

Visualizing multivariate data using lattice and direct labels  
<http://directlabels.r-forge.r-project.org>

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# Outline

The lattice system

Adding direct labels using the latticedl package

## Brief history of lattice

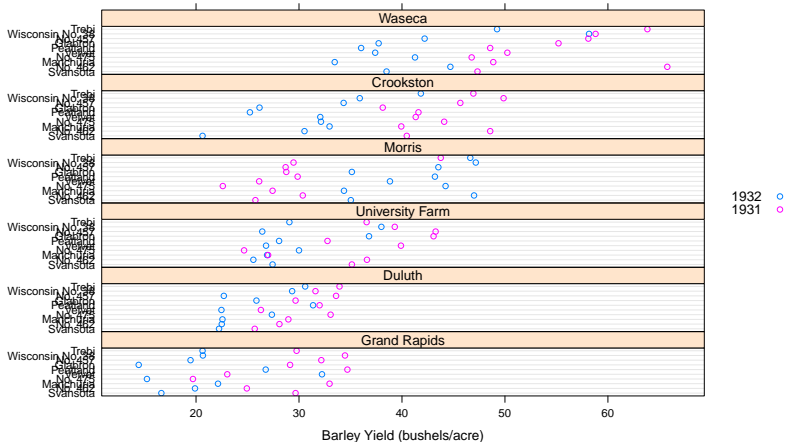
- ▶ Bill Cleveland, *Visualizing Data* (1993).
- ▶ Bill Cleveland, Rick Becker, Bell Labs, 1990s: trellis graphics system for S:  
http:  
[//cm.bell-labs.com/cm/ms/departments/sia/project/trellis/](http://cm.bell-labs.com/cm/ms/departments/sia/project/trellis/)
- ▶ Deepayan Sarkar, 2000s: the lattice package for R.
- ▶ Deepayan Sarkar (2008) *Lattice: Multivariate Data Visualization with R*, Springer.

## Installing the required packages

- ▶ I used R 2.9.2 for the following examples.
- ▶ lattice is preinstalled with R.
- ▶ `library(lattice)`
- ▶ `install.packages(c("latticeExtra", "latticed1"))`
- ▶ `library(latticeExtra)`
- ▶ `library(latticed1)`

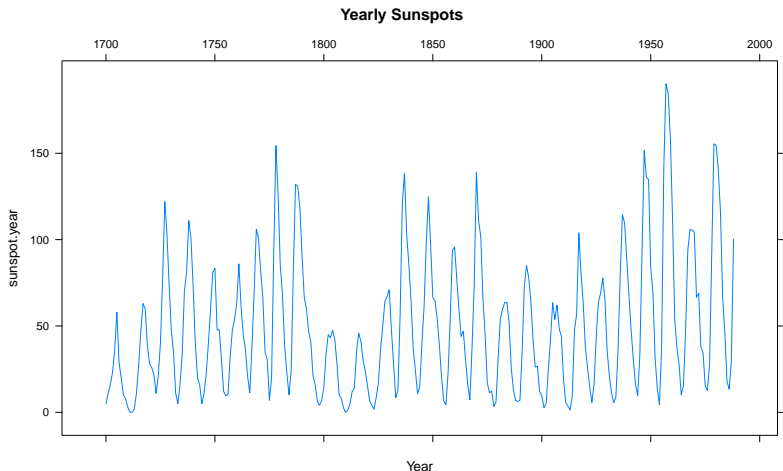
## Lattice allows easy visualization of many variables

```
> options(width = 55)
> library(lattice)
> dotplot(variety ~ yield | site, data = barley,
+         groups = year, auto.key = list(space = "right"),
+         layout = c(1, 6), xlab = "Barley Yield (bushels/acre)")
```



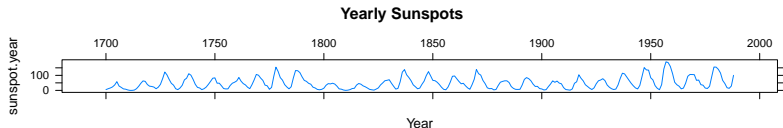
## Aspect ratio in scatterplots is important

```
> xyplot(sunspot.year ~ 1700:1988, xlab = "Year",  
+       type = "l", scales = list(x = list(alternating = 2)),  
+       main = "Yearly Sunspots")
```



## Lattice also automatically calculates aspect ratio for optimal decoding

```
> xyplot(sunspot.year ~ 1700:1988, xlab = "Year",  
+       type = "l", scales = list(x = list(alternating = 2)),  
+       main = "Yearly Sunspots", aspect = "xy")
```



## Load a data set

```
> data(Chem97, package = "mlmRev")  
> head(Chem97)
```

	lea	school	student	score	gender	age	gcsescore
1	1	1	1	4	F	3	6.625
2	1	1	2	10	F	-3	7.625
3	1	1	3	10	F	-4	7.250
4	1	1	4	10	F	-2	7.500
5	1	1	5	8	F	-1	6.444
6	1	1	6	10	F	4	7.750

