

# CorrelCon: Introduction to R

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## 1 Let's get started...

First, we read in the data from an `rda` file (typical for R). Of course, also other data format (e.g., Excel sheets, text files) can be read in. In RStudio, you can also use the **Import Dataset** button in the Environment window. It guides you through, and you can later paste the code into your script.

```
> load("student_pisa.rda")
```

We get a first overview.

```
> # first six rows
> head(dat)
```

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20	X21
743	1	0	0	1	1	1	1	0	0	1	1	1	0	0	0	1	0	1	0	0	1
801	0	1	0	1	1	1	0	0	0	1	0	1	1	0	1	0	0	1	0	1	0
3747	0	0	0	1	0	1	0	0	1	1	1	1	0	0	1	1	0	0	0	0	0
4273	0	0	1	1	1	0	1	0	1	0	0	1	0	0	1	1	1	0	0	0	1
4366	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1	1	1	1	0	1	0
5177	1	0	0	1	0	1	1	1	0	1	0	1	0	0	1	1	0	1	0	1	1

	X22	X23	X24	X25	X26	X27	X28	X29	X30	X31	X32	X33	X34	X35	X36	X37	X38	X39
743	1	1	1	0	0	0	0	0	0	0	1	1	1	1	1	1	1	0
801	1	1	1	0	1	1	0	0	1	0	1	1	0	0	1	1	1	0
3747	0	0	1	1	0	0	1	1	1	0	0	1	1	1	1	1	1	0
4273	1	0	1	1	0	0	1	1	0	0	0	1	1	1	1	0	1	1
4366	1	1	1	1	1	0	0	1	1	1	1	0	1	1	1	1	1	1
5177	0	1	0	0	0	1	1	0	1	0	1	1	1	1	1	0	1	1

	X40	X41	X42	X43	X44	X45	gender	age	semester	elite	spon
743	1	0	1	1	1	1	female	21		3	no 1-3/month
801	1	1	1	1	1	1	male	20		1	no 4-5/week
3747	1	0	1	1	1	1	female	25		9	no 1-3/month
4273	1	1	1	0	1	1	male	27		10	no never
4366	1	0	0	1	1	1	male	24		8	no 1/week
5177	1	1	1	0	1	1	male	20		1	yes 1-3/month

```
> # last six rows
> tail(dat)
```

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20
690775	0	1	0	1	1	1	1	1	0	0	0	0	1	0	0	0	1	1	0	0
691336	1	1	0	1	1	1	1	0	0	1	0	1	0	0	1	0	1	1	0	1
691506	0	1	1	1	1	0	1	1	1	0	0	1	0	0	1	1	0	0	1	0
691527	0	1	0	1	0	1	1	0	0	1	1	1	0	0	1	1	1	0	1	1
691849	0	0	0	1	1	1	1	0	0	0	0	0	0	0	1	0	1	0	0	0
691972	1	1	0	1	1	1	1	1	0	1	1	1	0	0	0	1	0	1	0	0

	X21	X22	X23	X24	X25	X26	X27	X28	X29	X30	X31	X32	X33	X34	X35	X36	X37	X38
690775	0	1	1	1	1	0	1	0	0	1	1	0	1	0	1	0	1	0
691336	1	1	0	0	0	1	1	0	0	1	1	1	1	1	1	1	1	1
691506	1	1	1	1	1	1	1	1	0	1	1	1	0	1	1	1	1	1
691527	1	1	1	1	1	1	1	0	0	0	1	1	1	0	1	1	1	1
691849	1	1	1	1	0	0	1	1	1	0	0	0	1	1	1	1	0	1
691972	0	1	0	1	0	1	1	1	0	0	0	1	1	1	1	1	1	1

