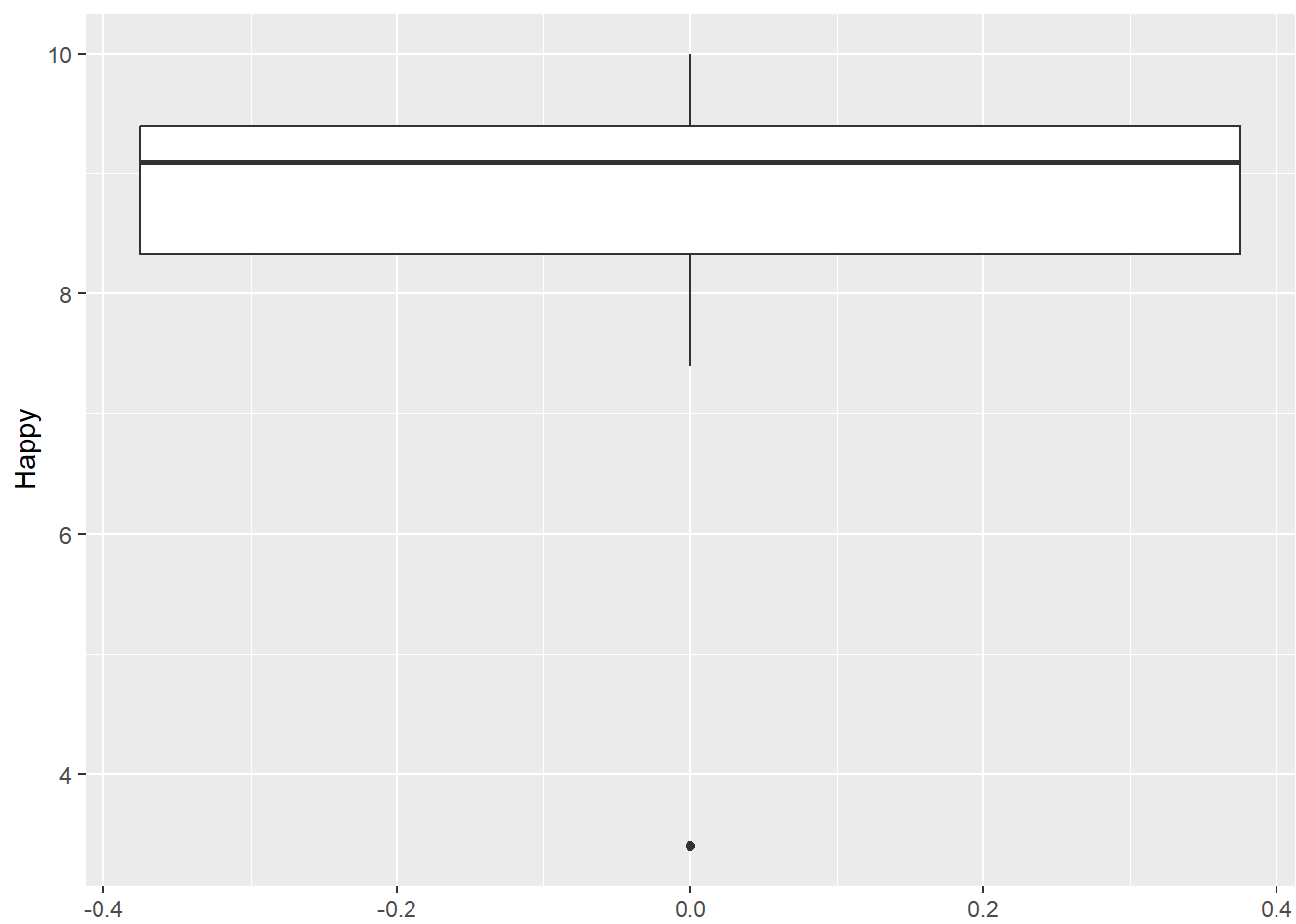
```
ggplot(data = dat,  
       mapping = aes(y = Need)) +  
  geom_boxplot() +  
  labs(y = "Need")
```



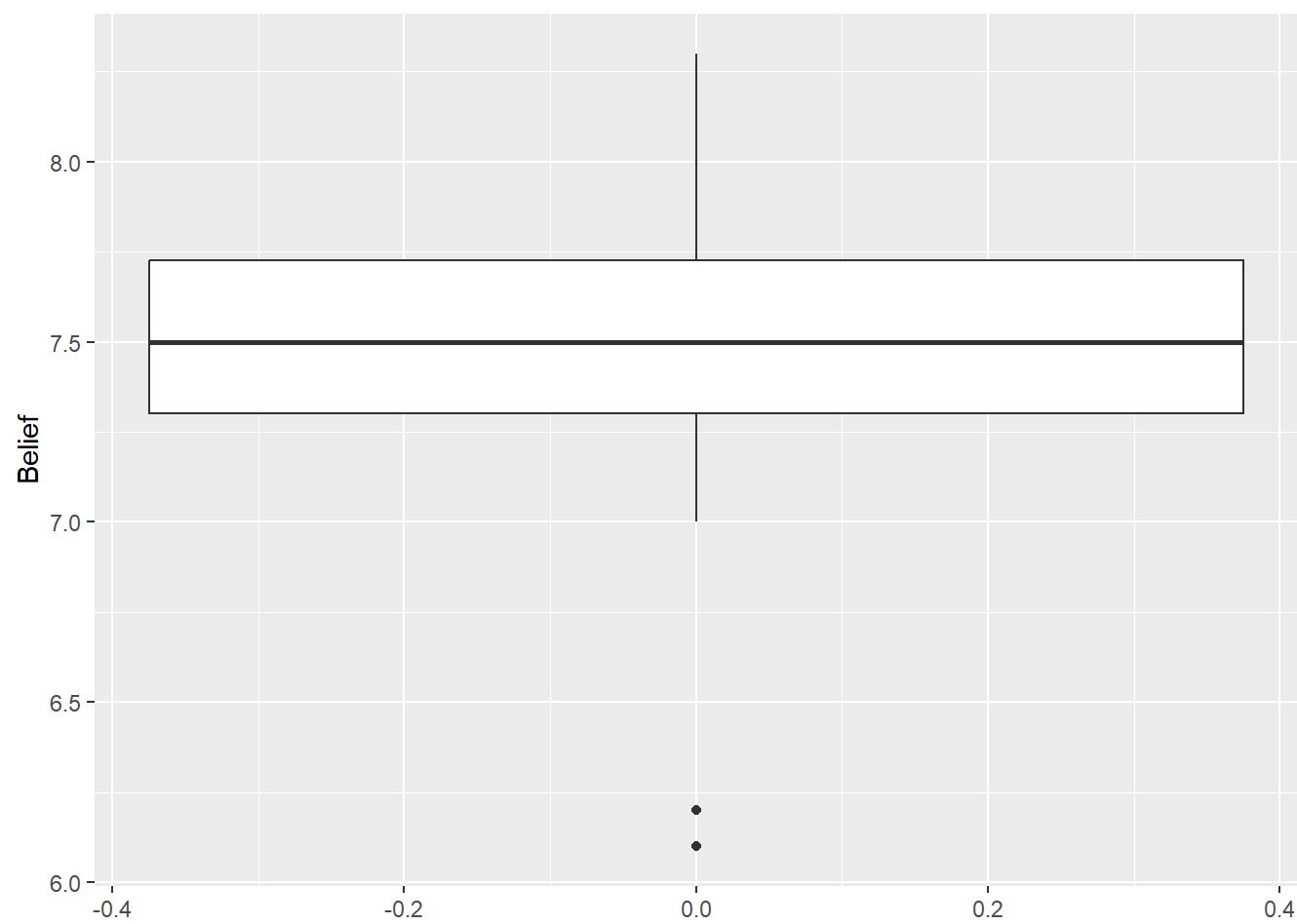
```
ggplot(data = dat,  
       mapping = aes(y = Interest)) +  
  geom_boxplot() +  
  labs(y = "Interest")
```



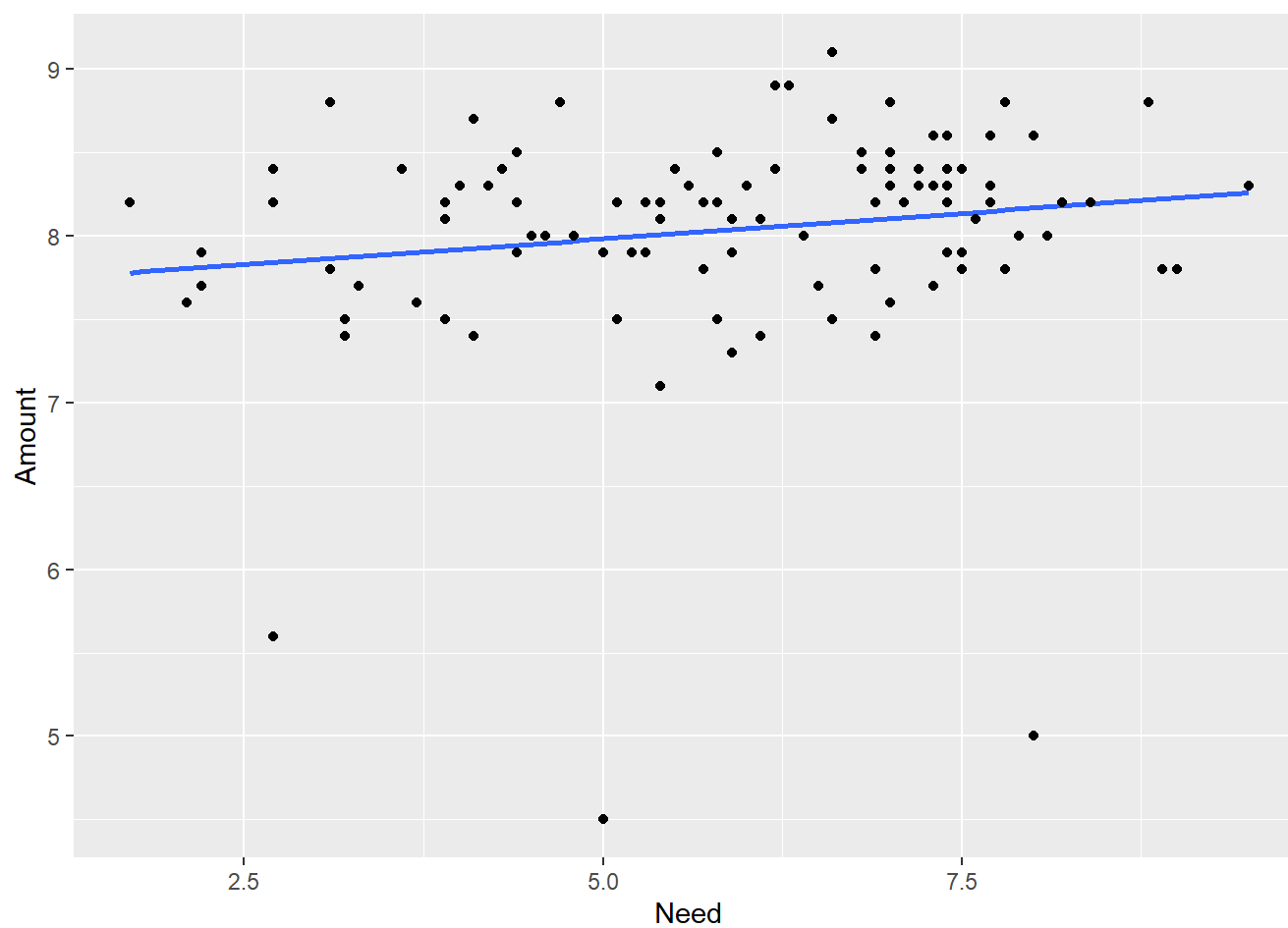
```
ggplot(data = dat,  
       mapping = aes(y = Happy)) +  
  geom_boxplot() +  
  labs(y = "Happy")
```



