

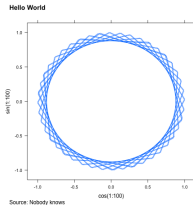
Lattice graphics

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December 14, 2016

Biometris

Wageningen University & Research



Displaying data (and models)

Good data graphics:
display data accurately and
clearly



Seminal reference:
"Visualizing data"
William S. Cleveland, (1993)

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William S. Cleveland, (1993)

"How to display data badly",

Howard Wainer, *The American Statistician* (1984)

1. show as few data as possible
2. hide what data you do show
3. ignore the visual metaphor
4. only order matters
5. graph data out of context
6. change scales in mid-axis
7. emphasize the trivial
8. jiggle the baseline
9. Austria first!
10. Label illegibly, incompletely, incorrectly and ambiguously
11. more is murkier
12. if it has been done well in the past, do it differently

Example: choice of colors

Matlab's colorjet:



- ▶ introduces artefacts
- ▶ highlights specific regions
- ▶ no BW print
- ▶ bad for colour-impaired people

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Alternatives:



From the **RColorBrewer** package

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base graphics, based on Cleveland principles

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1. **graphics** (R Core team, ...)
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Both **lattice** and **ggplot2** make extensive use of the **grid** package (Paul Murrel, R Core team)

Example: the `mtcars` data

```
data(mtcars)  
class(mtcars)
```

```
[1] "data.frame"
```

```
dim(mtcars)
```

```
[1] 32 11
```

Example: the `mtcars` data

```
data(mtcars)
class(mtcars)
```

```
[1] "data.frame"
```

```
dim(mtcars)
```

```
[1] 32 11
```

```
head(mtcars[,1:4])
```

	mpg	cyl	disp	hp
Mazda RX4	21.0	6	160	110
Mazda RX4 Wag	21.0	6	160	110
Datsun 710	22.8	4	108	93
Hornet 4 Drive	21.4	6	258	110
Hornet Sportabout	18.7	8	360	175
Valiant	18.1	6	225	105

Miles per gallon...

```
ncyl <- c(8, 6, 4)
par(mfrow = c(3,1))

for (nc in ncyl) {
  idx <- which(mtcars$cyl == nc)

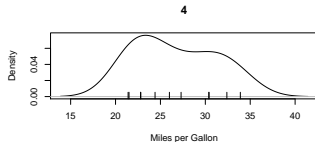
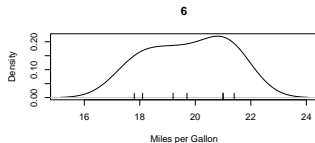
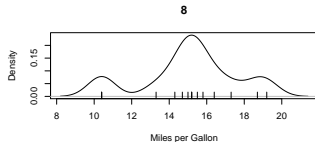
  plot(density(mtcars$mpg[idx]),
       main = nc,
       xlab = "Miles per Gallon",
       ylab = "Density")
  rug(mtcars$mpg[idx], ticksize = .1)
}
```

Miles per gallon...

```
ncyl <- c(8, 6, 4)
par(mfrow = c(3,1))

for (nc in ncyl) {
  idx <- which(mtcars$cyl == nc)

  plot(density(mtcars$mpg[idx]),
       main = nc,
       xlab = "Miles per Gallon",
       ylab = "Density")
  rug(mtcars$mpg[idx], ticksize = .1)
}
```

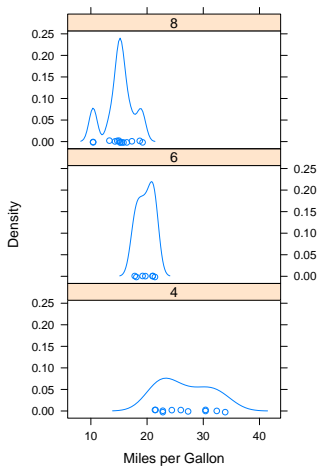


The same in lattice

```
mtcars$cyl.f <- factor(mtcars$cyl)
densityplot(~ mpg | cyl.f,
  data = mtcars,
  xlab = "Miles per Gallon",
  layout = c(1,3))
```