	X39	X40	X41	X42	X43	X44	X45	gender	age	semester	elite	spon
690775	0	0	1	1	1	1	1	male	22		no	daily
691336	1	1	1	1	1	0	1	female	23		no	daily
691506	0	1	1	1	1	1	1	male	21		yes	1-3/month
691527	1	0	0	1	1	1	1	male	23		yes	1/week
691849	1	1	1	1	0	0	0	female	27		no	1-3/month
691972	1	1	0	1	1	0	1	female	22		no	2-3/week

```
> # overview of the variables
> colnames(dat)
```

```
[1] "X1"      "X2"      "X3"      "X4"      "X5"      "X6"
[7] "X7"      "X8"      "X9"      "X10"     "X11"     "X12"
[13] "X13"     "X14"     "X15"     "X16"     "X17"     "X18"
[19] "X19"     "X20"     "X21"     "X22"     "X23"     "X24"
[25] "X25"     "X26"     "X27"     "X28"     "X29"     "X30"
[31] "X31"     "X32"     "X33"     "X34"     "X35"     "X36"
[37] "X37"     "X38"     "X39"     "X40"     "X41"     "X42"
[43] "X43"     "X44"     "X45"     "gender"  "age"     "semester"
[49] "elite"   "spon"
```

```
> str(dat)
```

```
'data.frame':      1075 obs. of  50 variables:
 $ X1      : num  1 0 0 0 1 1 0 1 1 0 ...
 $ X2      : num  0 1 0 0 1 0 1 1 0 0 ...
 $ X3      : num  0 0 0 1 1 0 0 1 0 0 ...
 $ X4      : num  1 1 1 1 1 1 1 0 1 1 ...
 $ X5      : num  1 1 0 1 1 0 1 1 0 1 ...
 $ X6      : num  1 1 1 0 1 1 1 1 0 0 ...
 $ X7      : num  1 0 0 1 1 1 0 1 1 0 ...
 $ X8      : num  0 0 0 0 1 1 0 0 0 1 ...
 $ X9      : num  0 0 1 1 1 0 0 0 1 0 ...
```

```

$ X10      : num  1 1 1 0 1 1 1 1 1 1 ...
$ X11      : num  1 0 1 0 1 0 0 0 0 1 ...
$ X12      : num  1 1 1 1 1 1 1 1 1 1 ...
$ X13      : num  0 1 0 0 1 0 1 0 0 0 ...
$ X14      : num  0 0 0 0 0 0 1 0 0 0 ...
$ X15      : num  0 1 1 1 1 1 1 0 1 0 ...
$ X16      : num  1 0 1 1 1 1 1 1 1 0 ...
$ X17      : num  0 0 0 1 1 0 0 0 1 0 ...
$ X18      : num  1 1 0 0 1 1 0 0 0 0 ...
$ X19      : num  0 0 0 0 0 0 0 0 0 0 ...
$ X20      : num  0 1 0 0 1 1 1 0 1 1 ...
$ X21      : num  1 0 0 1 0 1 1 0 0 0 ...
$ X22      : num  1 1 0 1 1 0 0 1 1 1 ...
$ X23      : num  1 1 0 0 1 1 0 0 1 1 ...
$ X24      : num  1 1 1 1 1 0 1 0 1 0 ...
$ X25      : num  0 0 1 1 1 0 1 0 0 0 ...
$ X26      : num  0 1 0 0 1 0 0 0 1 0 ...
$ X27      : num  0 1 0 0 0 1 1 1 1 0 ...
$ X28      : num  0 0 1 1 0 1 1 1 0 1 ...
$ X29      : num  0 0 1 1 1 0 0 1 0 1 ...
$ X30      : num  0 1 1 0 1 1 1 0 1 0 ...
$ X31      : num  1 0 0 0 1 0 0 1 1 1 ...
$ X32      : num  1 1 0 0 1 1 0 0 1 1 ...
$ X33      : num  1 1 1 1 0 1 1 1 1 1 ...
$ X34      : num  1 0 1 1 1 1 1 0 1 1 ...
$ X35      : num  1 0 1 1 1 1 1 1 1 1 ...
$ X36      : num  1 1 1 1 1 1 1 0 1 1 ...
$ X37      : num  1 1 1 0 1 0 1 1 1 1 ...
$ X38      : num  1 1 1 1 1 1 1 1 1 0 ...
$ X39      : num  0 0 0 1 1 1 0 0 0 0 ...
$ X40      : num  1 1 1 1 1 1 1 1 1 1 ...
$ X41      : num  0 1 0 1 0 1 0 0 1 0 ...
$ X42      : num  1 1 1 1 0 1 1 1 1 1 ...
$ X43      : num  1 1 1 0 1 0 0 1 1 1 ...
$ X44      : num  1 1 1 1 1 1 0 0 0 0 ...
$ X45      : num  1 1 1 1 1 1 1 1 1 1 ...
$ gender   : Factor w/ 2 levels "female","male": 1 2 1 2 2 2 2 2 1 ...
$ age      : num  21 20 25 27 24 20 22 22 21 20 ...
$ semester: Ord.factor w/ 11 levels "1"<"2"<"3"<"4"<...: 3 1 9 10 8 1 2 2 1 1 ...
$ elite    : Factor w/ 2 levels "no","yes": 1 1 1 1 1 2 1 1 1 1 ...
$ spon     : Ord.factor w/ 7 levels "never"<"<1/month"<...: 3 6 3 1 4 3 1 4 4 3 ...

```

## 2 Get some summary and plots

R provides some basic built-in function that return summary or visualizations. Here for the `elite` (coded as a factor). The `summary` function is a generic function. It checks the data type and then returns what it thinks is the best output (here summary is a table). Alternatively, the `table` function works fine as well.

```
> summary(dat$elite)
```

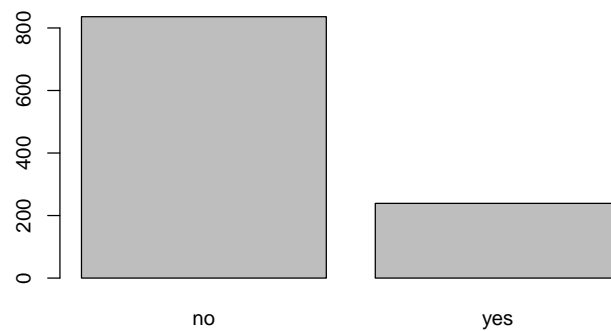
```
no yes  
836 239
```

```
> table(dat$elite)
```

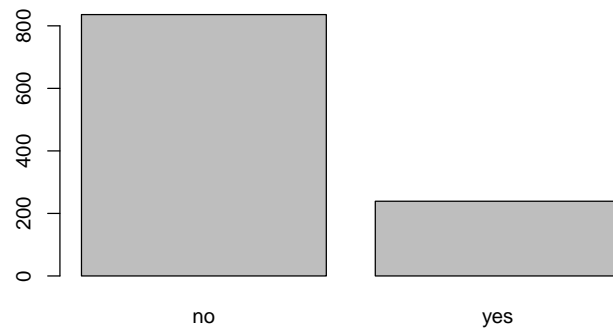
```
no yes  
836 239
```

When we call `plot` for a factor, we get a barplot (or we use `barplot` directly). But we can also easily get a `pie` chart!