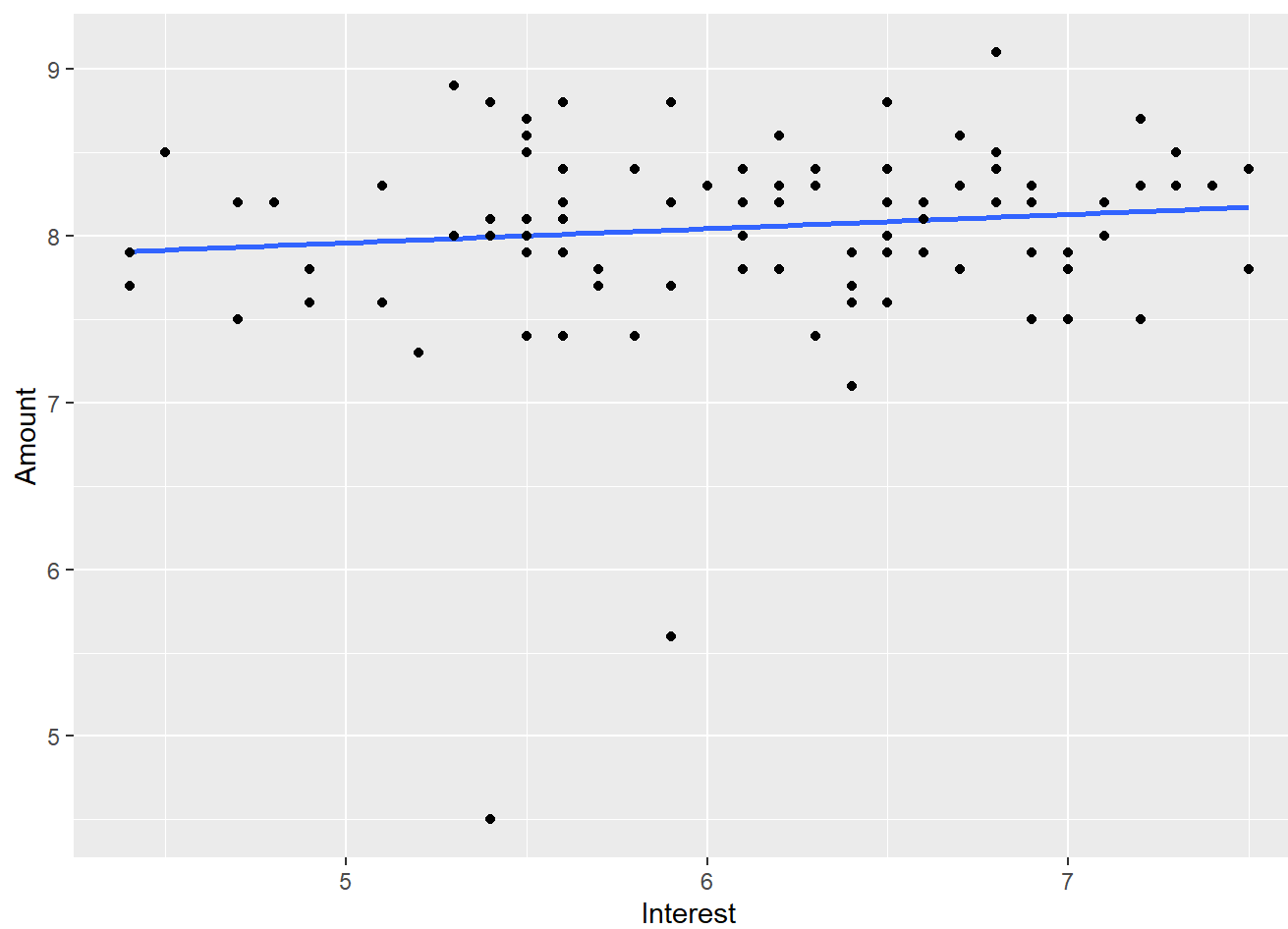
```
ggplot(data = dat,  
       mapping = aes(y = Belief)) +  
  geom_boxplot() +  
  labs(y = "Belief")
```



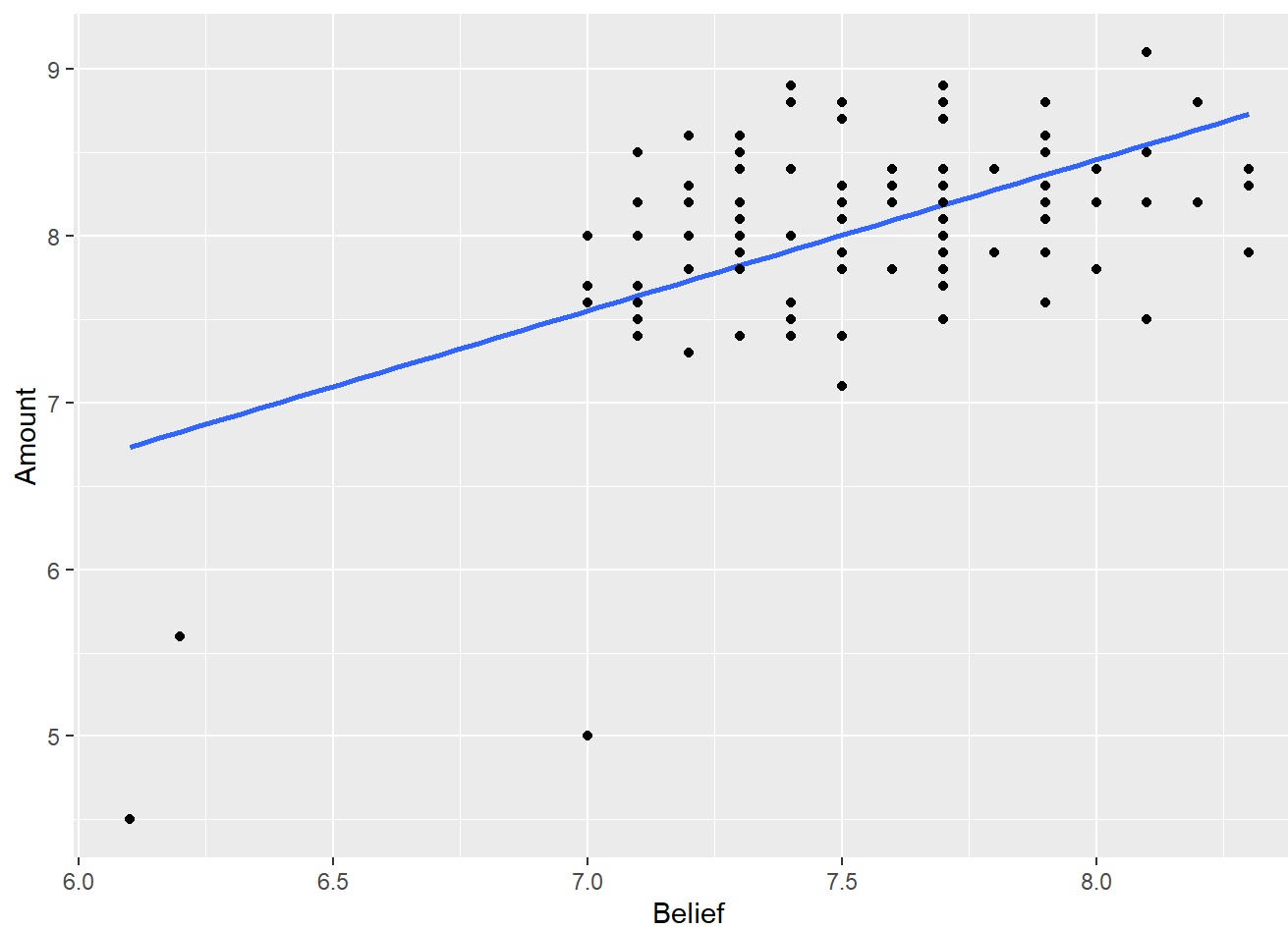
```
# Bivariate Scatterplots of Amount with Predictors  
ggplot(data = dat,  
       mapping = aes(y = Amount, x = Need)) +  
  geom_smooth(method = "lm", formula = y~x, se=FALSE) +  
  geom_point() +  
  labs(y = "Amount", x = "Need")
```



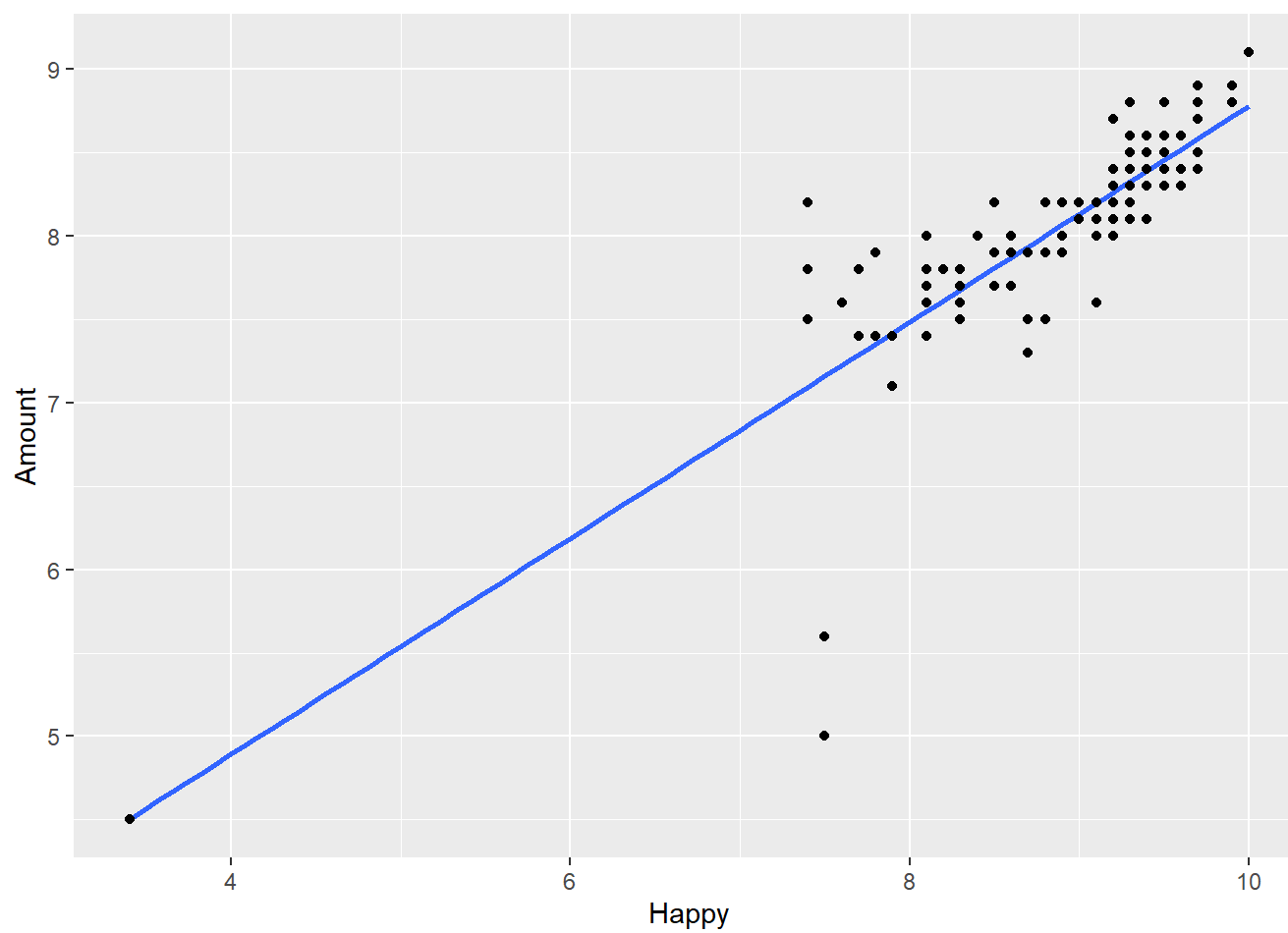
```
ggplot(data = dat,  
       mapping = aes(y = Amount, x = Interest)) +  
  geom_smooth(method = "lm", formula = y~x, se=FALSE) +  
  geom_point() +  
  labs(y = "Amount", x = "Interest")
```



```
ggplot(data = dat,  
       mapping = aes(y = Amount, x = Belief)) +  
  geom_smooth(method = "lm", formula = y~x, se=FALSE) +  
  geom_point() +  
  labs(y = "Amount", x = "Belief")
```



```
ggplot(data = dat,  
       mapping = aes(y = Amount, x = Happy)) +  
  geom_smooth(method = "lm", formula = y~x, se=FALSE) +  
  geom_point() +  
  labs(y = "Amount", x = "Happy")
```