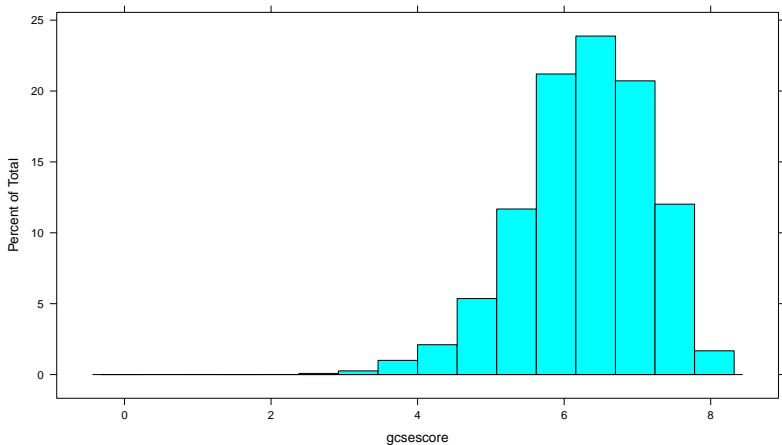
  

	gcsecnt
1	0.3393157
2	1.3393157
3	0.9643157
4	1.2143157
5	0.1583157
6	1.4643157



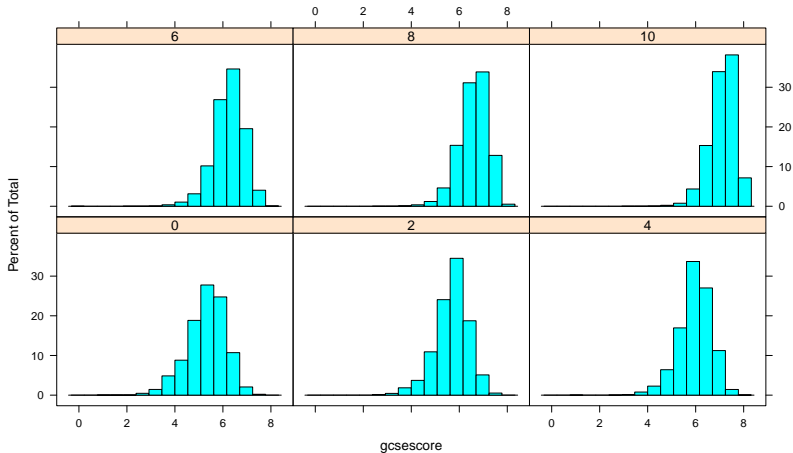
## Simple histogram

```
> histogram(~gcsescore, Chem97)
```



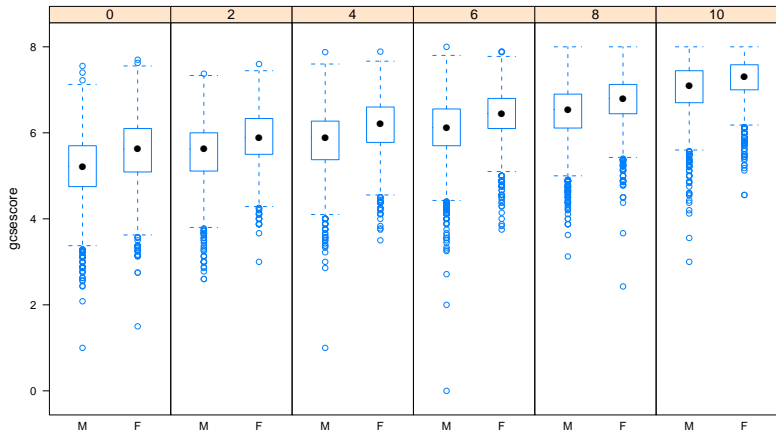
## Histograms conditional on a categorical variable

```
> histogram(~gcsescore | factor(score),  
+ Chem97)
```



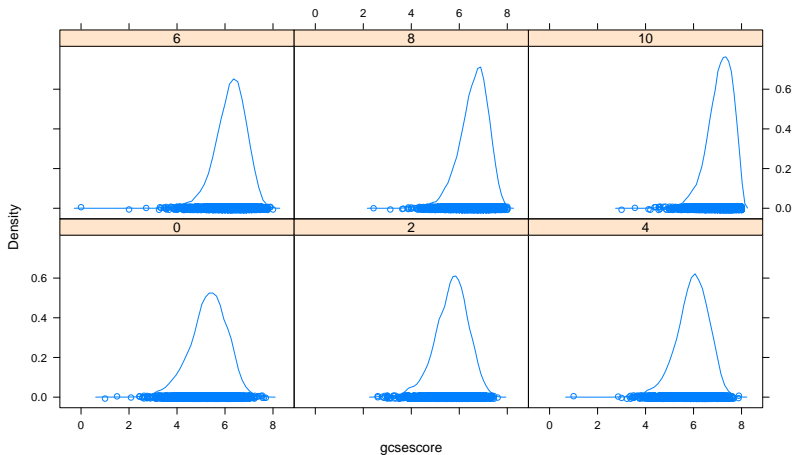
## Box and whisker plots

```
> bwplot(gcsescore ~ gender | factor(score),  
+       Chem97, layout = c(6, 1))
```