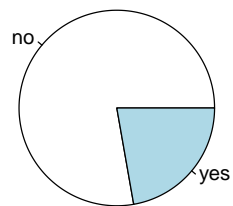
```
> plot(dat$elite)
```



```
> # alternative  
> barplot(table(dat$elite))
```



```
> pie(table(dat$elite))
```



The same applies to the variable `spon` (coded as an ordered factor). As `spon` is ordered, the order of the values in the plot is not alphabetically, but ordered as the factor order.

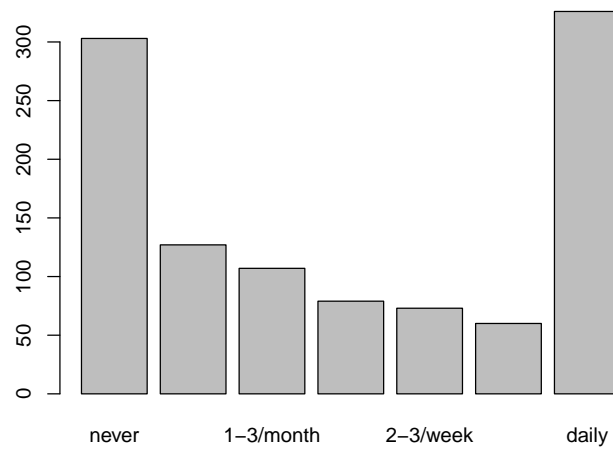
```
> summary(dat$spon)
```

never	<1/month	1-3/month	1/week	2-3/week	4-5/week	daily
303	127	107	79	73	60	326

```
> table(dat$spon)
```

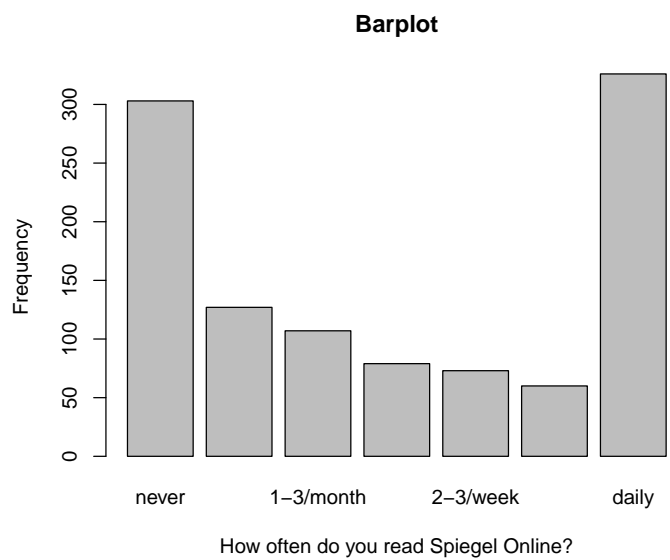
never	<1/month	1-3/month	1/week	2-3/week	4-5/week	daily
303	127	107	79	73	60	326

```
> plot(dat$spon)
```

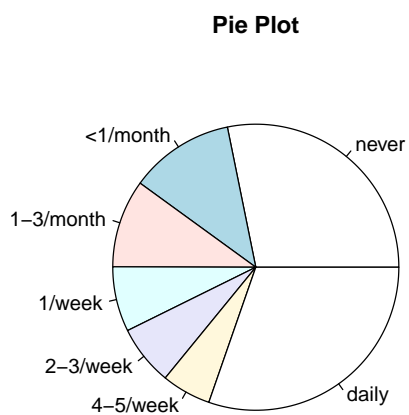


We can also add titles or axis labels

```
> barplot(table(dat$spon),  
+         xlab = "How often do you read Spiegel Online?", ylab = "Frequency",  
+         main = "Barplot")
```



```
> pie(table(dat$spon), main = "Pie Plot")
```



For `age` (continuous variable), we get different plots. But also here, *R* tries to find the best solution for us, when we just use `summary` or `plot`. But maybe a scatterplot is not what we want, so ask for a `histogram` instead.

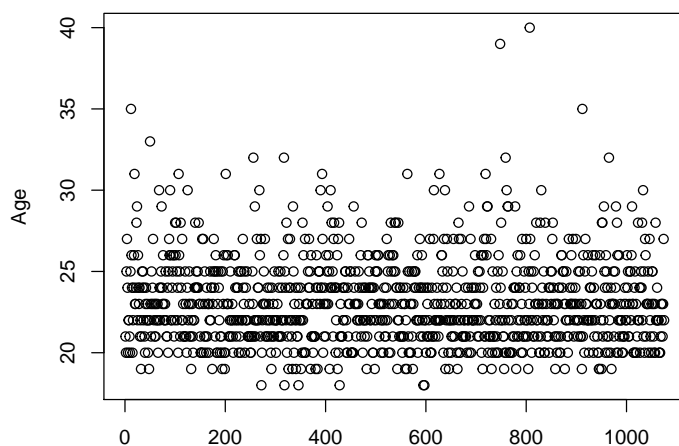
```
> table(dat$age)
```

18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	35	39	40
6	37	124	159	188	149	136	102	61	43	28	15	11	7	4	1	2	1	1

```
> summary(dat$age)
```