Generate Correlations Among Variables

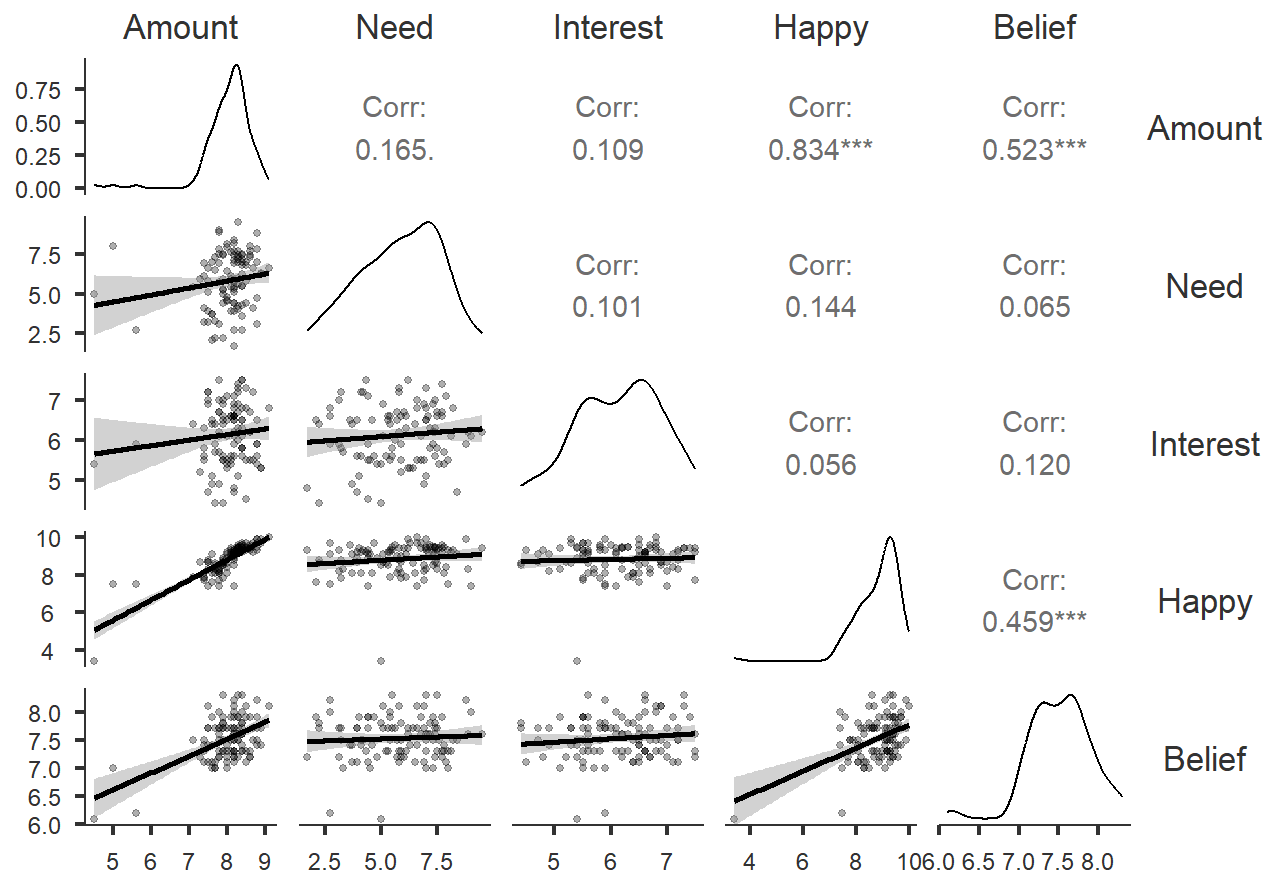
```
# `psych` Correlation analyses (Listwise Deletion)
cor_listwise <- corr.test(dat[2:6], use="complete")
print(cor_listwise, short=FALSE)
```

```
## Call:corr.test(x = dat[2:6], use = "complete")
## Correlation matrix
##           Amount Need Interest Happy Belief
## Amount      1.00 0.25      0.12 0.87  0.52
## Need        0.25 1.00      0.10 0.16  0.08
## Interest     0.12 0.10      1.00 0.08  0.15
## Happy        0.87 0.16      0.08 1.00  0.45
## Belief       0.52 0.08      0.15 0.45  1.00
## Sample Size
## [1] 102
## Probability values (Entries above the diagonal are adjusted for multiple tests.)
##           Amount Need Interest Happy Belief
## Amount      0.00 0.09      0.89 0.00  0.00
## Need        0.01 0.00      0.95 0.63  0.95
## Interest     0.22 0.32      0.00 0.95  0.68
## Happy        0.00 0.11      0.42 0.00  0.00
## Belief       0.00 0.41      0.14 0.00  0.00
##
## Confidence intervals based upon normal theory. To get bootstrapped values, try cor.ci
##           raw.lower raw.r raw.upper raw.p lower.adj upper.adj
## Amont-Need      0.05 0.25      0.42 0.01      -0.02      0.48
## Amont-Intrs     -0.07 0.12      0.31 0.22      -0.13      0.36
## Amont-Happy      0.81 0.87      0.91 0.00       0.78      0.92
## Amont-Belif      0.36 0.52      0.65 0.00       0.29      0.69
## Need-Intrs      -0.10 0.10      0.29 0.32      -0.14      0.33
## Need-Happy      -0.03 0.16      0.34 0.11      -0.10      0.40
## Need-Belif      -0.11 0.08      0.27 0.41      -0.14      0.30
## Intrs-Happy     -0.12 0.08      0.27 0.42      -0.12      0.27
## Intrs-Belif     -0.05 0.15      0.33 0.14      -0.11      0.39
## Happy-Belif      0.28 0.45      0.59 0.00       0.21      0.64
```

```
# `jmv` Correlation analyses with Plots
corrMatrix(dat[2:6],
  ci = TRUE,
  plots = TRUE,
  plotDens = TRUE,
  plotStats = TRUE)
```

```
##
## CORRELATION MATRIX
##
## Correlation Matrix
```

		Amount	Need	Interest	Happy	Belief
Amount	Pearson's r	—				
	df	—				
	p-value	—				
	95% CI Upper	—				
	95% CI Lower	—				
Need	Pearson's r	0.1645276	—			
	df	101	—			
	p-value	0.0967671	—			
	95% CI Upper	0.3470037	—			
	95% CI Lower	-0.0299506	—			
Interest	Pearson's r	0.1090415	0.1011541	—		
	df	101	100	—		
	p-value	0.2728962	0.3117232	—		
	95% CI Upper	0.2963133	0.2899256	—		
	95% CI Lower	-0.0863044	-0.0951934	—		
Happy	Pearson's r	0.8341407	0.1436484	0.0557898	—	
	df	103	102	102	—	
	p-value	< .0000001	0.1457294	0.5737694	—	
	95% CI Upper	0.8844102	0.3271850	0.2457377	—	
	95% CI Lower	0.7647387	-0.0503323	-0.1382842	—	
Belief	Pearson's r	0.5227888	0.0646370	0.1204032	0.4592751	—
	df	101	100	101	102	—
	p-value	< .0000001	0.5186476	0.2257197	0.0000009	—
	95% CI Upper	0.6505007	0.2558952	0.3067796	0.5988908	—
	95% CI Lower	0.3663263	-0.1314908	-0.0748659	0.2925645	—

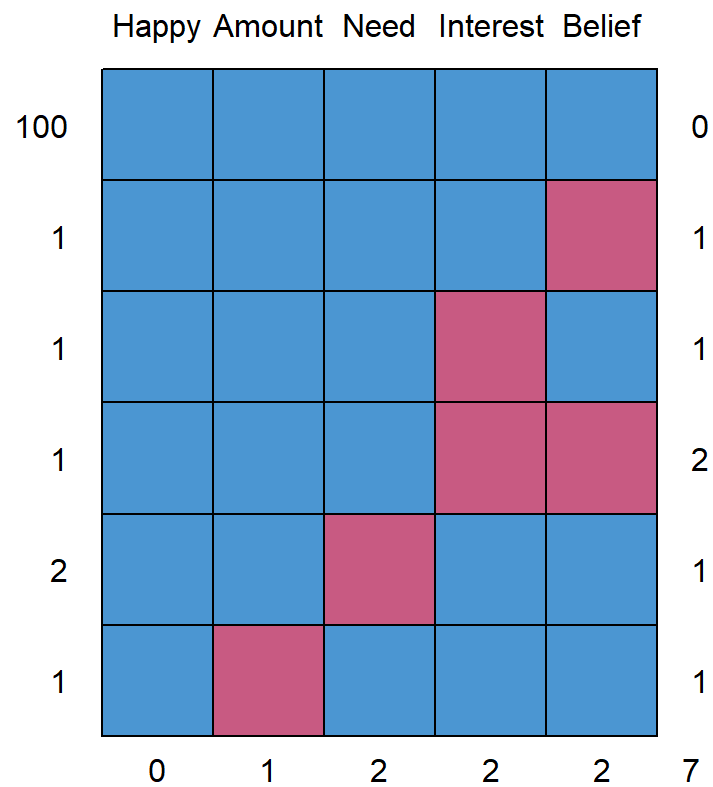


Assess Missing Data/Pattern

```
# Calculate Amount of Missing Data
sum(is.na(dat[2:6]))/prod(dim(dat[2:6]))
```

```
## [1] 0.01320755
```

```
# Assess Pattern of Missing Data
md.pattern(dat[2:6])
```



```
##      Happy Amount Need Interest Belief
## 100      1      1      1         1      1 0
## 1       1      1      1         1      0 1
## 1       1      1      1         0      1 1
## 1       1      1      1         0      0 2
## 2       1      1      0         1      1 1
## 1       1      0      1         1      1 1
##       0      1      2         2      2 7
```

Due to small amount of missing data we will use listwise deletion and proceed

List-wise deletion of missing data

```
# Listwise deletion results in 7 cases removed from data
# Will create new data object to compare analyses later
dat_no_NA <- na.omit(dat)
```