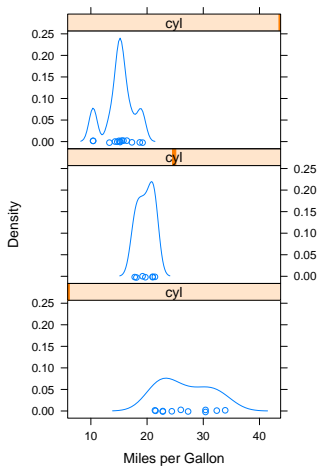

The same in lattice

```
mtcars$cyl.f <- factor(mtcars$cyl)
densityplot(~ mpg | cyl.f,
  data = mtcars,
  xlab = "Miles per Gallon",
  layout = c(1,3))
```



By the way...

```
densityplot(~ mpg | cyl,  
            data = mtcars,  
            xlab = "Miles per Gallon",  
            layout = c(1,3))
```



And a more complicated example...

How about two factors?

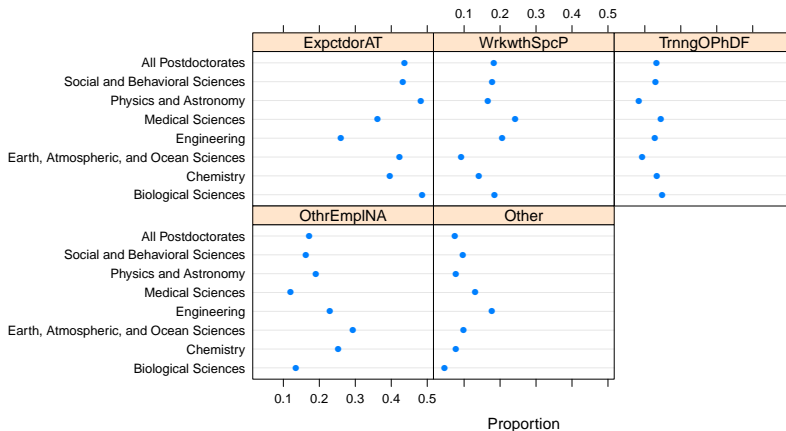
```
data(postdoc, package = "latticeExtra")  
rownames(postdoc)
```

```
[1] "Biological Sciences"  
[2] "Chemistry"  
[3] "Earth, Atmospheric, and Ocean Sciences"  
[4] "Engineering"  
[5] "Medical Sciences"  
[6] "Physics and Astronomy"  
[7] "Social and Behavioral Sciences"  
[8] "All Postdoctorates"
```

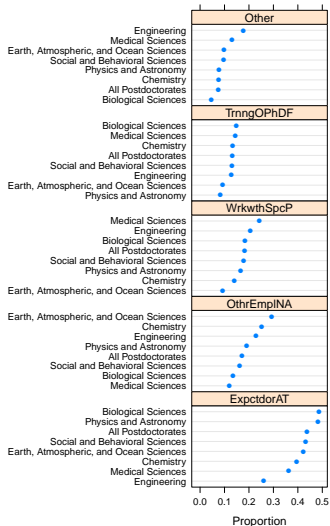
```
colnames(postdoc)
```

```
[1] "Expected or Additional Training" "Work with Specific Person"  
[3] "Training Outside PhD Field"    "Other Employment Not Available"  
[5] "Other"
```

The postdoc plot (first version)



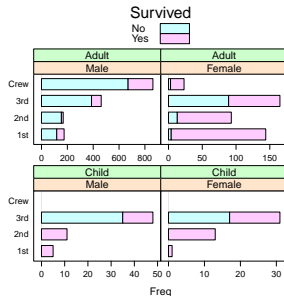
Another version...