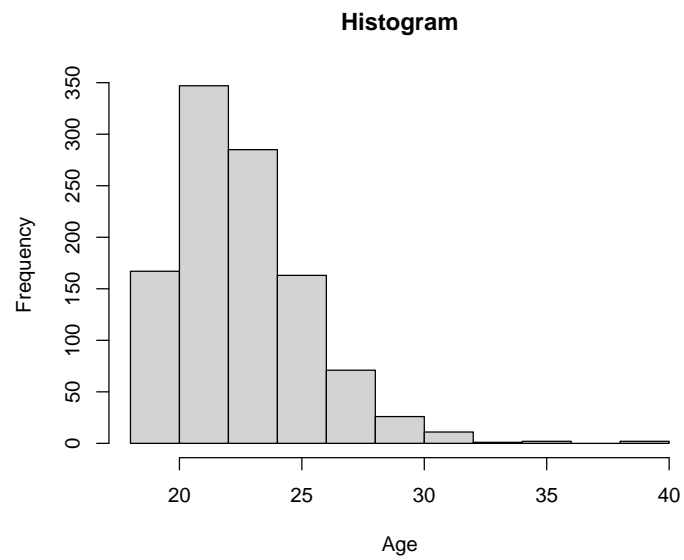
Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
18.0	21.0	23.0	23.1	25.0	40.0

```
> plot(dat$age,
+       xlab = "", ylab = "Age")
```





```
> hist(dat$age,  
+       xlab = "Age",  
+       main = "Histogram")
```



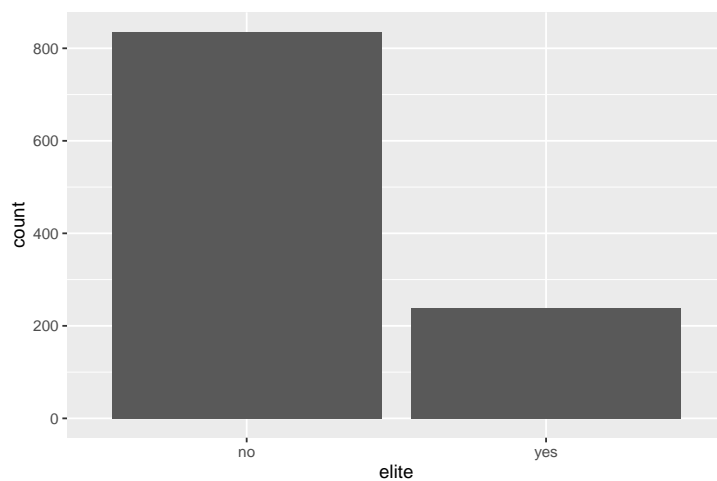
## 2.1 In another universe

The plot functions that we have used until now are R's built-in functions. Alternatively, we can use functions from packages from the **tidyverse** by Hadley Wickham. He has written many functions that some people find easier to use than R's built-in functions.

```
> # install.packages("ggplot2")  
> library(ggplot2)
```

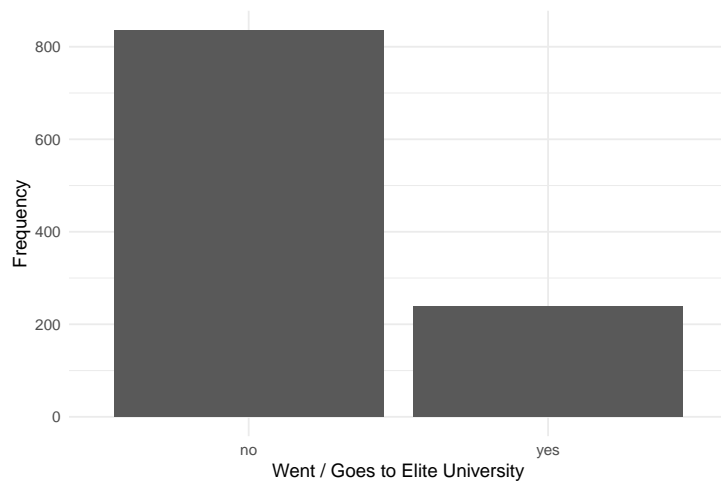
The plotting mechanism for **ggplot** works very differently from **base R**. We first define what will be on the x-axis and y-axis, and then add what kind of plot we want.

```
> ggplot(dat, aes(x = elite)) +  
+   geom_bar()
```



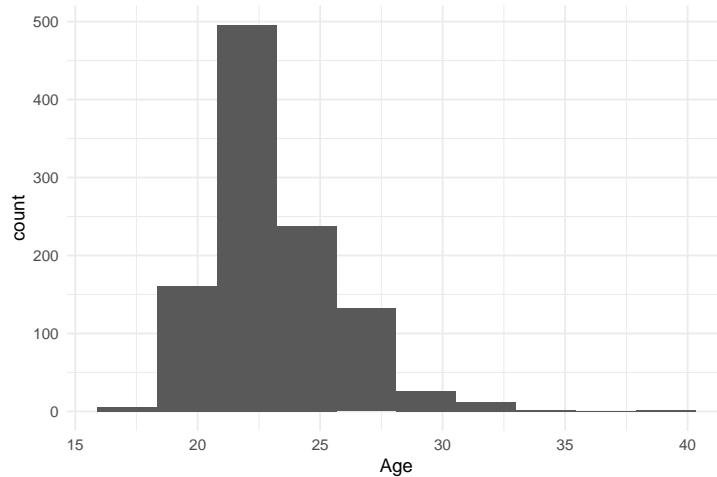
We can add more options, such as axis labels and another background theme

```
> ggplot(dat, aes(x = elite)) +  
+   geom_bar() +  
+   xlab("Went / Goes to Elite University") +  
+   ylab("Frequency") +  
+   theme_minimal()
```