Assessment of Univariate Outliers

```
#Identify outliers
dat_no_NA[abs(scale(dat_no_NA$Belief)) > 3.29, ]
```

```
##      ID Amount Need Interest Happy Belief
## 1 100    4.5  5.0     5.4   3.4    6.1
## 6 105    5.6  2.7     5.9   7.5    6.2
```

```
dat_no_NA[abs(scale(dat_no_NA$Need)) > 3.29, ]
```

```
## [1] ID      Amount    Need      Interest Happy     Belief
## <0 rows> (or 0-length row.names)
```

```
dat_no_NA[abs(scale(dat_no_NA$Interest)) > 3.29, ]
```

```
## [1] ID      Amount    Need      Interest Happy     Belief
## <0 rows> (or 0-length row.names)
```

```
dat_no_NA[abs(scale(dat_no_NA$Happy)) > 3.29, ]
```

```
##      ID Amount Need Interest Happy Belief
## 1 100    4.5   5     5.4   3.4    6.1
```

```
dat_no_NA[abs(scale(dat_no_NA$Amount)) > 3.29, ]
```

```
##      ID Amount Need Interest Happy Belief
## 1 100    4.5  5.0     5.4   3.4    6.1
## 6 105    5.6  2.7     5.9   7.5    6.2
```

```
#Belief has 2 univariate outliers (IDs 100 and 105)
#Need has 0
#Interest has 0
#Happy has 1 univariate outlier (ID 100)
#Amount has 2 univariate outliers (IDs 100 and 105)
```

Remove Univariate Outliers

```
# Step needs to be conducted sequentially as shown below (can be expanded with larger datasets)
dat.no.uni1 <- dat_no_NA[!abs(scale(dat_no_NA$Belief)) > 3.29, ]
dat.no.uni2 <- dat.no.uni1[!abs(scale(dat.no.uni1$Need)) > 3.29, ]
dat.no.uni3 <- dat.no.uni2[!abs(scale(dat.no.uni2$Interest)) > 3.29, ]
dat.no.uni4 <- dat.no.uni3[!abs(scale(dat.no.uni3$Happy)) > 3.29, ]
dat_no_NA_UNI <- dat.no.uni4[!abs(scale(dat.no.uni4$Amount)) > 3.29, ]
```

Reassess Univariate Normality After Univariate Outlier Removal

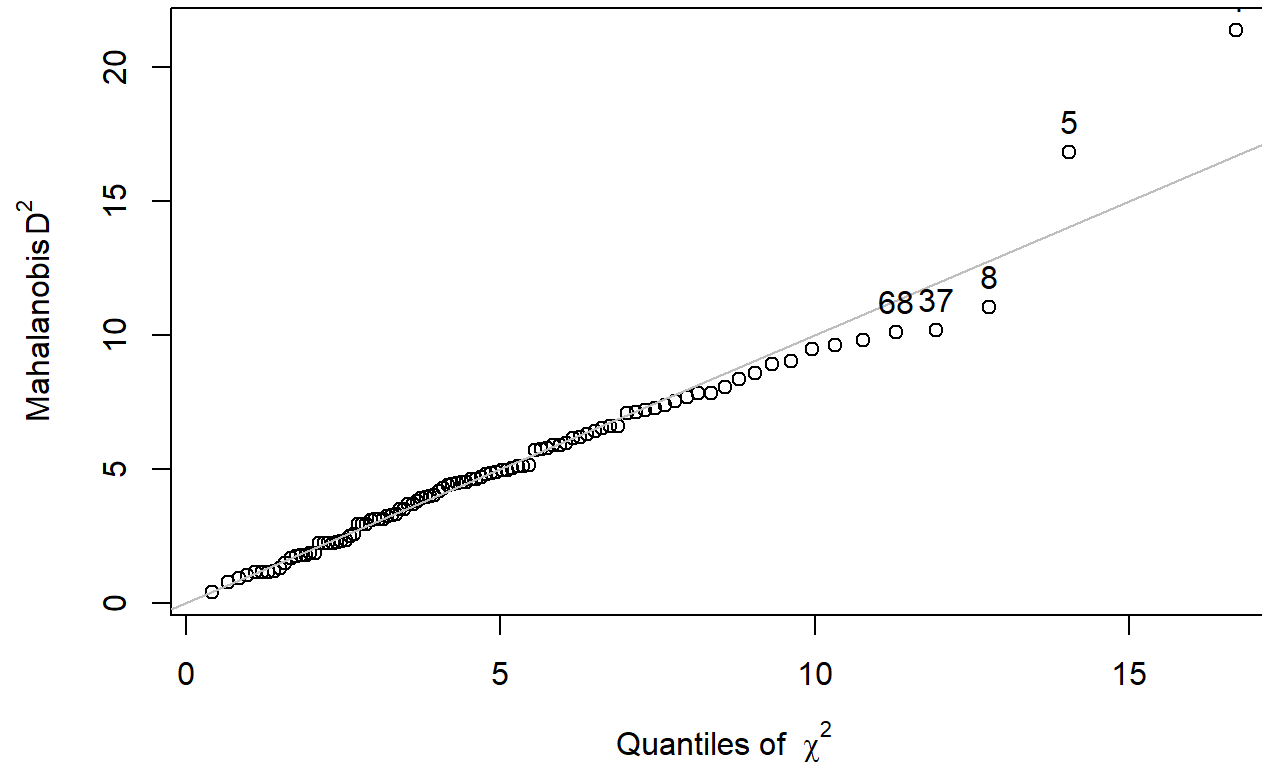
```
describe(dat_no_NA_UNI[2:6])
```

```
##          vars  n mean   sd median trimmed  mad min  max range  skew kurtosis
## Amount      1 98 8.12 0.41   8.20    8.12 0.44 7.1   9.1   2.0 -0.11   -0.49
## Need        2 98 5.82 1.77   5.90    5.89 2.08 1.7   9.5   7.8 -0.30   -0.71
## Interest    3 98 6.13 0.75   6.20    6.15 0.89 4.4   7.5   3.1 -0.28   -0.67
## Happy       4 98 8.93 0.63   9.10    8.98 0.59 7.4  10.0   2.6 -0.64   -0.49
## Belief      5 98 7.56 0.33   7.55    7.55 0.37 7.0   8.3   1.3  0.28   -0.68
##              se
## Amount    0.04
## Need      0.18
## Interest  0.08
## Happy     0.06
## Belief    0.03
```

Multivariate Outliers - Mahalanobis and Cook's Distances

```
# Mahalanobis Distance
dat_no_NA_UNI$mahal <- outlier(dat_no_NA_UNI[2:6])
```

Q-Q plot of Mahalanobis D^2 vs. quantiles of χ^2_{nvar}



```
dat_no_NA_UNI[abs(scale(dat_no_NA_UNI$mahal)) > 3.29, ]
```

```
##      ID Amount Need Interest Happy Belief      mahal
## 4 103    8.2  4.4     5.9   7.4    7.7 21.35014
## 5 104    7.8  9.0     6.1   7.4    8.0 16.80466
```

2 Multivariate Outliers Identified (IDs 103 and 104)

#Cook's Distance

```
lm<-lm(Amount~Belief + Need + Interest + Happy, data = dat_no_NA_UNI)
```

#Generate Cook's Distance

```
dat_no_NA_UNI$cooks <- cooks.distance(lm)
```

```
dat_no_NA_UNI[abs(scale(dat_no_NA_UNI$cooks)) > 3.29, ]
```



```
##      ID Amount Need Interest Happy Belief      mahal      cooks
## 4 103      8.2  4.4      5.9   7.4      7.7 21.35014 0.2715843
```

```
## 1 Multivariate Outlier Identified (ID 103)
```

```
#Remove multivariate outliers
```

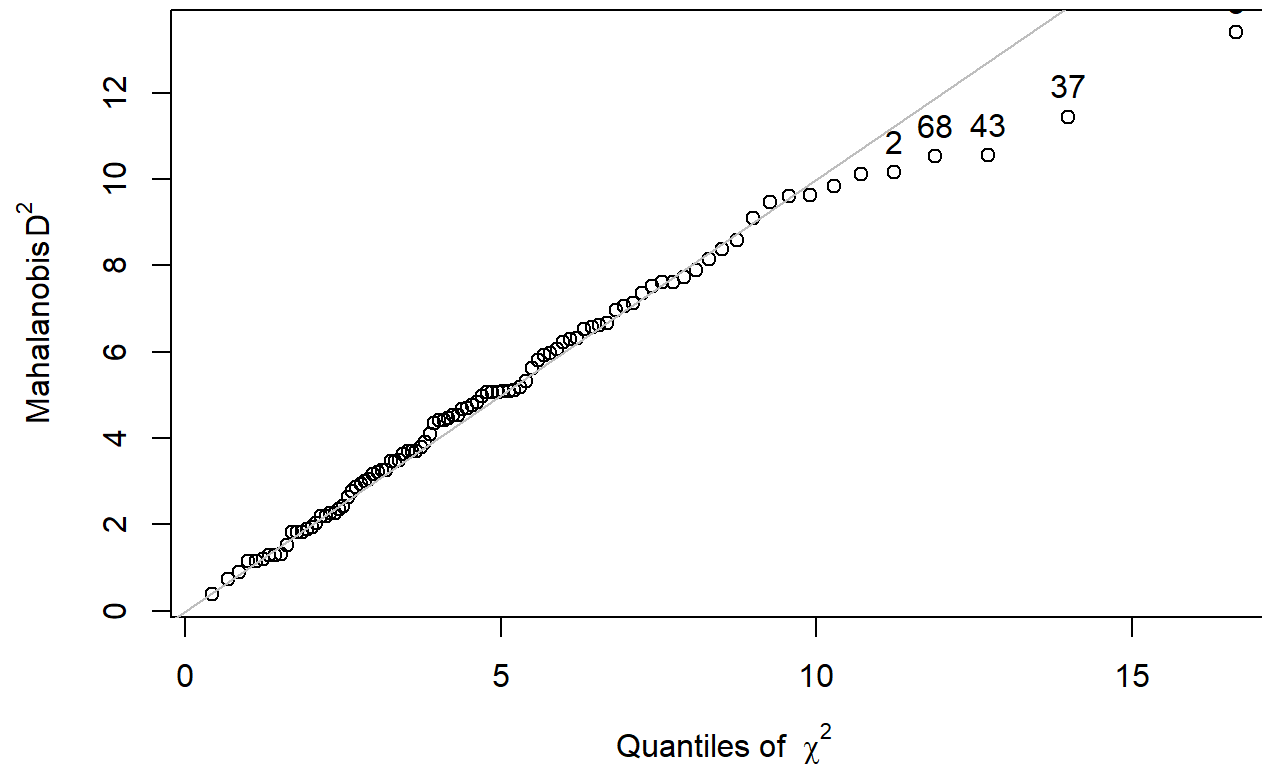
```
dat_no_NA_UMO <- dat_no_NA_UNI[!abs(scale(dat_no_NA_UNI$mahal)) > 3.29,]
```

Multivariate Outlier Iteration Checking Code

```
# Mahalanobis Distance
```

```
dat_no_NA_UMO$mahal <- outlier(dat_no_NA_UMO[2:6])
```

Q-Q plot of Mahalanobis D^2 vs. quantiles of χ^2_{nvar}



```
dat_no_NA_UMO[abs(scale(dat_no_NA_UMO$mahal)) > 3.29, ]
```

```
## [1] ID      Amount  Need    Interest Happy   Belief   mahal    cooks
## <0 rows> (or 0-length row.names)
```

```
# 0 Multivariate Outliers Identified
```

```
#Cook's Distance
```

```
lm<-lm(Amount~Belief + Need + Interest + Happy, data = dat_no_NA_UMO)
```

```
#Generate Cook's Distance
```

```
dat_no_NA_UMO$cooks <- cooks.distance(lm)
```

```
dat_no_NA_UMO[abs(scale(dat_no_NA_UMO$cooks)) > 3.29, ]
```

```
##      ID Amount Need Interest Happy Belief   mahal     cooks
## 43 142    7.5  3.2      7.2   8.8    7.7 10.56072 0.07604036
```

```
## Multivariate Outliers Identified IDs #1[142], #2[140,150], #3[111,136], #4[107,138],
## #5[101,173], #6[113,131,162], #7[102,205], #8[197], #9[187], #10[166]
```

```
#Remove multivariate outliers
```

```
dat_no_NA_UMO <- dat_no_NA_UMO[!abs(scale(dat_no_NA_UMO$cooks)) > 3.29,]
```