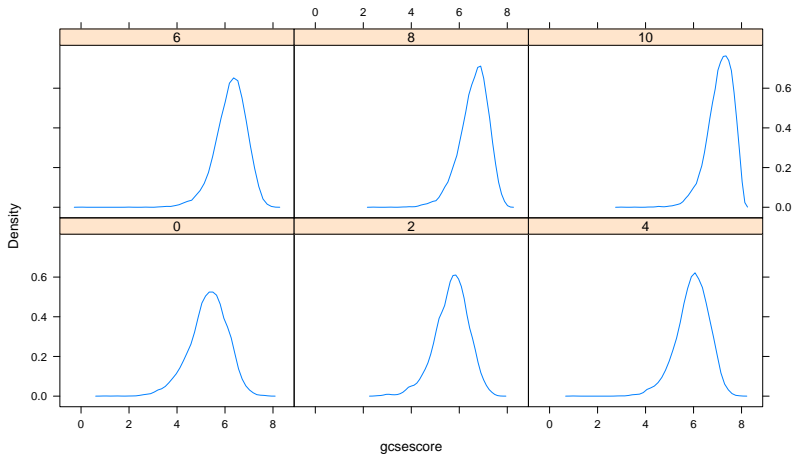
## Conditioned plots of kernel density estimates

```
> densityplot(~gcsescore | factor(score),  
+ Chem97)
```



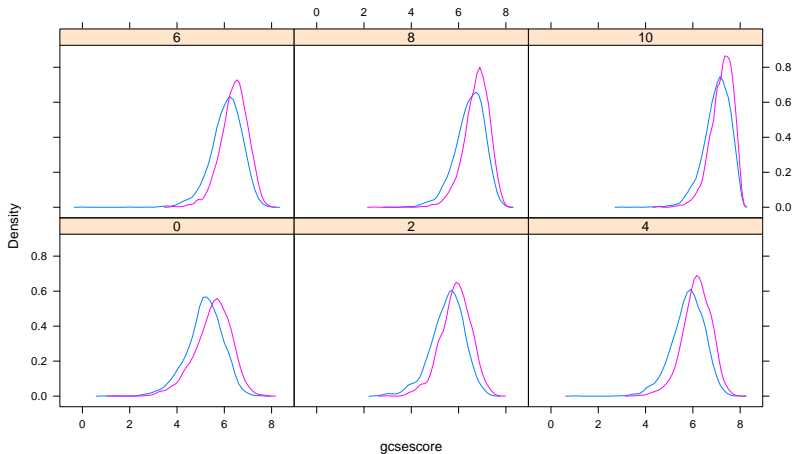
## Hide the actual points with the plot.points argument

```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE)
```



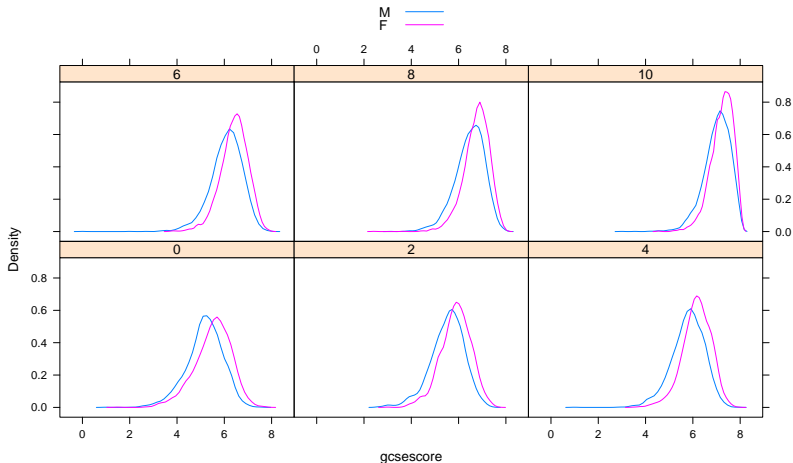
## Conditioned and grouped density plots

```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE, groups = gender)
```



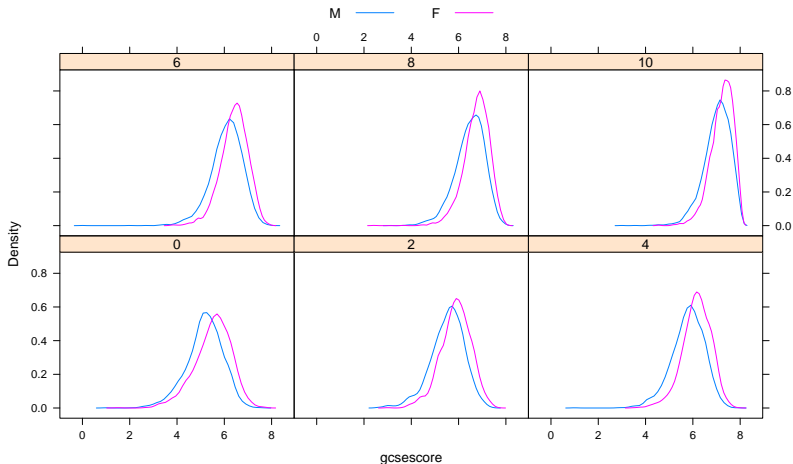
## Add a legend with the auto.key argument

```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE, groups = gender,  
+ auto.key = list())
```



## Legend layout with the columns argument

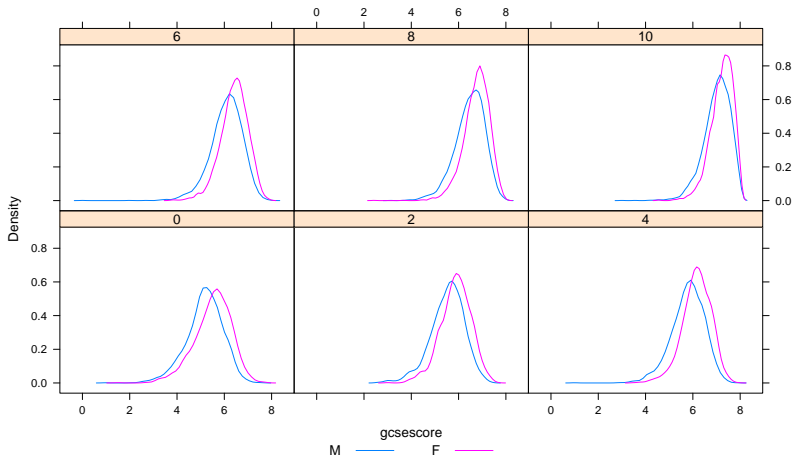
```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE, groups = gender,  
+ auto.key = list(columns = 2))
```





