

# Example of Output of R Markdown

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## Abstract

R includes some features that might simplify the writing process by making it possible to gather data analysis and writing in a single document. This makes it easier to reproduce ones analysis in case the data is changed and to produce presentable papers for self-publishing. This presentation gives a brief example of some of these features.

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## 1 Introduction

There are a number of ways to incorporate R code and ordinary text into a single coherent paper. The most popular way is to write code and text in the Markdown format. Markdown is a mark-up language, meaning that code is written in an ordinary text file and then compiled into a desirable format, for example a pdf file or a Word document. This is a short example of the R Markdown features available in R using the packages `rmarkdown` and `knitr`. It gives examples of the following:

- Lists
- Figures
- Tables
- References

The data `oats` from the `MASS` package is used as example data. We load the dataset with the following code. In this document the R code is shown, but it is easy to hide the code and only show the results.

```
dat <- MASS::oats
```

## 2 Example output

### 2.1 Descriptive statistics

The data can be presented in table form, summarized for each variety and nitrogen combination. The following code summarizes the dataset and presents it as table using the `kable` function from the `knitr` package.

```
dat_agg <- aggregate(Y ~ V + N, dat, function(x) c(mean(x), sd(x)))
dat_agg[,1] <- as.character(dat_agg[,1])
dat_agg <- data.frame(dat_agg[, 1:2], dat_agg[, 3])
names(dat_agg) <- c("Variety", "Nitrogen", "Mean", "Std.dev")
dat_agg[dat_agg$Variety == "Golden.rain", 1] <- "Golden Rain"
dat_agg <- dat_agg[order(dat_agg$Variety), ]

knitr::kable(dat_agg, digits = 2,
              caption = "Mean and standard deviation of yield
                        per variety and nitrogen combination.",
              row.names = F)
```

Table 1: Mean and standard deviation of yield per variety and nitrogen combination.

Variety	Nitrogen	Mean	Std.dev
Golden Rain	0.0cwt	80.00	21.00
Golden Rain	0.2cwt	98.50	13.47
Golden Rain	0.4cwt	114.67	29.94
Golden Rain	0.6cwt	124.83	20.88
Marvellous	0.0cwt	86.67	16.57
Marvellous	0.2cwt	108.50	26.85
Marvellous	0.4cwt	117.17	9.79
Marvellous	0.6cwt	126.83	20.29
Victory	0.0cwt	71.50	20.60
Victory	0.2cwt	89.67	22.51
Victory	0.4cwt	110.83	26.01
Victory	0.6cwt	118.50	30.09

This can be visualized in a bar chart. The `ggplot2` package is used to construct the plot.

```
library(ggplot2)
barchart <- ggplot(dat_agg, aes(Variety, Mean, fill = Nitrogen)) +
  geom_bar(stat = "identity", position = position_dodge(),
           col = "black") +
  geom_errorbar(aes(ymin = Mean - Std.dev,
```

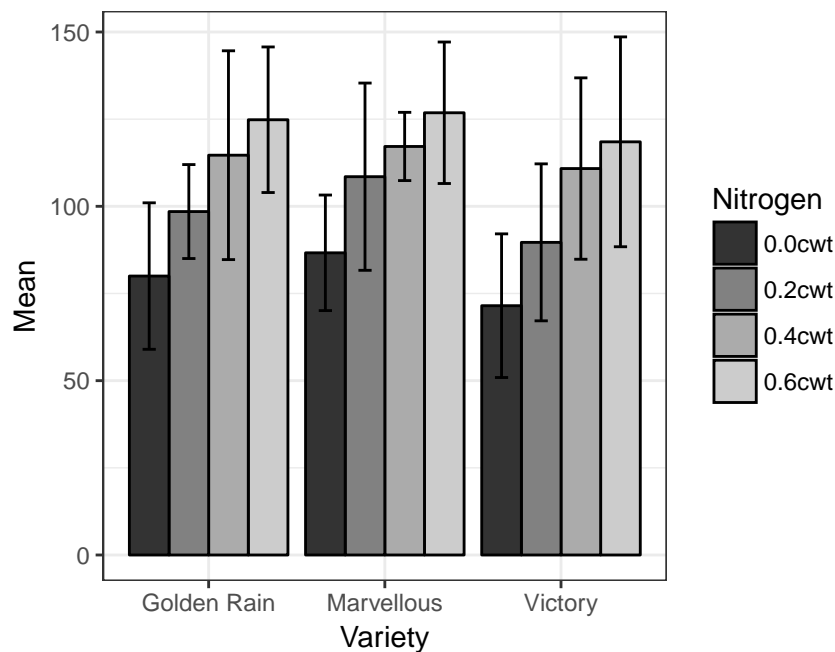


Figure 1: Mean of yield per variety and nitrogen combination. Error bars give standard deviation.

```

      ymax = Mean + Std.dev),
      position = position_dodge(0.9),
      width = 0.3) +
  theme_bw() +
  scale_fill_grey()
print(barchart)

```

## 2.2 Anova model of split-plot experiment

The data is analyzed as a mixed model using variety and nitrogen level as fixed factors, and block and interaction between block and variety as random factors. The results are given below. The model specifies a significant effect of nitrogen level, but not of variety. Once again, the `kable` function prints the table in a presentable form.

```

library(lme4, quietly = T)
library(lmerTest, quietly = T)

names(dat) <- c("Block", "Variety", "Nitrogen", "Yield")
mod <- lmer(Yield ~ Variety + Nitrogen + (1 | Block / Variety), dat)

```

```
knitr::kable(anovaTable <- anova(mod), digits = 2,
             caption = "Anova results of analysis of yield.",
             row.names = T)
```

Table 2: Anova results of analysis of yield.

	Sum Sq	Mean Sq	NumDF	DenDF	F.value	Pr(>F)
Variety	482.91	241.46	2	10	1.49	0.27
Nitrogen	20020.50	6673.50	3	51	41.05	0.00

The estimated mean values of the nitrogen levels are calculated and tested for differences.

```
library(emmeans, quietly = T)

knitr::kable(emmeansTable <- cld(emmeans(mod, "Nitrogen")),
             digits = 2,
             caption = "Comparison between levels of nitrogen.")
```

Table 3: Comparison between levels of nitrogen.

Nitrogen	emmean	SE	df	lower.CL	upper.CL	.group
0.0cwt	79.39	7.13	6.64	62.34	96.44	1
0.2cwt	98.89	7.13	6.64	81.84	115.94	2
0.4cwt	114.22	7.13	6.64	97.17	131.28	3
0.6cwt	123.39	7.13	6.64	106.34	140.44	3

These tables could obviously need a little more cleaning up.

## 2.3 A reference

The Markdown format can handle references in a simple way. For example, the anova model was estimated with the `lmer` function from the package `lme4` (Bates et al. 2015).

The code to construct the reference is simply `[@lme4]` where the part in square brackets is linked to the full reference, which is printed in the last part of the document. In this case the reference is specified in the introduction of the `.Rmd` file, but references can also be stored in a separate file and handled using special software.

## References

Bates, Douglas, Martin Machler, Ben Bolker, and Steve Walker. 2015. “Fitting Linear Mixed-Effects Models Using Lme4.” *Journal of Statistical Software* 67 (1):1–48.