- ▶ Each panel sorted: easy interpretation
- ▶ Common x scale: easy comparison
- ▶ Include zero on x scale

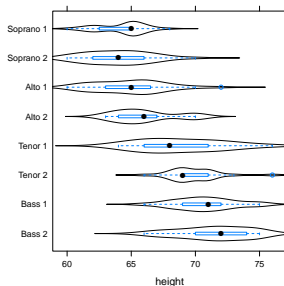
Basic lattice plots: univariate

- `barchart`
- `bwplot`
- `densityplot`
- `stripplot`
- `dotplot`
- `histogram`
- `qqmath`



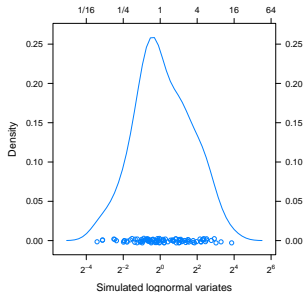
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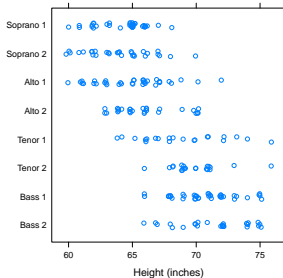
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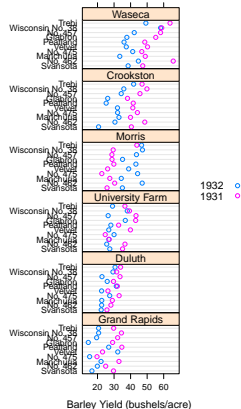
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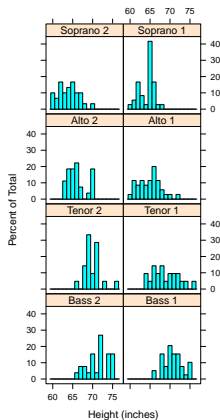
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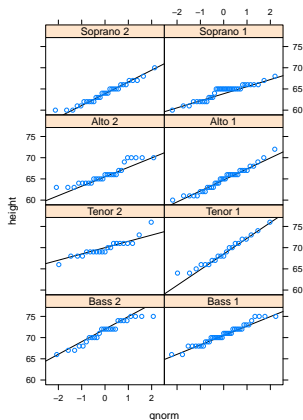
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Basic lattice plots: univariate

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Further lattice plots

Bivariate, trivariate

- ▶ `qq`
- ▶ `xyplot`
- ▶ `levelplot`
- ▶ `contourplot`
- ▶ `cloud`
- ▶ `wireframe`

Hypervariate, miscellaneous

- ▶ `splom`
- ▶ `parallel`
- ▶ `rfs`
- ▶ `tmd`

Key features

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Key features

- ▶ formula interface
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- ▶ also methods for other variable types
- ▶ automatic splits into panels
- ▶ defined by factor levels or variable ranges
- ▶ indications of groups
- ▶ automatic keys/legends
- ▶ highly customizable

Standard plot customizations

Data: locations and
characteristics of earthquakes
near Fiji

Simplest possible case: show
locations

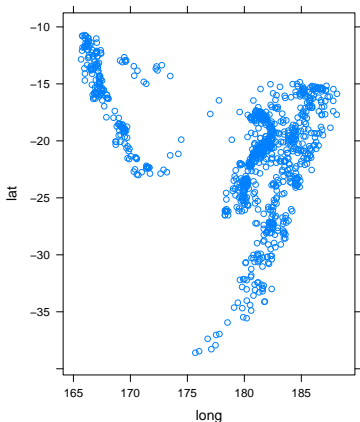
```
xyplot(lat ~ long,  
       data = quakes,  
       aspect = "iso")
```

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Data: locations and characteristics of earthquakes near Fiji

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xyplot(lat ~ long,  
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Standard plot customizations