## Legend positioning with the space argument

```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE, groups = gender,  
+ auto.key = list(columns = 2, space = "bottom"))
```



# Show all default settings

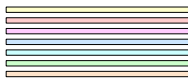
```
> show.settings()
```



superpose.symbol



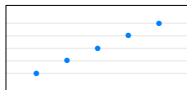
superpose.line



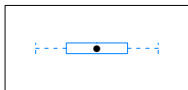
strip.background



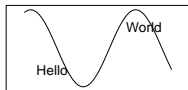
strip.shingle



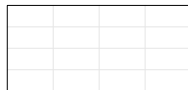
dot.[symbol, line]



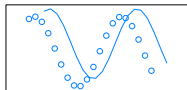
box.[dot, rectangle, umbrella]



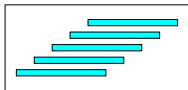
add.[line, text]



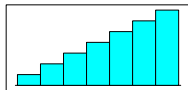
reference.line



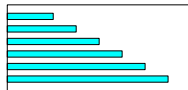
plot.[symbol, line]



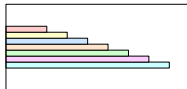
plot.shingle[plot.polygon]



histogram[plot.polygon]



barchart[plot.polygon]



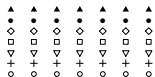
superpose.polygon



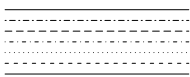
regions

## Show settings good for printout

```
> show.settings(standard.theme(color = FALSE))
```



superpose.symbol



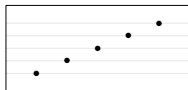
superpose.line



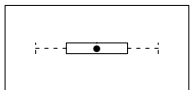
strip.background



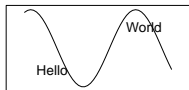
strip.shingle



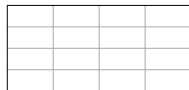
dot.[symbol, line]



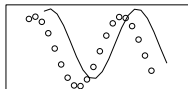
box.[dot, rectangle, umbrella]



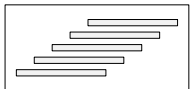
add.[line, text]



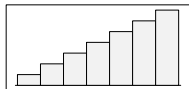
reference.line



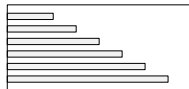
plot.[symbol, line]



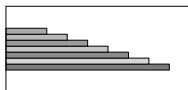
plot.shingle[plot.polygon]



histogram[plot.polygon]



barchart[plot.polygon]



superpose.polygon



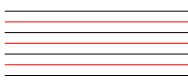
regions

## Change the settings

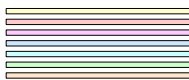
```
> br <- simpleTheme(col = c("black", "red"))  
> show.settings(br)
```



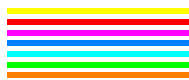
superpose.symbol



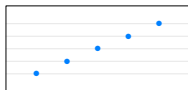
superpose.line



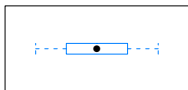
strip.background



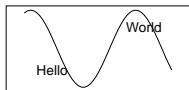
strip.shingle



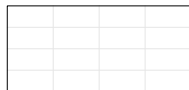
dot.[symbol, line]



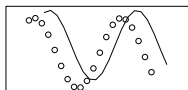
box.[dot, rectangle, umbrella]



add.[line, text]



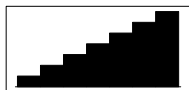
reference.line



plot.[symbol, line]



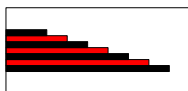
plot.shingle[plot.polygon]



histogram[plot.polygon]



barchart[plot.polygon]



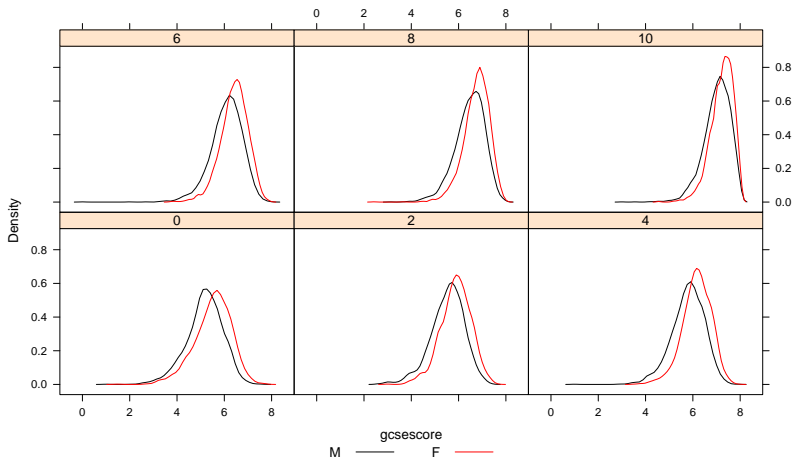
superpose.polygon



regions

## Change group colors with par.settings

```
> densityplot(~gcsescore | factor(score),  
+ Chem97, plot.points = FALSE, groups = gender,  
+ auto.key = list(columns = 2, space = "bottom"),  
+ par.settings = br)
```