Rename Data Object for Additional Assumption Checks

```
dat_final <- dat_no_NA_UMO
```

```
# Remove extra datasets
```

```
rm(dat.no.uni1,
   dat.no.uni2,
   dat.no.uni3,
   dat.no.uni4,
   dat_no_NA,
   dat_no_NA_UMO,
   dat_no_NA_UNI)
```

Multivariate Normality

```
# Mardia's Test of Multivariate Normality
mardia <- mvn(dat_final[2:6],
             mvnTest = "mardia",
             desc = FALSE)

# Henze-Zirkler's Test of Multivariate Normality
hz <- mvn(dat_final[2:6],
          mvnTest = "hz",
          desc = FALSE)

# Energy Test of Multivariate Normality
energy <- mvn(dat_final[2:6],
              mvnTest = "energy",
              desc = FALSE)

# Doornik-Hansen's Test of Multivariate Normality
dh <- mvn(dat_final[2:6],
          mvnTest = "dh",
          desc = FALSE)

mardia$multivariateNormality
```

##	Test	Statistic	p value	Result
## 1	Mardia Skewness	33.9618014701476	0.518086248634635	YES
## 2	Mardia Kurtosis	-1.17166969389339	0.241329695001204	YES
## 3	MVN	<NA>	<NA>	YES

```
hz$multivariateNormality
```

##	Test	HZ	p value	MVN
## 1	Henze-Zirkler	1.127156	0.000575138	NO

```
energy$multivariateNormality
```

##	Test	Statistic	p value	MVN
## 1	E-statistic	1.447145	0.001	NO

```
dh$multivariateNormality
```

```
##           Test           E df           p value MVN
## 1 Doornik-Hansen 53.28651 10 6.587082e-08 NO
```

Homoscedasticity

```
bplm<-lm(Amount~Belief+Need+Interest+Happy, data = dat_final[2:6])
```

```
#Breusch-Pagan test
bptest(bplm, studentize=FALSE)
```

```
##
## Breusch-Pagan test
##
## data:  bplm
## BP = 5.7009, df = 4, p-value = 0.2226
```

```
bptest(bplm, studentize=TRUE)
```

```
##
## studentized Breusch-Pagan test
##
## data:  bplm
## BP = 4.0224, df = 4, p-value = 0.403
```

Multicollinearity

```
#VIF and Tolerance
collin <- linReg(data = dat_final,
                 dep = Amount,
                 cov = c(Belief, Need, Interest, Happy),
                 blocks = list(list('Belief','Need','Interest','Happy')),
                 r=FALSE,
                 r2=FALSE,
                 collin = TRUE)

collin
```

```
##
## LINEAR REGRESSION
##
## MODEL SPECIFIC RESULTS
##
## MODEL 1
##
## Model Coefficients - Amount
##
```

	Predictor	Estimate	SE	t	p
##	Intercept	2.45654948	0.54450316	4.5115431	0.0000194
##	Belief	0.01671975	0.06935396	0.2410785	0.8100427
##	Need	0.01429280	0.01285053	1.1122346	0.2689994
##	Interest	0.04465911	0.02928590	1.5249354	0.1307817
##	Happy	0.57908575	0.03854724	15.0227545	< .0000001

```
##
##
## ASSUMPTION CHECKS
##
## Collinearity Statistics
##
```

		VIF	Tolerance
##	Belief	1.118742	0.8938611
##	Need	1.054752	0.9480898
##	Interest	1.031024	0.9699099
##	Happy	1.139251	0.8777698

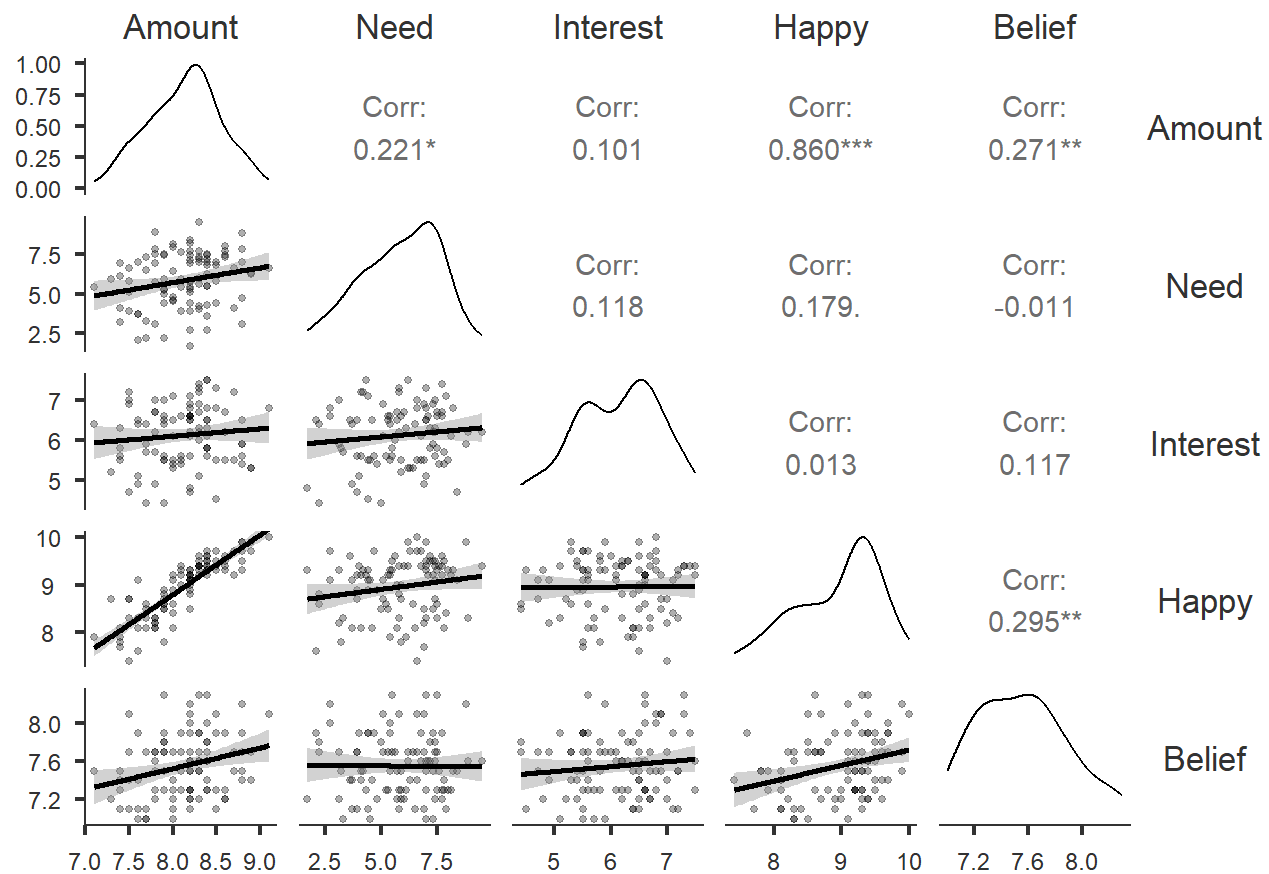
```
##
```

```
corrMatrix(dat_final[2:6],
           ci = TRUE)
```

```
##
## CORRELATION MATRIX
##
## Correlation Matrix
```

		Amount	Need	Interest	Happy	Belief
Amount	Pearson's r	—				
	df	—				
	p-value	—				
	95% CI Upper	—				
	95% CI Lower	—				
Need	Pearson's r	0.2206200	—			
	df	93	—			
	p-value	0.0316806	—			
	95% CI Upper	0.4041909	—			
	95% CI Lower	0.0199647	—			
Interest	Pearson's r	0.1010147	0.1179069	—		
	df	93	93	—		
	p-value	0.3300415	0.2551309	—		
	95% CI Upper	0.2965209	0.3120349	—		
	95% CI Lower	-0.1026174	-0.0856720	—		
Happy	Pearson's r	0.8597107	0.1789614	0.0128636	—	
	df	93	93	93	—	
	p-value	< .0000001	0.0826971	0.9015336	—	
	95% CI Upper	0.9045274	0.3672580	0.2138522	—	
	95% CI Lower	0.7961083	-0.0234266	-0.1891698	—	
Belief	Pearson's r	0.2714395	-0.0106415	0.1172407	0.2951672	—
	df	93	93	93	93	—
	p-value	0.0077957	0.9184786	0.2578443	0.0036862	—
	95% CI Upper	0.4484493	0.1913118	0.3114251	0.4688206	—
	95% CI Lower	0.0739416	-0.2117304	-0.0863425	0.0995460	—

```
corrMatrix(dat_final[2:6],
  ci = TRUE,
  plots = TRUE,
  plotDens = TRUE,
  plotStats = TRUE)$plot
```



Multiple Regression Cleaned vs. Uncleaned Data

Multiple regression not clean

```
linReg(data = dat,  
      dep = 'Amount',  
      covs = c('Belief', 'Need', 'Interest', 'Happy'),  
      blocks = list(list('Belief', 'Need', 'Interest', 'Happy')),  
      r2Adj = TRUE,  
      modelTest=TRUE,  
      ci = TRUE,  
      stdEst = TRUE,  
      ciStdEst = TRUE)
```



```
#Multiple regression clean
linReg(data = dat_final,
      dep = 'Amount',
      covs = c('Belief', 'Need', 'Interest', 'Happy'),
      blocks = list(list('Belief', 'Need', 'Interest', 'Happy')),
      r2Adj = TRUE,
      modelTest=TRUE,
      ci = TRUE,
      stdEst = TRUE,
      ciStdEst = TRUE)
```

##									
## LINEAR REGRESSION									
##									
## Model Fit Measures									
##									
##	Model	R	R ²	Adjusted R ²	F	df1	df2	p	
##									
##	1	0.8664207	0.7506848	0.7396041	67.74719	4	90	< .0000001	
##									
## Note. Models estimated using sample size of N=95									
##									
##									
## MODEL SPECIFIC RESULTS									
##									
## MODEL 1									
##									
## Model Coefficients - Amount									
##									
##	Predictor	Estimate	SE	Lower	Upper	t	p	Stand. Estimate	L
ower									
##									
##									
##									
##	Intercept	2.45654948	0.54450316	1.37479892	3.53830004	4.5115431	0.0000194		
##	Belief	0.01671975	0.06935396	-0.12106399	0.15450348	0.2410785	0.8100427	0.01342074	-
	0.09717665	0.1240181							
##	Need	0.01429280	0.01285053	-0.01123701	0.03982262	1.1122346	0.2689994	0.06012081	-
	0.04726704	0.1675087							
##	Interest	0.04465911	0.02928590	-0.01352245	0.10284066	1.5249354	0.1307817	0.08149649	-
	0.02467653	0.1876695							
##	Happy	0.57908575	0.03854724	0.50250492	0.65566657	15.0227545	< .0000001	0.84394165	
	0.73233513	0.9555482							
##									
##									

Model Pruning and Model Comparisons/Parsimonious Models

```
#Simple Bivariate Regression (Amount ~ Belief)
lm1 <- lm(Amount~Belief, data = dat_final)

#Simple Standardized Bivariate Regression (Amount ~ Belief)
zlm1 <- lm(scale(Amount) ~ scale(Belief), data = dat_final)

#Call for standardized regression coefficients/summary
summary(zlm1)
```

```
##
## Call:
## lm(formula = scale(Amount) ~ scale(Belief), data = dat_final)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.4444 -0.6748  0.1282  0.5759  1.9865
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -7.875e-16  9.928e-02   0.00   1.0000
## scale(Belief)  2.714e-01  9.980e-02   2.72   0.0078 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.9676 on 93 degrees of freedom
## Multiple R-squared:  0.07368,    Adjusted R-squared:  0.06372
## F-statistic: 7.397 on 1 and 93 DF,  p-value: 0.007796
```

```
#Call for standardized regression confidence intervals
confint(zlm1)
```

```
##              2.5 %    97.5 %
## (Intercept) -0.19714112 0.1971411
## scale(Belief) 0.07325257 0.4696265
```