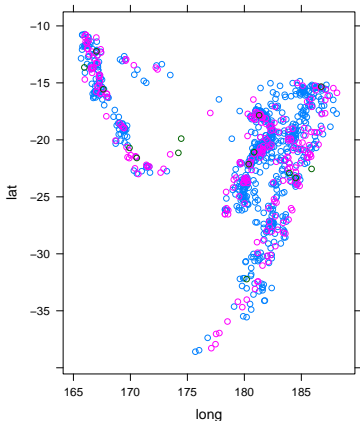
Now: location and magnitude

```
xyplot(lat ~ long,  
  data = quakes,  
  groups = cut(mag, 3),  
  aspect = "iso")
```


Standard plot customizations

Now: location and magnitude

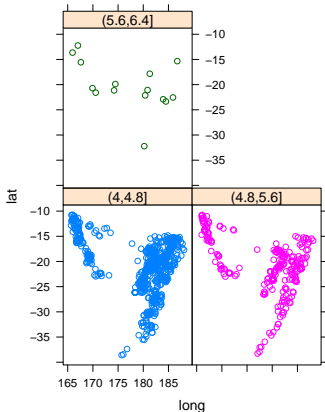
```
xyplot(lat ~ long,  
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  groups = cut(mag, 3),  
  aspect = "iso")
```



Standard plot customizations

Now: location and magnitude

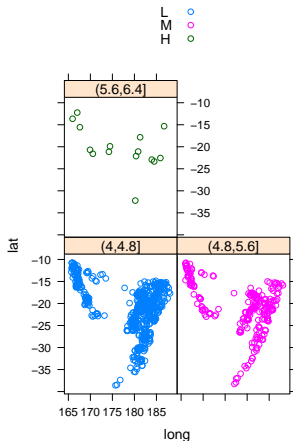
```
xyplot(lat ~ long | cut(mag, 3),  
  data = quakes,  
  groups = cut(mag, 3),  
  aspect = "iso")
```



Standard plot customizations

Now: location and magnitude

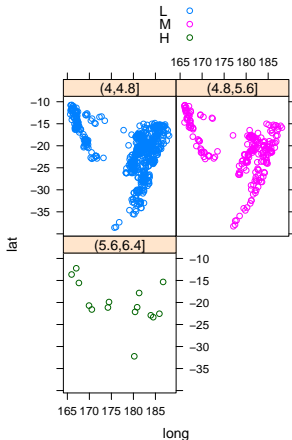
```
xyplot(lat ~ long | cut(mag, 3),
  data = quakes,
  groups = cut(mag, 3),
  aspect = "iso",
  auto.key =
    list(text = c("L", "M", "H"),
         space = "top"))
```



Standard plot customizations

Now: location and magnitude

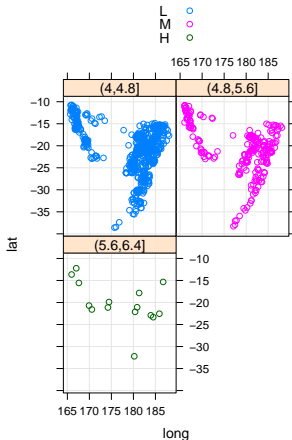
```
xyplot(lat ~ long | cut(mag, 3),
  data = quakes,
  groups = cut(mag, 3),
  aspect = "iso",
  as.table = TRUE,
  auto.key =
    list(text = c("L", "M", "H"),
         space = "top"))
```



Standard plot customizations

Now: location and magnitude

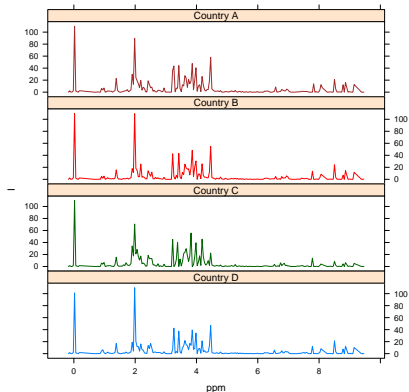
```
xyplot(lat ~ long | cut(mag, 3),
  data = quakes,
  groups = cut(mag, 3),
  aspect = "iso",
  as.table = TRUE,
  auto.key =
    list(text = c("L", "M", "H"),
         space = "top"),
  type = c("p", "g"))
```