## Load a tabular data set

```
> print(VADeaths)
```

	Rural Male	Rural Female	Urban Male	Urban Female
50-54	11.7	8.7	15.4	8.4
55-59	18.1	11.7	24.3	13.6
60-64	26.9	20.3	37.0	19.3
65-69	41.0	30.9	54.6	35.1
70-74	66.0	54.3	71.1	50.0

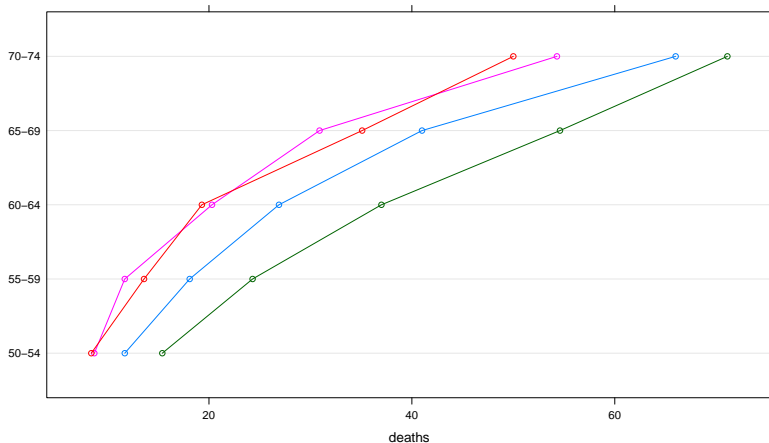
## Convert to data frame to work with lattice

```
> vad <- as.data.frame.table(VADeaths)
> names(vad) <- c("age", "demographic", "deaths")
> head(vad)
```

	age	demographic	deaths
1	50-54	Rural Male	11.7
2	55-59	Rural Male	18.1
3	60-64	Rural Male	26.9
4	65-69	Rural Male	41.0
5	70-74	Rural Male	66.0
6	50-54	Rural Female	8.7

## Grouped dotplots work well for these data

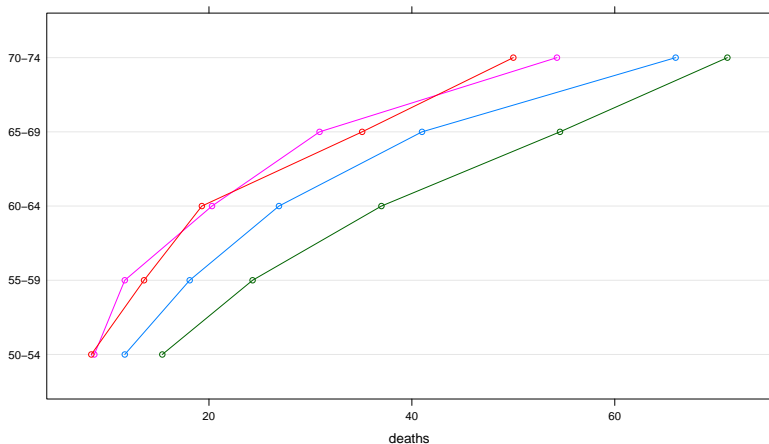
```
> dotplot(age ~ deaths, vad, groups = demographic,  
+         type = "o")
```





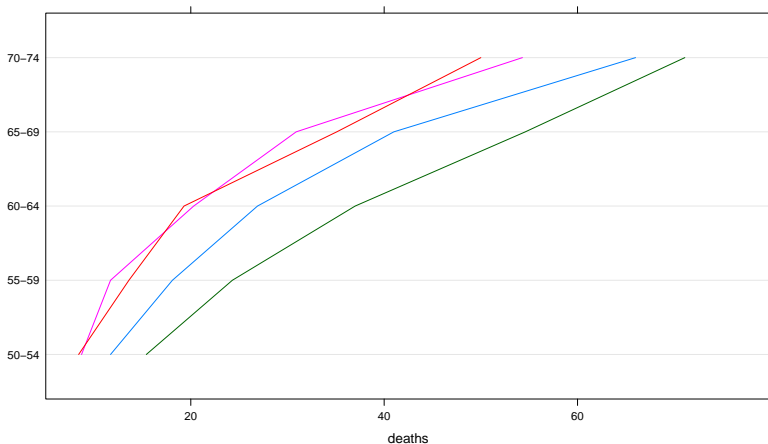
## Plots can be saved as R objects

```
> dots <- dotplot(age ~ deaths, vad, groups = demographic,  
+   type = "o")  
> dots
```



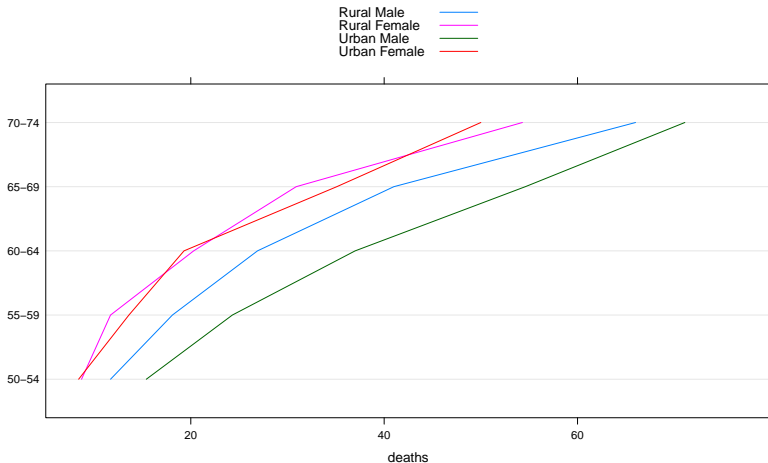
## Saved plots can be updated later

```
> dots2 <- update(dots, type = "l", xlim = c(5,  
+ 80))  
> dots2
```