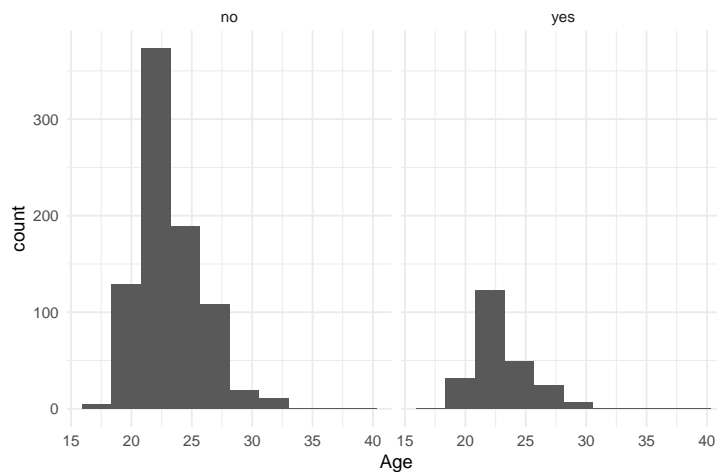
Similarly for the continuous variable `age`, we can ask for a histogram

```
> ggplot(dat, aes(x = age)) +  
+   geom_histogram(bins = 10) + # with "bins", we can set the number of bins  
+   xlab("Age") +  
+   theme_minimal()
```



With `ggplot`, you can very easily create faceted plots. There are very helpful for data exploration. Here, we explore whether the distribution of `Age` is different between people that went/go to a elite university:

```
> ggplot(dat, aes(x = age)) +  
+   geom_histogram(bins = 10) +  
+   facet_wrap(~elite) +  
+   xlab("Age") +  
+   theme_minimal()
```



Some people like the **base R** way of plotting more, some the **ggplot** way of plotting. I use both, with a tendency to **base R** because I think it is more flexible. Some people find **ggplot** easier to use or to get used to. It is up to you to decide which one you choose, both are great and provide much more options than we could cover in this workshop! Maybe you find these cheatsheets helpful when you work with visualizations in the future:

- base R: <http://publish.illinois.edu/johnrgallagher/files/2015/10/BaseGraphicsCheatsheet.pdf>
- ggplot: <https://statsandr.com/blog/files/ggplot2-cheatsheet.pdf>

### 3 Subsetting & computing new variables

To compute a sum score of items correct, we have to extract the test items. There is several options to do so:

```
> # subset data
> testItems <- dat[, 1:45]
> # alternativ:
> testItems <- dat[, paste("X", 1:45, sep = "")]
> # alternativ:
> testItems <- dat[, grep("X", colnames(dat))]
```

Then we can compute a sumscore. We use the `rowSums` function, that computes a sum row-wise (there is also `colSums`):

```
> dat$sumscore <- rowSums(testItems)
> head(dat)
```

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20	X21
743	1	0	0	1	1	1	1	0	0	1	1	1	0	0	0	1	0	1	0	0	1
801	0	1	0	1	1	1	0	0	0	1	0	1	1	0	1	0	0	1	0	1	0
3747	0	0	0	1	0	1	0	0	1	1	1	1	0	0	1	1	0	0	0	0	0
4273	0	0	1	1	1	0	1	0	1	0	0	1	0	0	1	1	1	0	0	0	1
4366	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1	1	1	1	0	1	0
5177	1	0	0	1	0	1	1	1	0	1	0	1	0	0	1	1	0	1	0	1	1

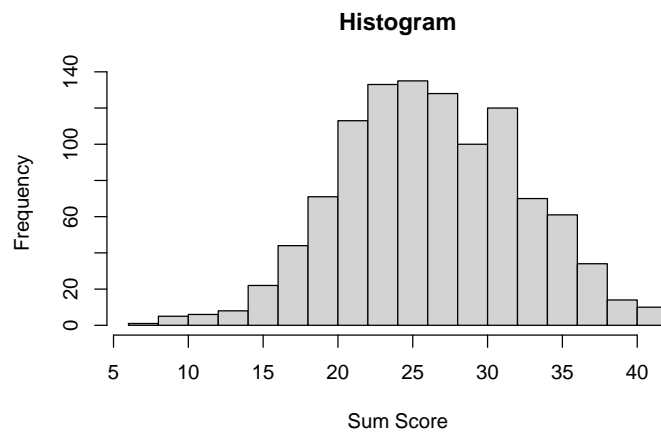
	X22	X23	X24	X25	X26	X27	X28	X29	X30	X31	X32	X33	X34	X35	X36	X37	X38	X39
743	1	1	1	0	0	0	0	0	0	0	1	1	1	1	1	1	1	0
801	1	1	1	0	1	1	0	0	1	0	1	1	0	0	1	1	1	0
3747	0	0	1	1	0	0	1	1	1	0	0	1	1	1	1	1	1	0
4273	1	0	1	1	0	0	1	1	0	0	0	1	1	1	1	0	1	1
4366	1	1	1	1	1	0	0	1	1	1	1	0	1	1	1	1	1	1
5177	0	1	0	0	0	1	1	0	1	0	1	1	1	1	1	0	1	1