An elegant example: NMR spectra

“Naive” plot leads to an inverted x axis:

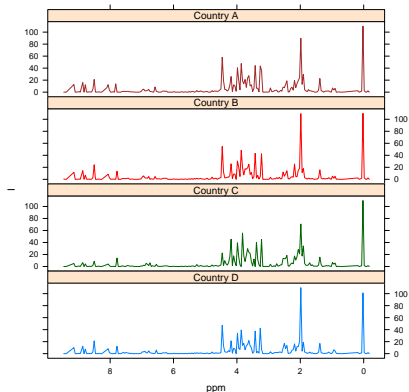
```
xyplot(I ~ ppm | Origin,  
       groups = Sample,  
       data = NMR.df,  
       type = "l",  
       layout = c(1, 4),  
       as.table = TRUE)
```



An elegant example: NMR spectra

... but this is easily remedied:

```
xyplot(I ~ ppm | Origin,
  groups = Sample,
  data = NMR.df,
  type = "l",
  layout = c(1,4),
  as.table = TRUE,
  xlim =
    rev(extendrange(NMR.df$ppm)) )
```

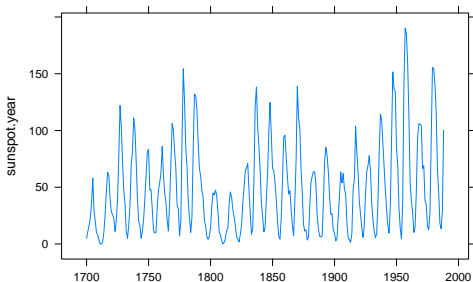


Banking: the sunspots data

```
myplot <-  
  xyplot(sunspot.year ~  
    1700:1988,  
    xlab = "",  
    type = "l")
```

Default:

```
aspect = "fill"
```



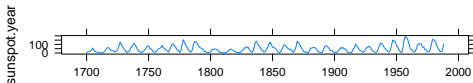
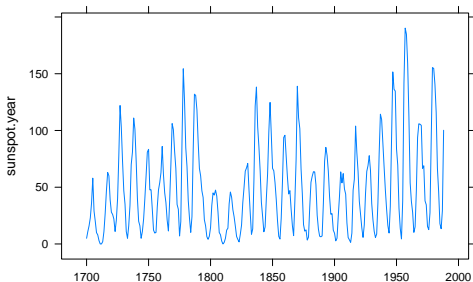
Banking: the `sunspots` data

```
myplot <-  
  xyplot(sunspot.year ~  
    1700:1988,  
    xlab = "",  
    type = "l")
```

Default:

```
aspect = "fill"
```

```
myplot2 <-  
  update(myplot,  
    aspect = "xy")
```



The `prepanel` and `panel` functions

“For the little things...
that take soooo much time”

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- ▶ `packet`: group of data points to be shown in one “shingle”
- ▶ `prepanel` function: determines rectangle to display the packet

The `prepanel` and `panel` functions

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- ▶ `packet`: group of data points to be shown in one “shingle”
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- ▶ `panel` function: determines how to display the packet

The `prepanel` and `panel` functions

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The `prepanel` and `panel` functions

“For the little things...
that take soooo much time”

- ▶ `packet`: group of data points to be shown in one “shingle”
- ▶ `prepanel` function: determines rectangle to display the packet
- ▶ `panel` function: determines how to display the packet

For many plot types there are basic “panel” functions:

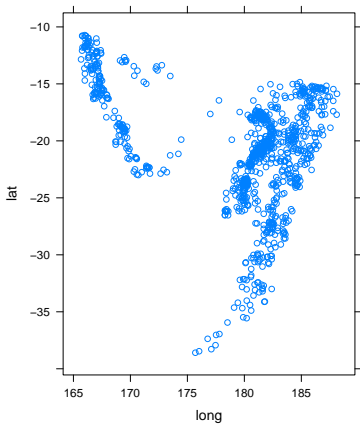
`panel.xyplot`,
`panel.stripplot`,
`panel.splom`, ...