## Add a confusing legend ... how can we label more intuitively?

```
> update(dots2, auto.key = list(points = FALSE,  
+       lines = TRUE))
```



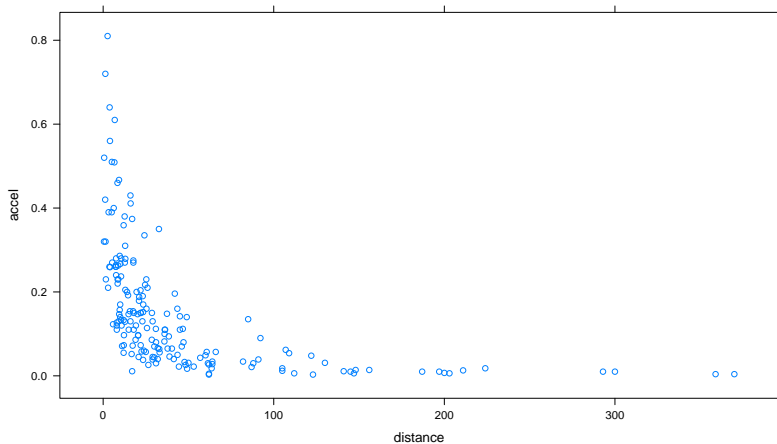
## Load some earthquake measurements

```
> data(Earthquake, package = "nlme")  
> head(Earthquake)
```

	Quake	Richter	distance	soil	accel
132	20	5	7.5	1	0.264
133	20	5	8.8	1	0.263
134	20	5	8.9	1	0.230
135	20	5	9.4	1	0.147
136	20	5	9.7	1	0.286
137	20	5	9.7	1	0.157

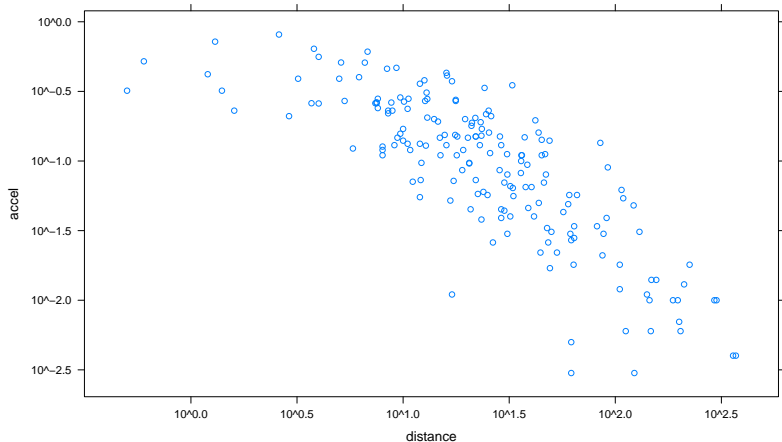
## Scatterplot with xyplot

```
> xyplot(accel ~ distance, Earthquake)
```



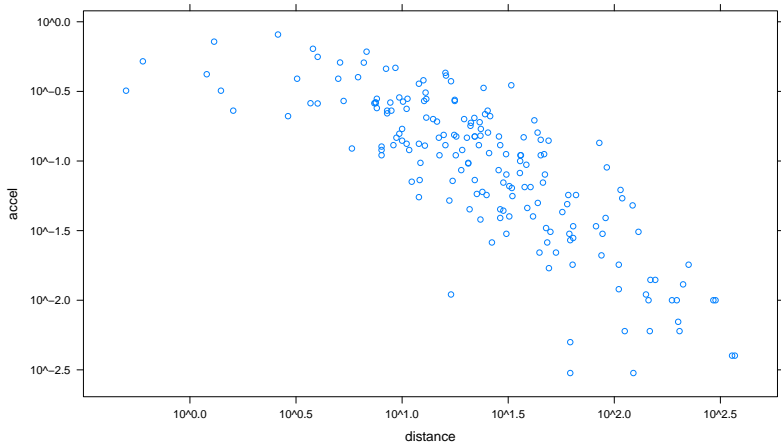
## Log scales with scales argument

```
> xyplot(accel ~ distance, Earthquake,  
+       scales = list(log = TRUE))
```



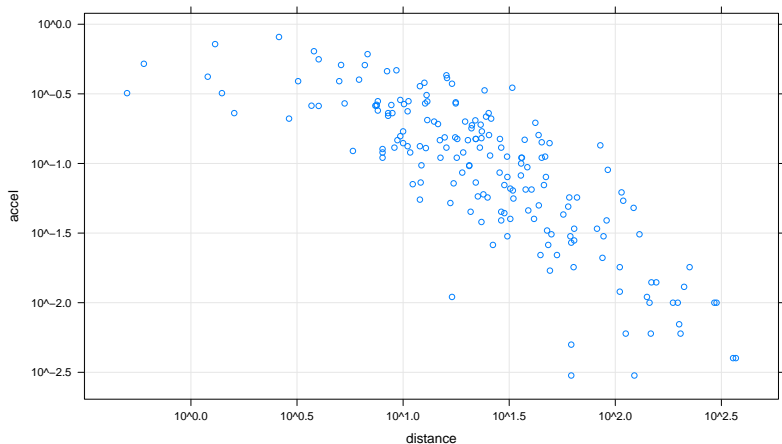
## Type "p" is the default

```
> xyplot(accel ~ distance, Earthquake,  
+       scales = list(log = TRUE), type = c("p"))
```