#Multiple Linear Regression

```
lm2 <- lm(Amount~Belief+Need, data = dat_final)
```

#Standardized Multiple Regression

```
zlm2 <- lm(scale(Amount) ~ scale(Belief) + scale(Need), data = dat_final)
```

#Call for standardized MR coefficients/summary

```
summary(zlm2)
```

```
##
```

```
## Call:
```

```
## lm(formula = scale(Amount) ~ scale(Belief) + scale(Need), data = dat_final)
```

```
##
```

```
## Residuals:
```

```
##      Min       1Q   Median       3Q      Max
```

```
## -2.38828 -0.65393  0.03925  0.61153  1.92773
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error t value Pr(>|t|)
```

```
## (Intercept)  -7.334e-16  9.708e-02   0.000  1.00000
```

```
## scale(Belief)  2.738e-01  9.760e-02   2.805  0.00613 **
```

```
## scale(Need)    2.235e-01  9.760e-02   2.290  0.02430 *
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
```

```
## Residual standard error: 0.9463 on 92 degrees of freedom
```

```
## Multiple R-squared:  0.1236, Adjusted R-squared:  0.1046
```

```
## F-statistic: 6.49 on 2 and 92 DF,  p-value: 0.002309
```

```
summary(lm2)
```

```
##
## Call:
## lm(formula = Amount ~ Belief + Need, data = dat_final)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.98845 -0.27064  0.01625  0.25310  0.79784
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  5.24303     0.93098   5.632 1.93e-07 ***
## Belief       0.34113     0.12160   2.805  0.00613 **
## Need        0.05314     0.02320   2.290  0.02430 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3916 on 92 degrees of freedom
## Multiple R-squared:  0.1236, Adjusted R-squared:  0.1046
## F-statistic:  6.49 on 2 and 92 DF,  p-value: 0.002309
```

```
#Call for standardized MR confidence intervals
confint(zlm2)
```

```
##              2.5 %    97.5 %
## (Intercept) -0.19281789 0.1928179
## scale(Belief)  0.07996649 0.4676701
## scale(Need)    0.02968202 0.4173856
```

```
confint(lm2)
```

```
##              2.5 %    97.5 %
## (Intercept) 3.394036695 7.09203114
## Belief      0.099623346 0.58262973
## Need        0.007056446 0.09922702
```

Direct Model Comparison

```
#Model Comparison
anova(lm1, lm2)
```

```
## Analysis of Variance Table
##
## Model 1: Amount ~ Belief
## Model 2: Amount ~ Belief + Need
##   Res.Df    RSS Df Sum of Sq    F Pr(>F)
## 1      93 14.915
## 2      92 14.111   1   0.80446 5.245 0.0243 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
anova(zlm1, zlm2)
```

```
## Analysis of Variance Table
##
## Model 1: scale(Amount) ~ scale(Belief)
## Model 2: scale(Amount) ~ scale(Belief) + scale(Need)
##   Res.Df    RSS Df Sum of Sq    F Pr(>F)
## 1      93 87.074
## 2      92 82.378   1   4.6964 5.245 0.0243 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Run and Visualize Final Model

```
linReg(data = dat_final,
       dep = 'Amount',
       covs = c('Belief', 'Need'),
       blocks = list(list('Belief', 'Need')),
       r2Adj = TRUE,
       modelTest=TRUE,
       ci = TRUE,
       stdEst = TRUE,
       ciStdEst = TRUE)
```

```
##
## LINEAR REGRESSION
##
## Model Fit Measures
##
```

Model	R	R ²	Adjusted R ²	F	df1	df2	p
1	0.3516264	0.1236411	0.1045898	6.489912	2	92	0.0023087

```
##
## Note. Models estimated using sample size of N=95
##
## MODEL SPECIFIC RESULTS
##
## MODEL 1
##
## Model Coefficients - Amount
##
```

Predictor	Estimate	SE	Lower	Upper	t	p	Stand. Estimate	Lower	Upper
Intercept	5.24303392	0.93097526	3.394036695	7.09203114	5.631765	0.0000002			
Belief	0.34112654	0.12159753	0.099623346	0.58262973	2.805374	0.0061327	0.2738183	0.07	0.996649
Need	0.05314174	0.02320407	0.007056446	0.09922702	2.290190	0.0242958	0.2235338	0.02	0.968202

```
set.seed(13)
apa.reg.boot.table(lm2,
  number.samples = 1000)
```

```
##
##
## apa.reg.boot.table is a beta version.
## Block 1: Generating 1000 bootstrap samples
```



```
## Warning: The `x` argument of `as_tibble.matrix()` must have unique column names if
## `.name_repair` is omitted as of tibble 2.0.0.
## i Using compatibility `.name_repair`.
## i The deprecated feature was likely used in the apaTables package.
## Please report the issue at <https://github.com/dstanley4/apaTables/issues>.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

```
## Bootstrap for Delta RSQ in progress
```

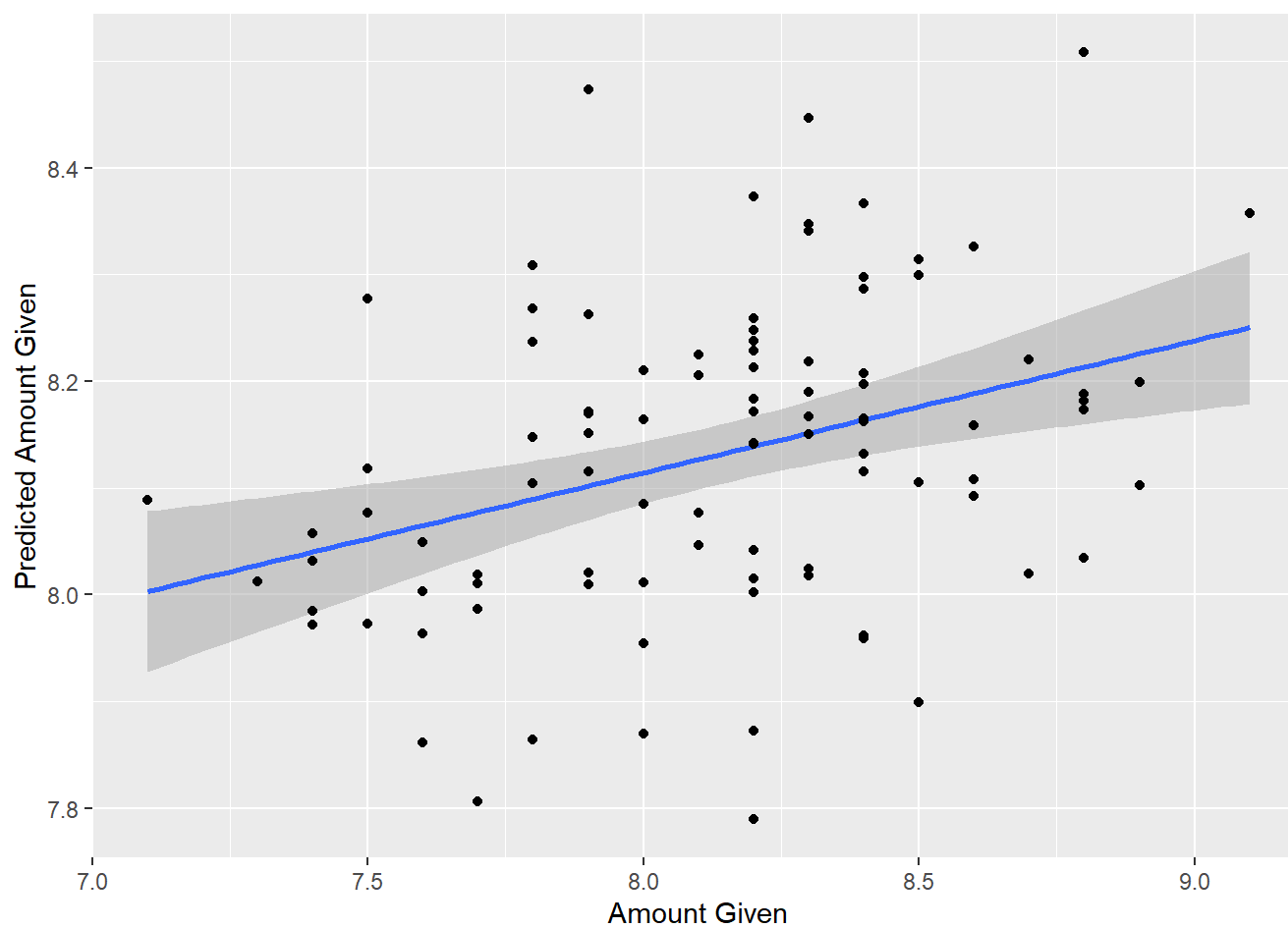
```
##
##
## Regression results using Amount as the criterion
##
##
## Predictor      b      b_95%_CI beta  beta_95%_CI sr2 sr2_95%_CI      r
## (Intercept) 5.24** [3.44, 6.97]
## Belief 0.34** [0.12, 0.57] 0.27 [0.10, 0.44] .07 [.01, .19] .27**
## Need 0.05* [0.01, 0.09] 0.22 [0.04, 0.38] .05 [.00, .15] .22*
##
##
##
## Fit
##
##
##
## R2 = .124**
## 95% CI[.04,.26]
##
##
## Note. A significant b-weight indicates the beta-weight and semi-partial correlation are also significant.
## b represents unstandardized regression weights. beta indicates the standardized regression weights.
## sr2 represents the semi-partial correlation squared. r represents the zero-order correlation.
## Square brackets are used to enclose the lower and upper limits of a confidence interval.
## * indicates p < .05. ** indicates p < .01.
##
```

```
apa.cor.table(dat_final[2:6])
```

```
##
##
## Means, standard deviations, and correlations with confidence intervals
##
##
## Variable      M      SD   1          2          3          4
## 1. Amount     8.13 0.41
##
## 2. Need       5.83 1.74 .22*
##                [.02, .40]
##
## 3. Interest   6.12 0.76 .10          .12
##                [-.10, .30] [-.09, .31]
##
## 4. Happy      8.96 0.60 .86**          .18          .01
##                [.80, .90] [-.02, .37] [-.19, .21]
##
## 5. Belief     7.56 0.33 .27**          -.01          .12          .30**
##                [.07, .45] [-.21, .19] [-.09, .31] [.10, .47]
##
##
## Note. M and SD are used to represent mean and standard deviation, respectively.
## Values in square brackets indicate the 95% confidence interval.
## The confidence interval is a plausible range of population correlations
## that could have caused the sample correlation (Cumming, 2014).
## * indicates  $p < .05$ . ** indicates  $p < .01$ .
##
```

```
# create predicted values from three predictors and save in object
dat_final$predictedF <- fitted(lm2)
dat_final$residuals <- resid(lm2)

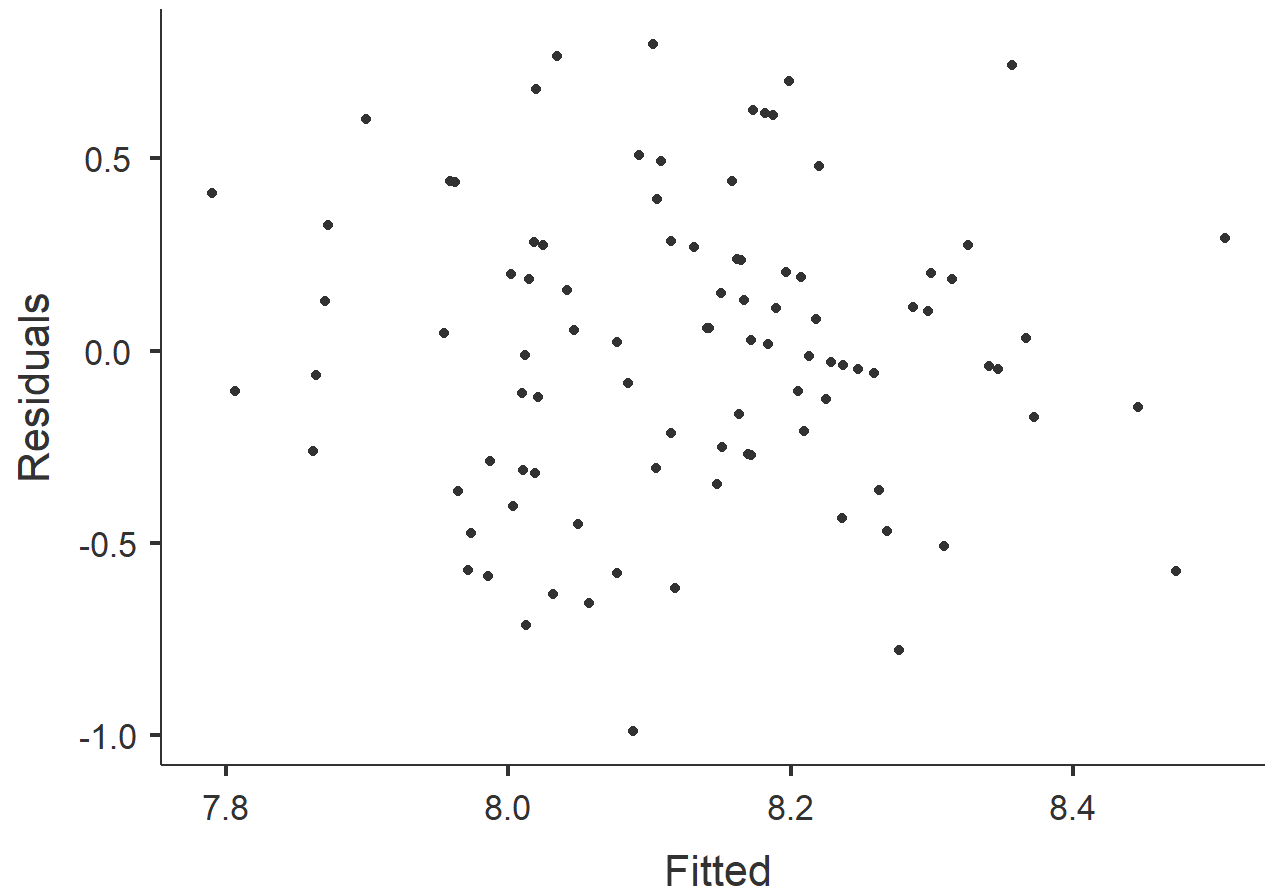
# plot predicted line
ggplot(data = dat_final,
       mapping = aes(x = Amount, y = predictedF)) +
  geom_smooth(method = "lm", formula = y ~ x) +
  geom_point() +
  labs(x = "Amount Given", y = "Predicted Amount Given")
```