Many supporting plotting functions also have “panel” variants:

`panel.points`, `panel.lines`,
`panel.text`

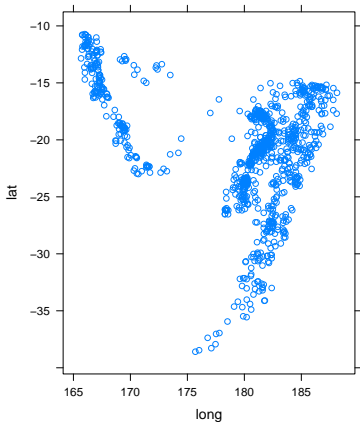
Building it up

```
xyplot(lat ~ long,  
data = quakes,  
aspect = "iso")
```



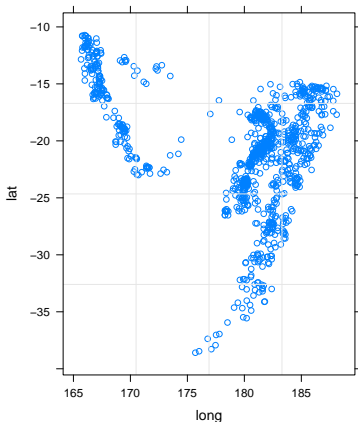
Building it up

```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.xyplot(...)  
  })
```



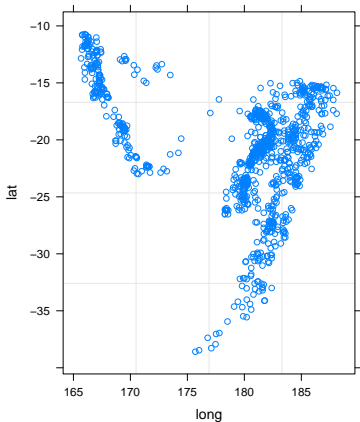
Building it up

```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.xyplot(...)  
    panel.grid()  
  })
```



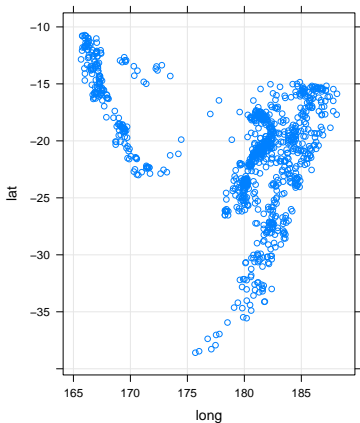
Building it up

```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.grid()  
  
    panel.xyplot(...)  
  })
```



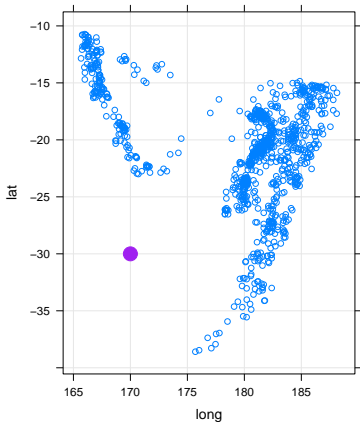
Building it up

```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.grid(h=-1, v=-1)  
  
    panel.xyplot(...)  
  })
```