## Type "g" adds a grid

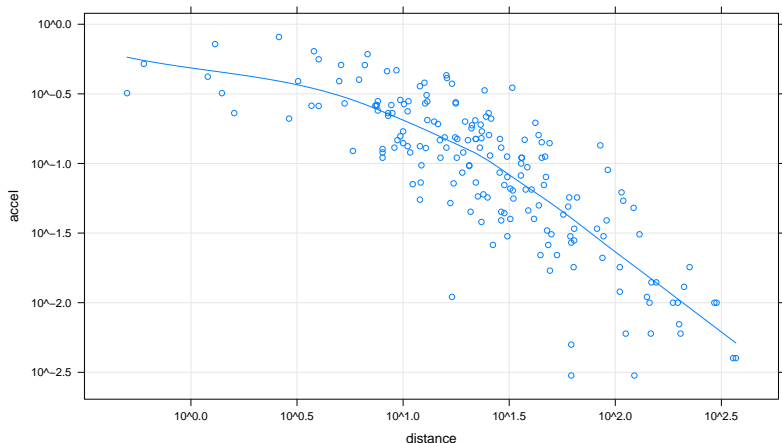
```
> xyplot(accel ~ distance, Earthquake,  
+       scales = list(log = TRUE), type = c("p",  
+       "g"))
```





## Type "smooth" adds a smooth line

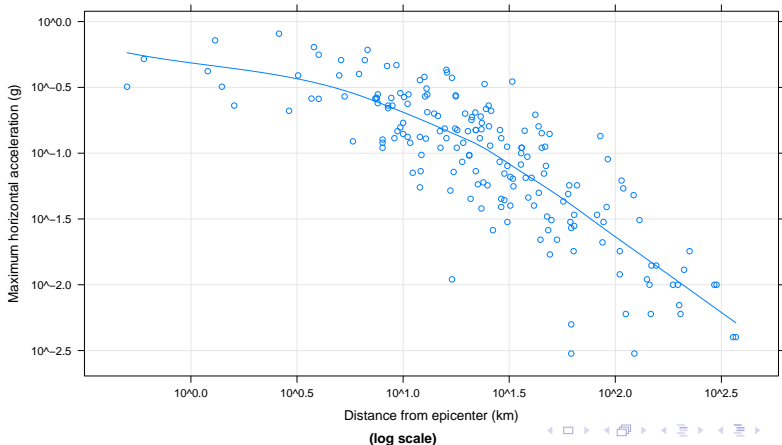
```
> xyplot(accel ~ distance, Earthquake,  
+       scales = list(log = TRUE), type = c("p",  
+       "g", "smooth"))
```



## Add some labels

```
> xyplot(accel ~ distance, Earthquake,  
+       scales = list(log = TRUE), type = c("p",  
+       "g", "smooth"), sub = "(log scale)",  
+       xlab = "Distance from epicenter (km)",  
+       ylab = "Maximum horizontal acceleration (g)",  
+       main = "Larger quakes are felt closer to the epicenter")
```

**Larger quakes are felt closer to the epicenter**



## Volcano elevation data in matrix form

```
> dim(volcano)
```

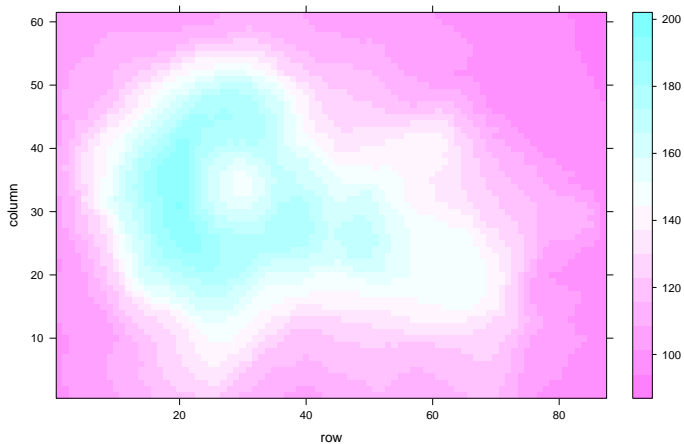
```
[1] 87 61
```

```
> print(volcano[1:5, 1:5])
```

	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]	100	100	101	101	101
[2,]	101	101	102	102	102
[3,]	102	102	103	103	103
[4,]	103	103	104	104	104
[5,]	104	104	105	105	105