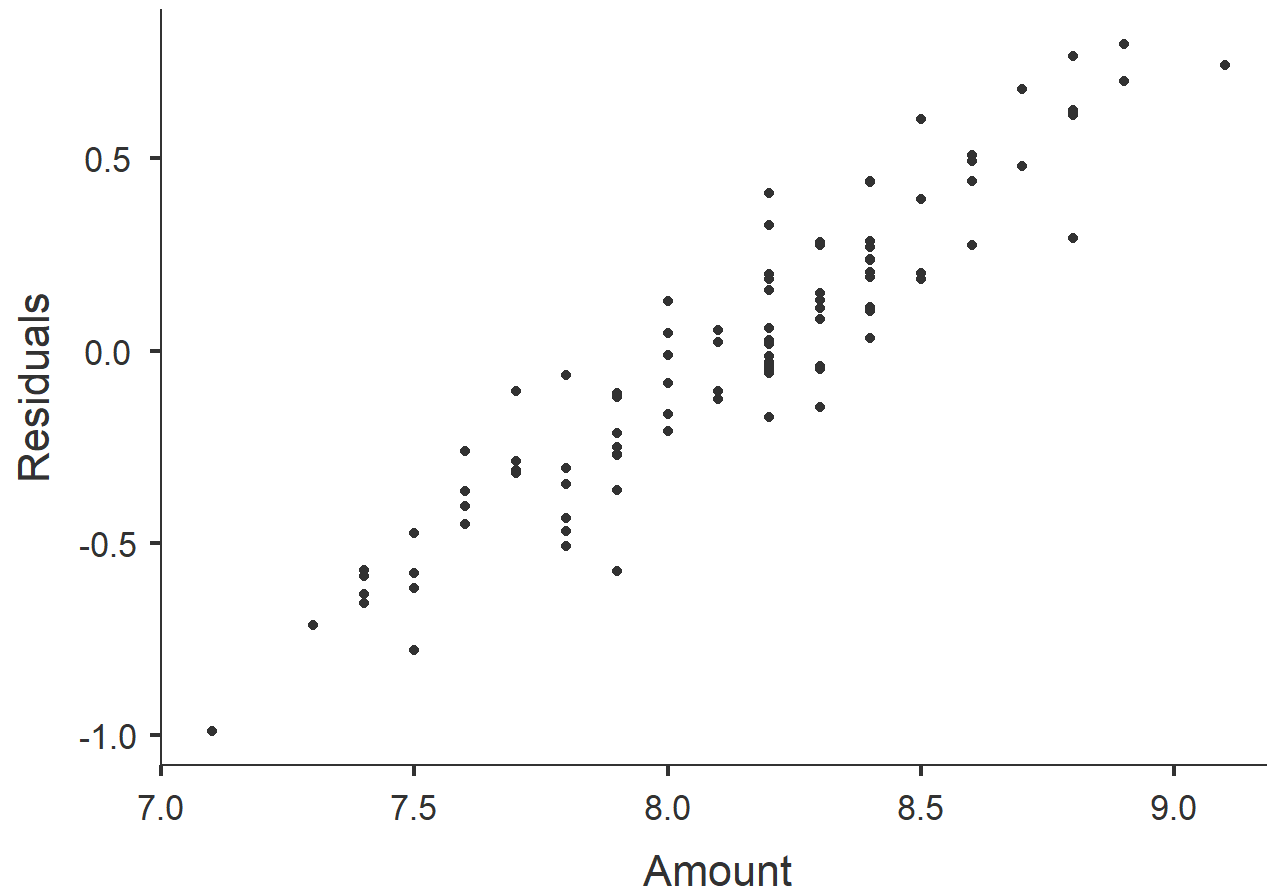


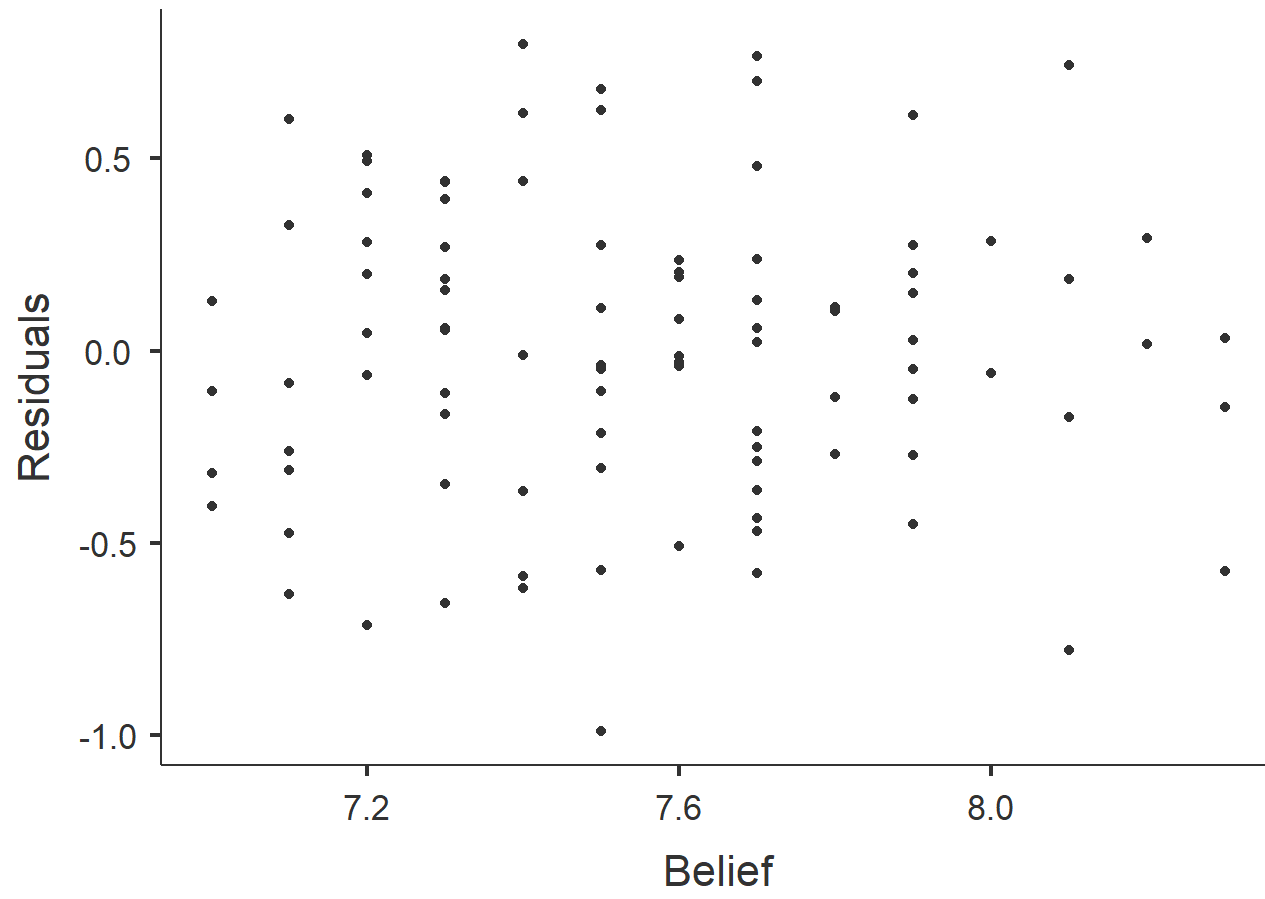
Checking residual plots

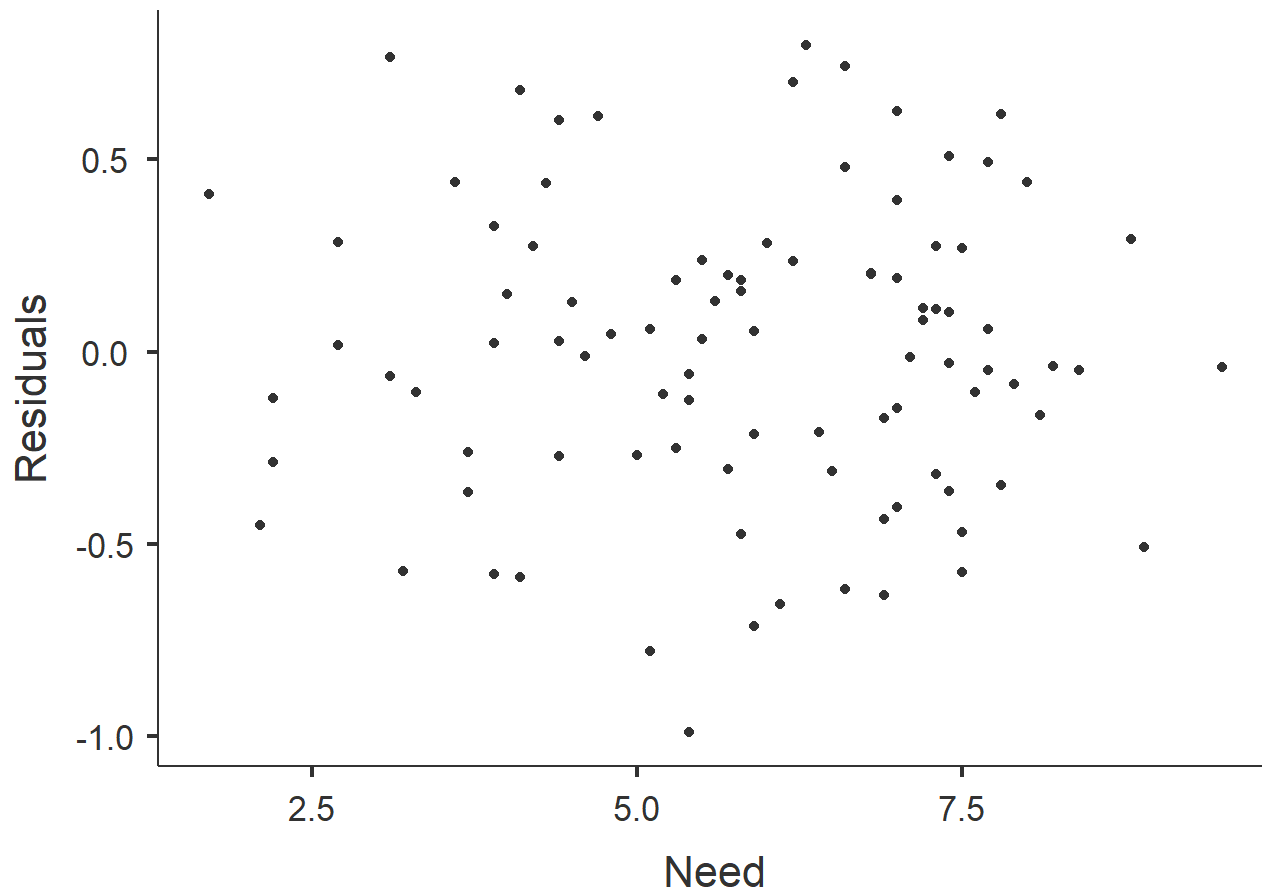
```
linReg(data = dat_final,  
       dep = Amount,  
       covs = vars(Belief,Need),  
       blocks = list(list('Belief','Need')),  
       modelTest = FALSE,  
       r=FALSE,  
       r2=FALSE,  
       resPlots=TRUE)
```

```
##
## LINEAR REGRESSION
##
## MODEL SPECIFIC RESULTS
##
## MODEL 1
##
## Model Coefficients - Amount
##
## | Predictor | Estimate | SE | t | p |
## |-----|-----|-----|-----|-----|
## | Intercept | 5.24303392 | 0.93097526 | 5.631765 | 0.0000002 |
## | Belief | 0.34112654 | 0.12159753 | 2.805374 | 0.0061327 |
## | Need | 0.05314174 | 0.02320407 | 2.290190 | 0.0242958 |
## |-----|-----|-----|-----|-----|
##
##
##
## ASSUMPTION CHECKS
```