	X40	X41	X42	X43	X44	X45	gender	age	semester	elite	spon	sumscore
743	1	0	1	1	1	1	female	21		3	no 1-3/month	27
801	1	1	1	1	1	1	male	20		1	no 4-5/week	27
3747	1	0	1	1	1	1	female	25		9	no 1-3/month	24
4273	1	1	1	0	1	1	male	27		10	no never	26
4366	1	0	0	1	1	1	male	24		8	no 1/week	37
5177	1	1	1	0	1	1	male	20		1	yes 1-3/month	28

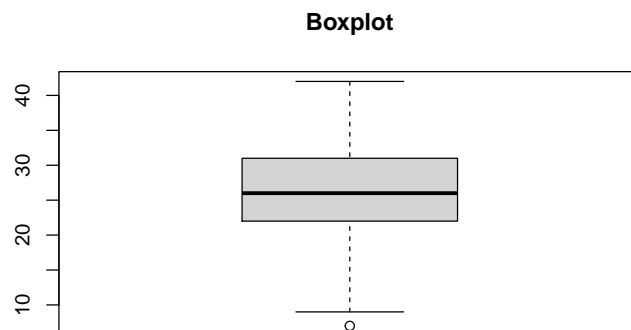
Now again we can explore the sum score by looking at a histogram

```
> hist(dat$sumscore,  
+       xlab = "Sum Score",  
+       main = "Histogram",  
+       breaks = 20)
```



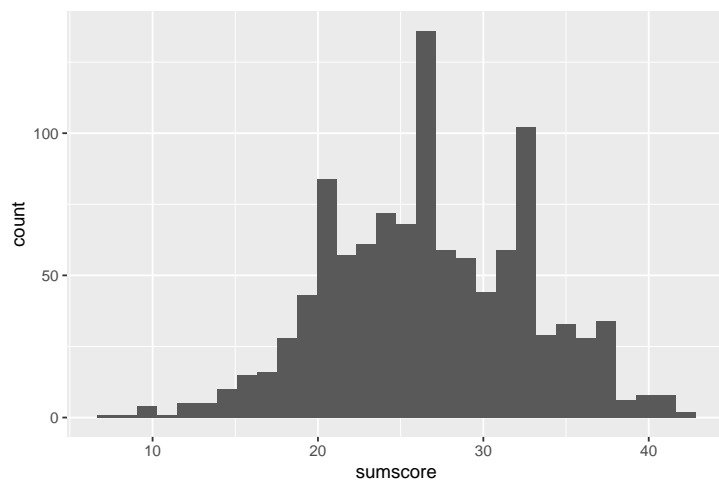
or a boxplot of the new variable.

```
> boxplot(dat$sumscore,  
+         main = "Boxplot")
```

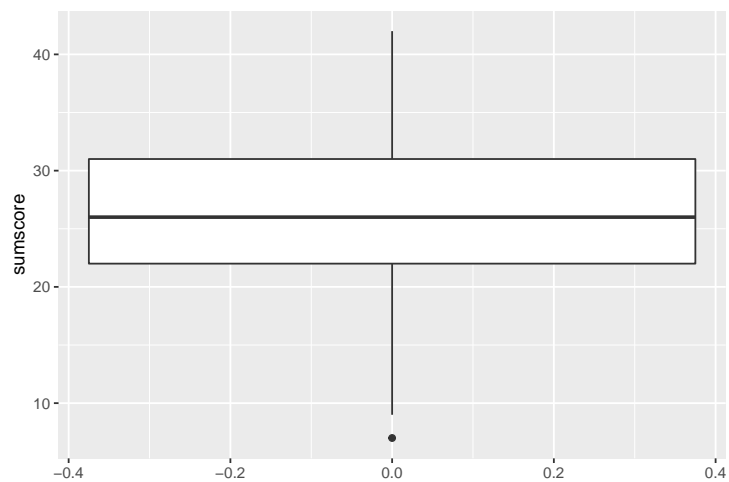


Or we use `ggplot` for plotting:

```
> ggplot(dat, aes(x = sumscore)) +  
+   geom_histogram()
```



```
> ggplot(dat, aes(y = sumscore)) +  
+   geom_boxplot()
```





