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ggplot2 facet : split a plot into a matrix of panels



- [Data](#)
- [Basic box plot](#)
- [Facet with one variable](#)
- [Facet with two variables](#)
- [Facet scales](#)
- [Facet labels](#)
- [facet_wrap](#)
- [Infos](#)

The **facet** approach partitions a plot into a matrix of panels. Each panel shows a different subset of the data. This **R tutorial** describes how to split a graph using **ggplot2** package.

There are two main functions for faceting :

- `facet_grid()`
- `facet_wrap()`

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Data

ToothGrowth data is used in the following examples.

```
# Convert dose from numeric to factor variables
ToothGrowth$dose <- as.factor(ToothGrowth$dose)
df <- ToothGrowth
head(df)
```

```
##      len supp dose
## 1  4.2   VC  0.5
## 2 11.5   VC  0.5
## 3  7.3   VC  0.5
## 4  5.8   VC  0.5
## 5  6.4   VC  0.5
## 6 10.0   VC  0.5
```



Make sure that the variable *dose* is converted as a factor using the above R script.

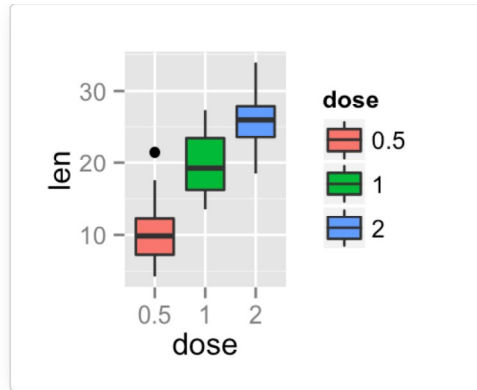
Basic box plot

Create a basic box plot filled by groups :

```
library(ggplot2)
bp <- ggplot(df, aes(x=dose, y=len, group=dose)) +
  geom_boxplot(aes(fill=dose))
bp
```

R Packages

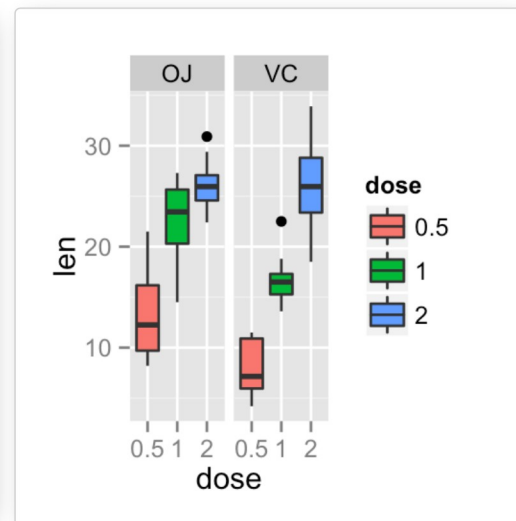
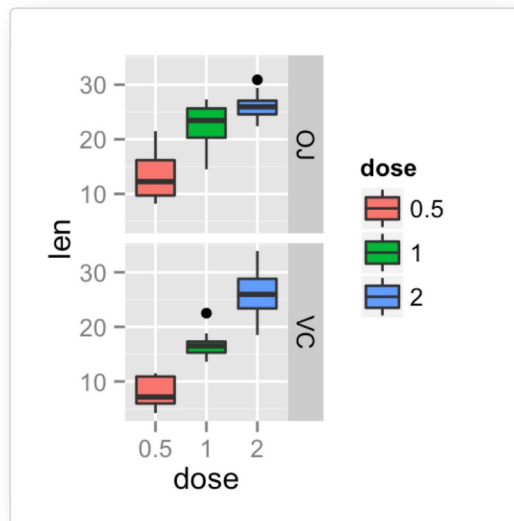
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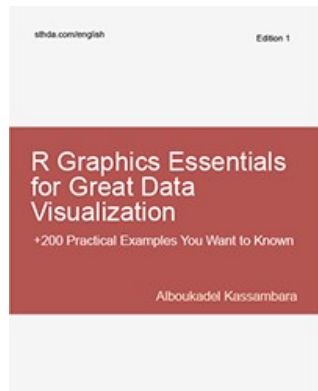
Facet with one variable

The graph is partitioned in multiple panels by levels of the group "supp":

```
# Split in vertical direction  
bp + facet_grid(supp ~ .)  
  
# Split in horizontal direction  
bp + facet_grid(. ~ supp)
```



Our Books



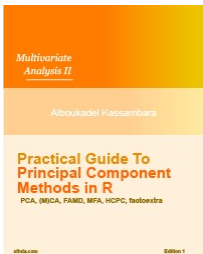
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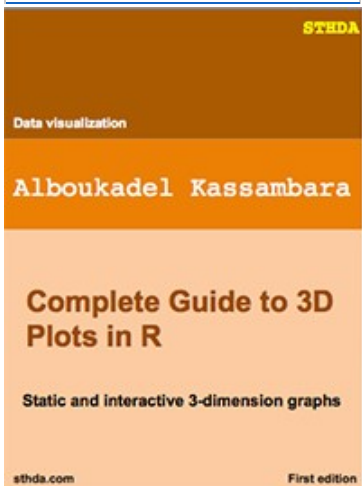
3D Plots in R