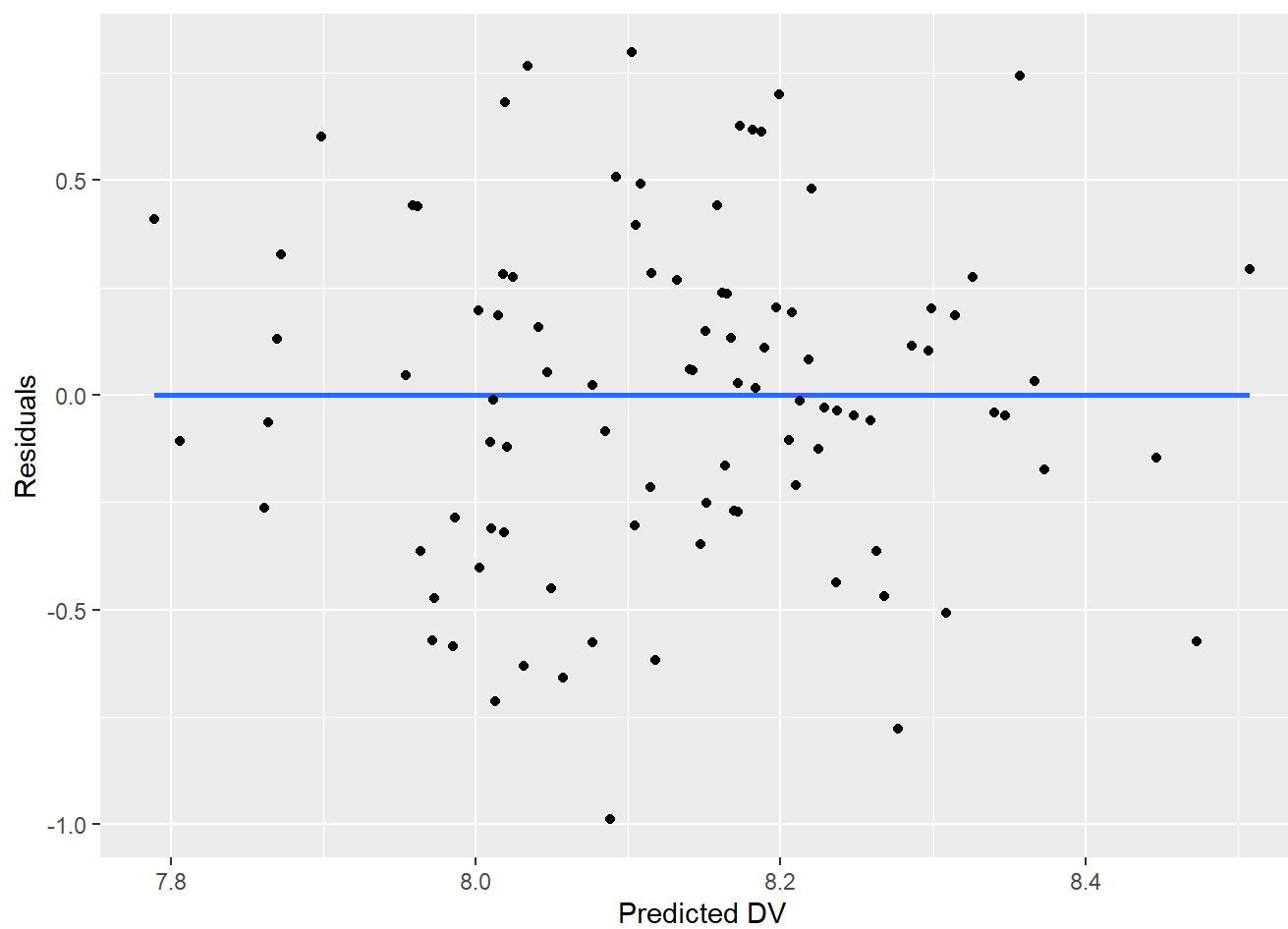




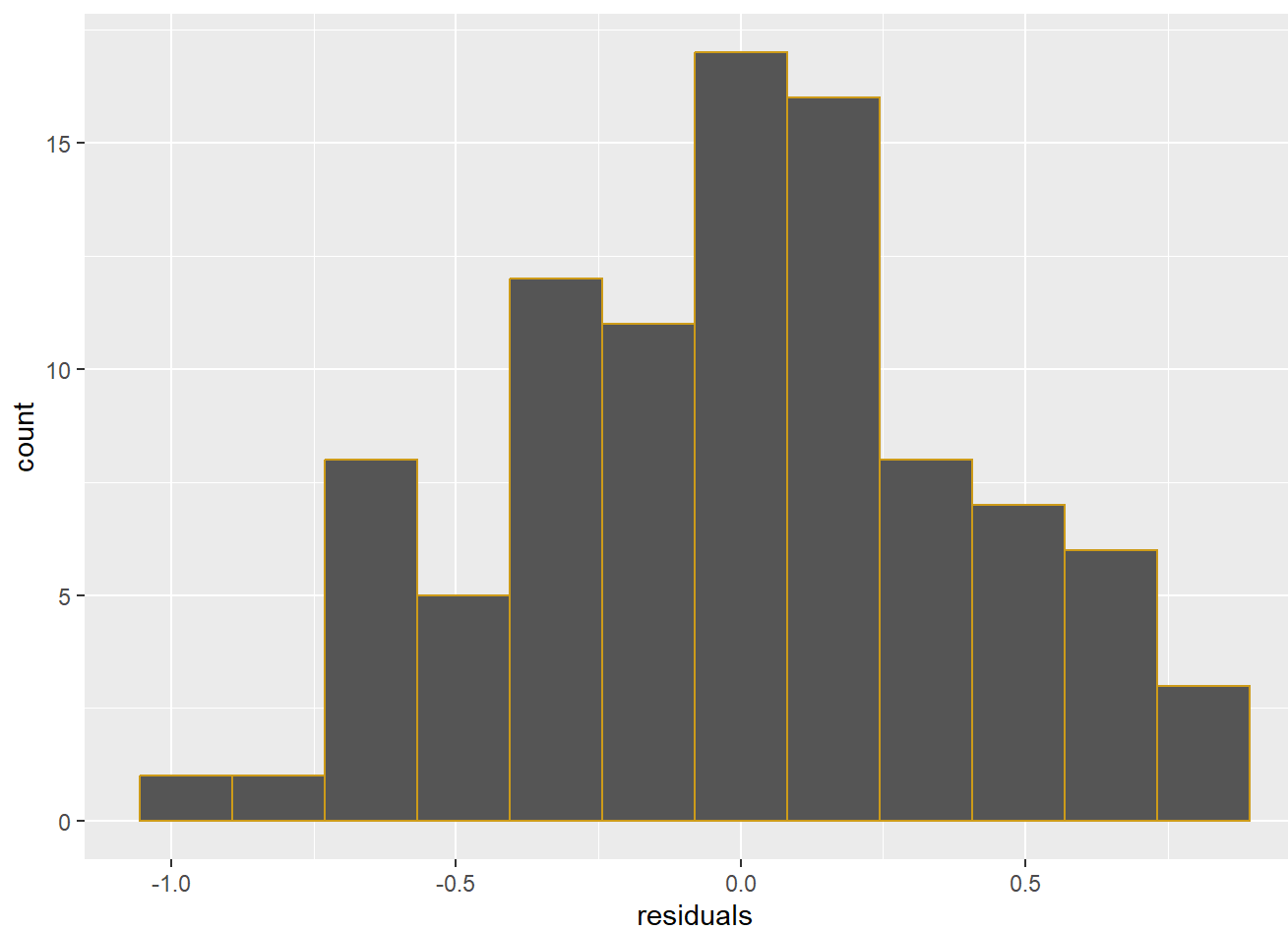




```
ggplot(data = dat_final,  
       mapping = aes(x = predictedF, y = residuals)) +  
  geom_smooth(method = "lm", formula = y ~ x, se = FALSE) +  
  geom_point() +  
  labs(x = "Predicted DV", y = "Residuals")
```

```
ggplot(data = dat_final,  
       mapping = aes(x = residuals)) +  
  geom_histogram(bins = 12, color = "goldenrod3")
```



```
describe(dat_final$residuals)
```

```
##      vars  n mean   sd median trimmed  mad   min max range  skew kurtosis   se
## X1      1 95    0 0.39  0.02      0 0.39 -0.99 0.8  1.79 -0.06   -0.49 0.04
```