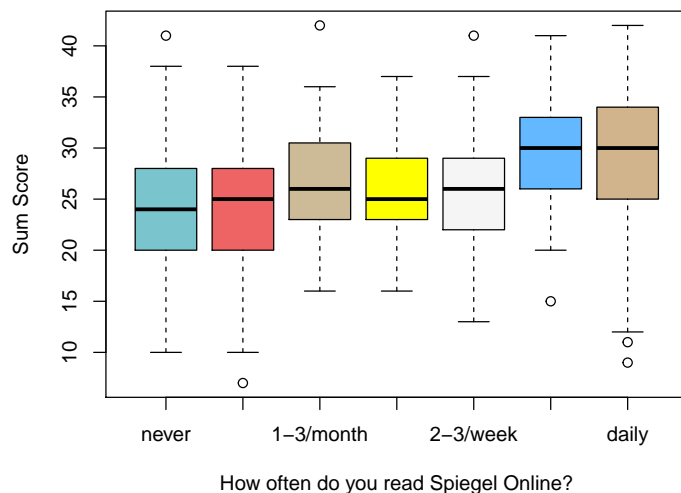
## 4 Bivariate Plots

Now, we want to look at bivariate plots to get more insights into the data. For example, boxplots of the sumscore for each group of Spiegel Online readers (spon). With the `colors` function, we can get the names of many nice colors. There are also color palettes, or you use HEX or rgb codes.

```
> head(colors())

[1] "white"          "aliceblue"      "antiquewhite"   "antiquewhite1"
[5] "antiquewhite2" "antiquewhite3"

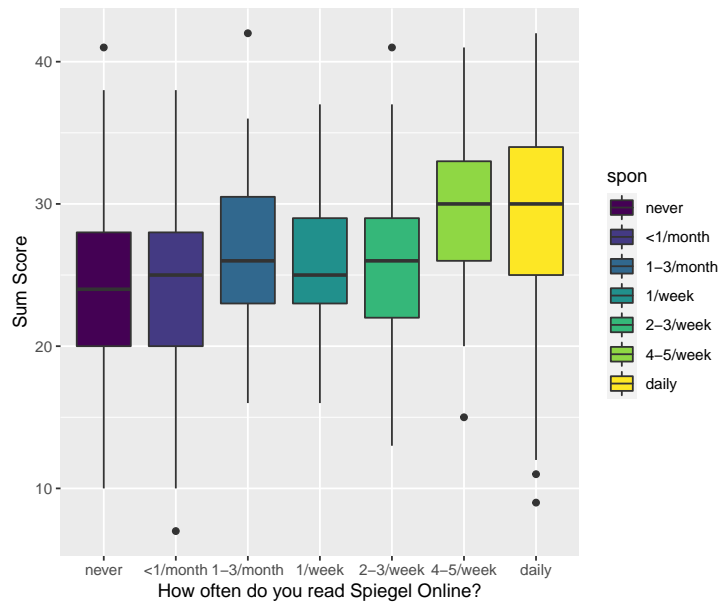
> # boxplot for each group of spon readers
> boxplot(sumscore ~ spon, data = dat,
+         col = c("cadetblue3", "indianred2", "wheat3",
+               "yellow1", "whitesmoke", "steelblue1", "tan"),
+         ylab = "Sum Score",
+         xlab = "How often do you read Spiegel Online?")
```



From a visual inspection, it seems as if regular readers of Spiegel Online had better results in the test.

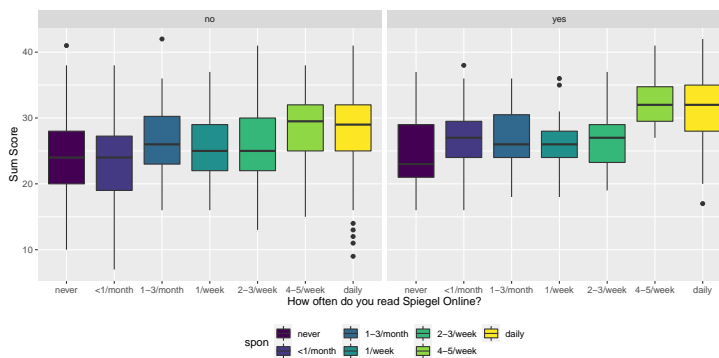
And a `ggplot` graphic (it uses pretty standard palettes for the fill colors):

```
> ggplot(dat, aes(x = spon, y = sumscore, fill = spon)) +
+   geom_boxplot() +
+   ylab("Sum Score") +
+   xlab("How often do you read Spiegel Online?")
```



As I said above, it is so easy to use facetting with `ggplot`:

```
> ggplot(dat, aes(x = spon, y = sumscore, fill = spon)) +
+   geom_boxplot() +
+   facet_wrap(~elite) +
+   theme(legend.position = "bottom") +
+   ylab("Sum Score") +
+   xlab("How often do you read Spiegel Online?")
```



## 5 Linear regression model

We want to fit a linear regression model, using the variables `age` and `elite` as predictor variables, and the `sumscore` as a dependent variable. Of course, we could add more predictors, or also interaction effects (using `*` or `:`). The `summary` function gives us the best output for the regression object.

```
> # fit model and save it in the object "lmod"
> lmod <- lm(sumscore ~ age + elite, data = dat)
> summary(lmod)
```

Call:

```
lm(formula = sumscore ~ age + elite, data = dat)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-19.1326	-4.1326	-0.1326	4.4577	15.4577

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	21.62635	1.56811	13.791	< 2e-16 ***
age	0.20483	0.06722	3.047	0.002368 **
eliteyes	1.59303	0.43929	3.626	0.000301 ***

---

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

Residual standard error: 5.988 on 1072 degrees of freedom

Multiple R-squared: 0.02017, Adjusted R-squared: 0.01834

F-statistic: 11.04 on 2 and 1072 DF, p-value: 1.804e-05

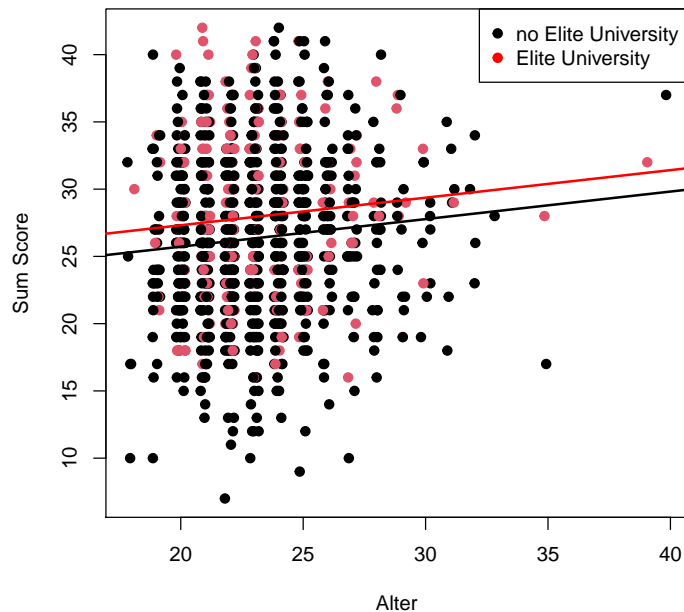
We want to visualize the results of the regression. So we create a scatterplot that plot `age` against `sumscore` and the points are colored by `elite`.

First, we extract the regression weights from the regression object.