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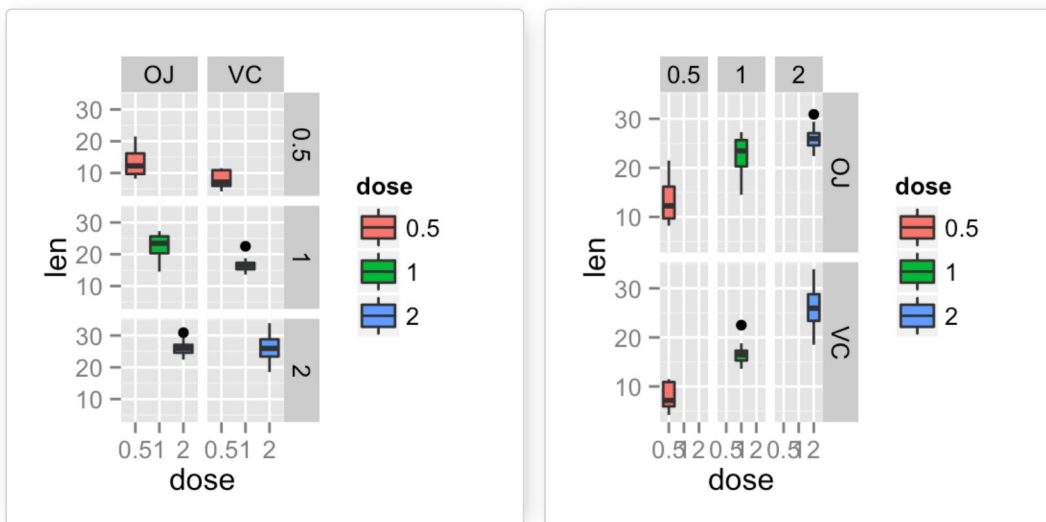


Facet with two variables

The graph is partitioned by the levels of the groups “dose” and “supp” :

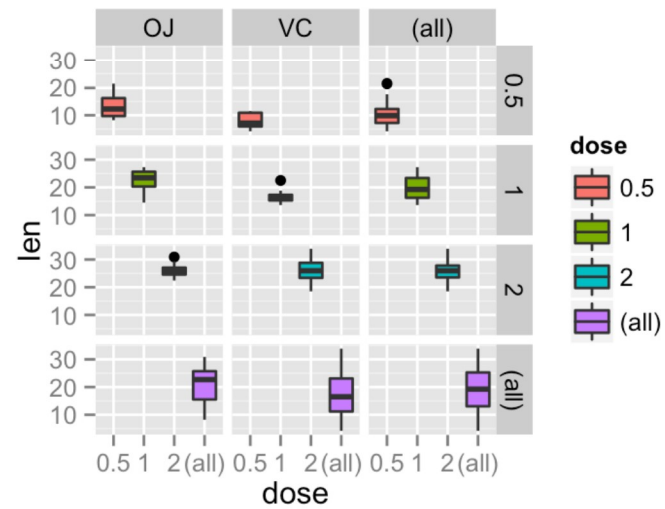
```
# Facet by two variables: dose and supp.
# Rows are dose and columns are supp
bp + facet_grid(dose ~ supp)

# Facet by two variables: reverse the order of the 2 variables
# Rows are supp and columns are dose
bp + facet_grid(supp ~ dose)
```



⚠ Note that, you can use the argument *margins* to add additional facets which contain all the data for each of the possible values of the faceting variables

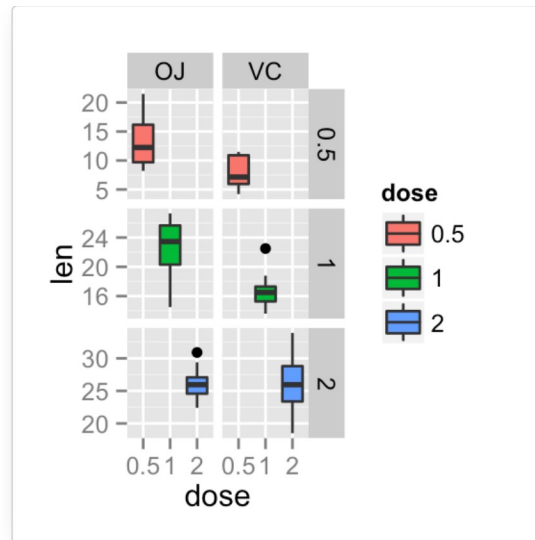
```
bp + facet_grid(dose ~ supp, margins=TRUE)
```