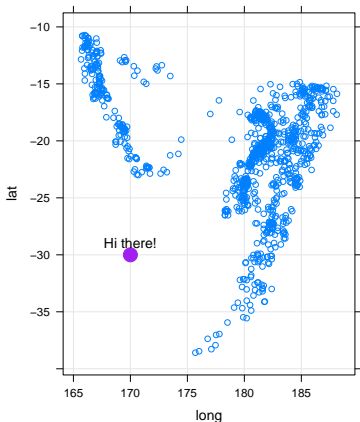
Building it up

```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.grid(h=-1, v=-1)  
    panel.points(170, -30,  
      cex = 2, col = "purple",  
      pch = 19)  
  
    panel.xyplot(...)  
  })
```



Building it up

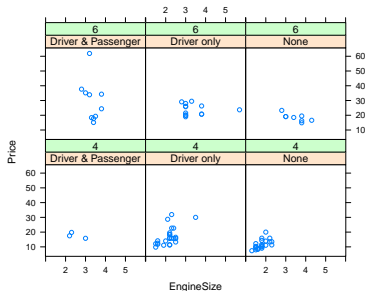
```
xyplot(lat ~ long,  
  data = quakes,  
  aspect = "iso",  
  panel = function(...) {  
    panel.grid(h=-1, v=-1)  
    panel.points(170, -30,  
      cex = 2, col = "purple",  
      pch = 19)  
    panel.text(170, -30,  
      "Hi there!",  
      pos = 3)  
  
    panel.xyplot(...)  
  })
```



The latticeExtra package

Normal plot:

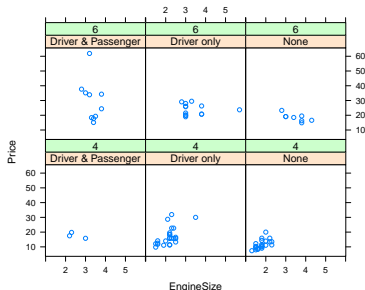
```
(pcars <-  
  xyplot(Price ~ EngineSize |  
    AirBags + Cylinders,  
    data = Cars93,  
    subset = Cylinders %in%  
      c(4, 6)) )
```



The latticeExtra package

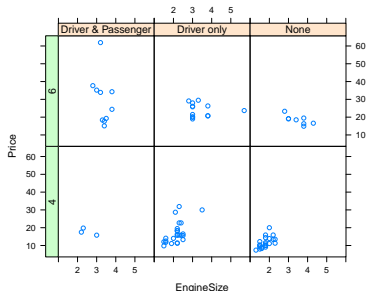
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    AirBags + Cylinders,  
    data = Cars93,  
    subset = Cylinders %in%  
      c(4,6)))
```



Now move one set of labels to the side...

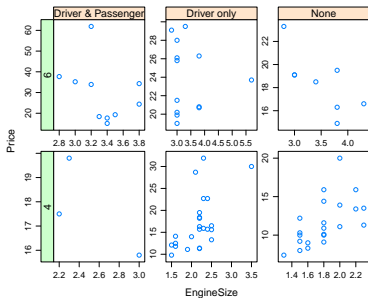
```
(pcars2 <-  
  useOuterStrips(pcars))
```



Combine axis ranges...

Every packet its own axes...

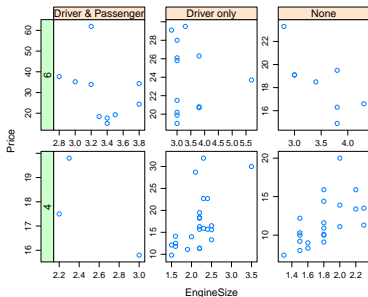
```
(pcars3 <-  
  update(pcars2,  
    scale = "free") )
```



Combine axis ranges...

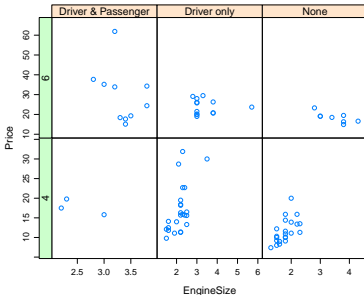
Every packet its own axes...