```
> # extract regression coefficients
> intercept <- coefficients(lmod)[1]
> age <- coefficients(lmod)[2]
> eliteyes <- coefficients(lmod)[3]
```

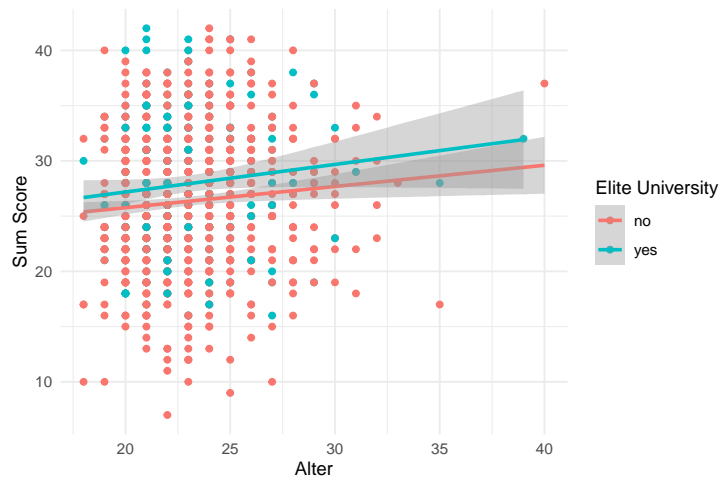
Then, we can create a scatterplot. As age is an integer variable, the scatterplot looks strange and the points are on top of each other. Using `jitter`, we can randomly jitter the points so that we can see them all. We add a `legend` to the plot, so we know what the colors of the points mean. Then, we can add the regression lines, one for students from an elite university, one from non-elite universities.

```
> # create plot
> plot(jitter(dat$age), dat$sumscore,
+      col = dat$elite,
+      pch = 19,
+      xlab = "Alter", ylab = "Sum Score")
> # add legend
> legend("topright", # Position of the legend
+      col = c("black", "red"),
+      legend = c("no Elite University", "Elite University"),
+      pch = 19)
> # add regression lines
> abline(a = intercept, b = age,
+      lwd = 2) # line width
> abline(a = intercept + eliteyes, b = age,
+      col = "red",
+      lwd = 2)
```



Of course we can create a similar plot using ggplot:

```
> # create plot
> ggplot(dat, aes(x = age, y = sumscore,
+               # unintuitively the argument is no named color,
+               # not fill as for the boxplot above
+               color = elite)) +
+   geom_point() +
+   # this how ggplot automatically adds the regression lines
+   geom_smooth(method = "lm",
+               formula = y ~ x) +
+   xlab("Alter") +
+   ylab("Sum Score") +
+   theme_minimal() +
+   # also quite unintuitively: to change legend title, the syntax has
+   # nothing to do with "legend":
+   labs(color = "Elite University")
```



With ggplot, I also search for tutorial (e.g., <https://www.datanovia.com/en/blog/ggplot-legend-title-position-and-labels/> or <https://statisticsglobe.com/add-regression-line-to-ggplot2-plot-in-r>) to adjust labels, legends, add regression lines, because in the `tidyverse`, there is a syntax for everything, but often (I find it) not easy to guess.

When we want to save the R `base` plot in an external file, we must open a device before we draw the plot, and close it afterwards. Here, we export a `png`, but of course other formats are possible. If you want to know which, click on `png` and press F1. alternatively run `?png` in the Console.

```
> # open device
> png("myRegressionPlot.png", height = 500, width = 500)
> # create plot
> plot(jitter(dat$age), dat$sumscore,
+      col = dat$elite,
+      pch = 19,
+      xlab = "Alter", ylab = "Sum Score")
> # add legend
> legend("topright", # Position of the legend
+      col = c("black", "red"),
+      legend = c("no Elite University", "Elite University"),
+      pch = 19)
> # add regression lines
> abline(a = intercept, b = age,
+      lwd = 2) # line width
> abline(a = intercept + eliteyes, b = age,
+      col = "red",
+      lwd = 2)
> # close device
> dev.off()

null device
      1
```

For `ggplot`, we first save the plot as an R object, and then save that object:

```
> # save plot:
> regressionPlot <-
+   ggplot(dat, aes(x = age, y = sumscore, color = elite)) +
+   geom_point() +
+   geom_smooth(method = "lm", formula = y ~ x) +
+   xlab("Alter") +
+   ylab("Sum Score") +
+   theme_minimal() +
+   labs(color = "Elite University")
> ggsave(filename = "myRegressionggPlot.png", regressionPlot)
```

Of course, also pdf files are possible using `filename = "myRegressiongg-Plot.pdf"`.