Facet scales

By default, all the panels have the same scales (`scales="fixed"`). They can be made independent, by setting scales to `free`, `free_x`, or `free_y`.

```
bp + facet_grid(dose ~ supp, scales='free')
```

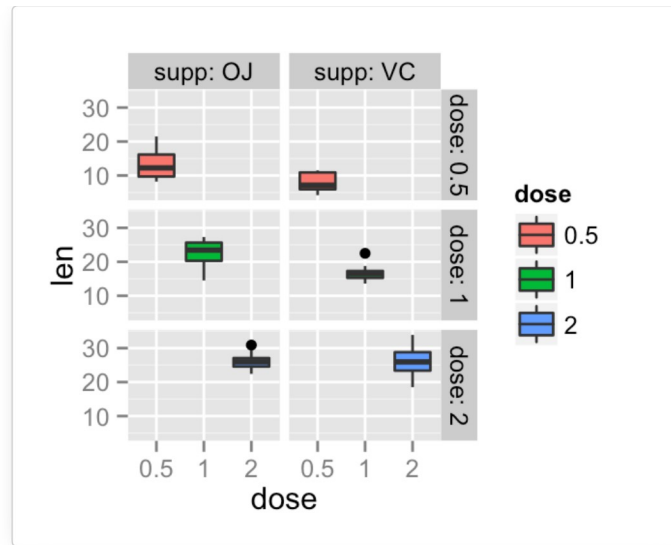


✓ As you can see in the above plot, y axis have different scales in the different panels.

Facet labels

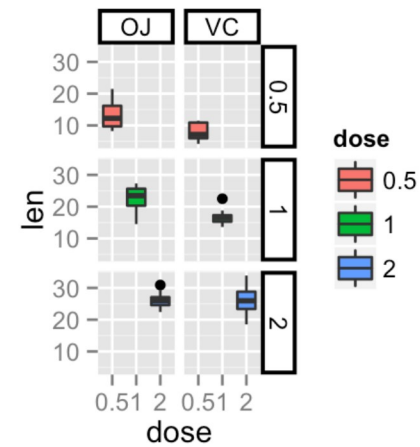
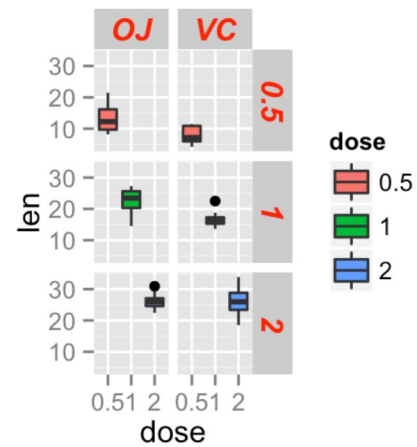
The argument *labeller* can be used to control the labels of the panels :

```
bp + facet_grid(dose ~ supp, labeller=label_both)
```



The appearance of facet labels can be modified as follow :

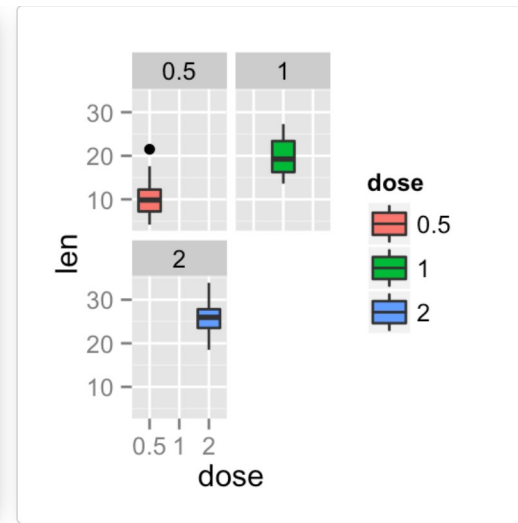
[illegible]



facet_wrap

Facets can be placed side by side using the function **facet_wrap()** as follow :

```
bp + facet_wrap(~ dose)
bp + facet_wrap(~ dose, ncol=2)
```

Infos



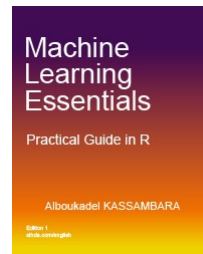
This analysis has been performed using **R software** (ver. 3.1.2) and **ggplot2** (ver. 1.0.0)



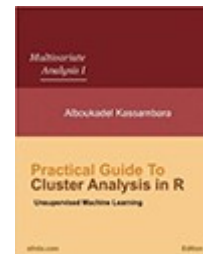
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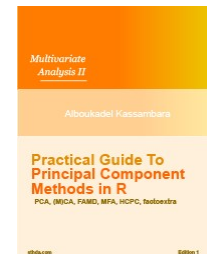
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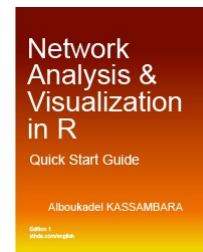
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