```
(pcars3 <-  
  update(pcars2,  
    scale = "free"))
```



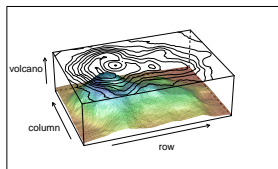
Combined in a meaningful way

```
combineLimits(pcars3)
```



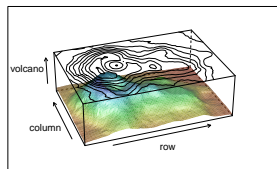
Conclusions

- You can do quite complicated things...



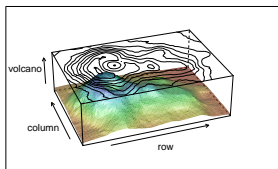
Conclusions

- ▶ You **can** do quite complicated things...
- ▶ Resist!



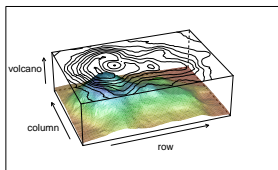
Conclusions

- ▶ You **can** do quite complicated things...
- ▶ Resist!
- ▶ Only if you **must**



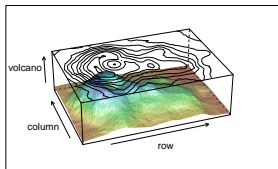
Conclusions

- ▶ You **can** do quite complicated things...
- ▶ Resist!
- ▶ Only if you **must**
- ▶ Which means: if it makes the plot better

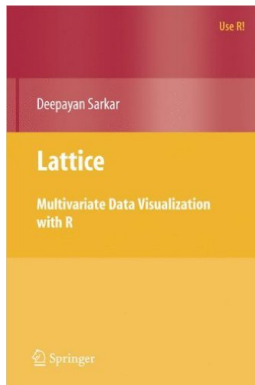


Conclusions

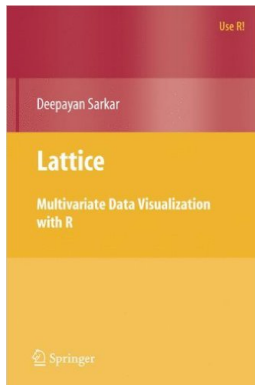
- ▶ You **can** do quite complicated things...
- ▶ Resist!
- ▶ Only if you **must**
- ▶ Which means: if it makes the plot better
- ▶ Not treated:
 - ▶ themes
 - ▶ grouping (e.g., **panel.superpose** instead of **panel.xyplot**)
 - ▶ labels and legends
 - ▶ axis coordinates and annotation
 - ▶ ...



Acknowledgements

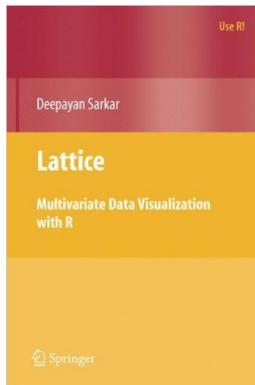


Acknowledgements



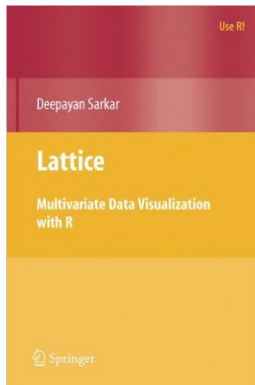
- `lmdvr.r-forge.r-project.org` contains all code to reproduce the book figures

Acknowledgements



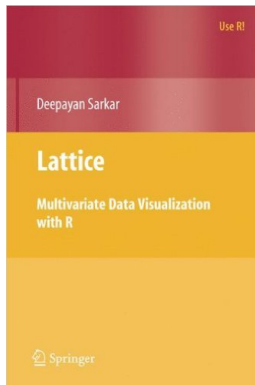
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