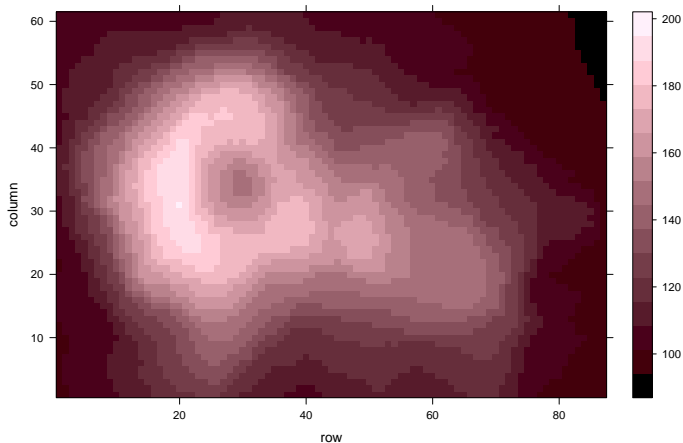
## Plot volcano elevations in a matrix using color

```
> levelplot(volcano)
```



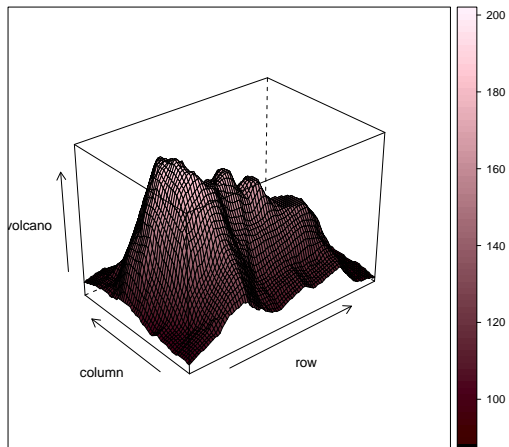
## Use a different color scale

```
> my.colors <- sapply(0:100, function(l) hcl(l = l))  
> levelplot(volcano, col.regions = my.colors)
```



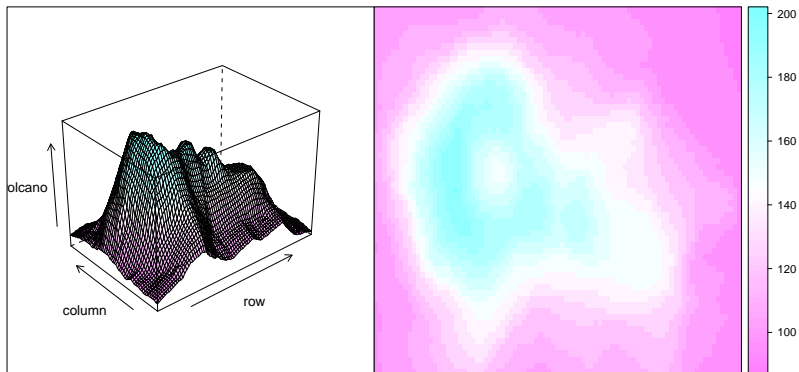
## Use 3d wireframe plots

```
> wireframe(volcano, drape = TRUE, col.regions = my.colors)
```



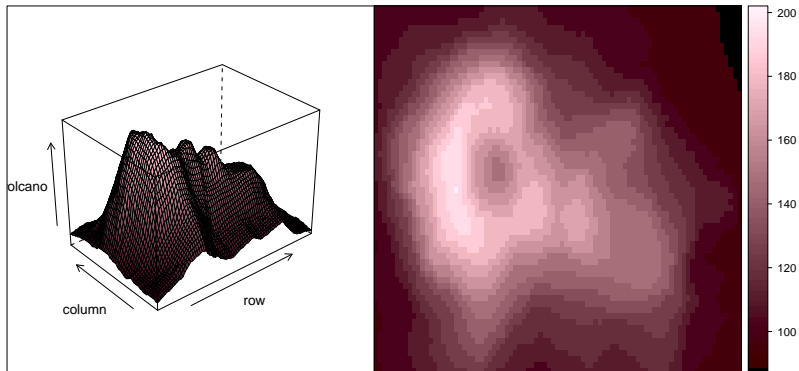
## Combine plots using latticeExtra

```
> library(latticeExtra)
> both <- c(wireframe(volcano, drape = TRUE),
+          levelplot(volcano))
> both
```



## Globally change the plot parameters

```
> trellis.par.set(regions = list(col = my.colors))  
> both
```





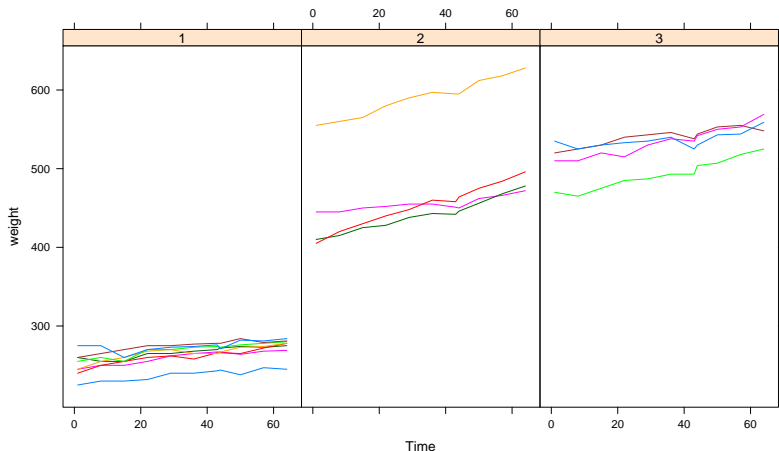
## Longitudinal data

```
> data(BodyWeight, package = "nlme")  
> head(BodyWeight)
```

	weight	Time	Rat	Diet
1	240	1	1	1
2	250	8	1	1
3	255	15	1	1
4	260	22	1	1
5	262	29	1	1
6	258	36	1	1

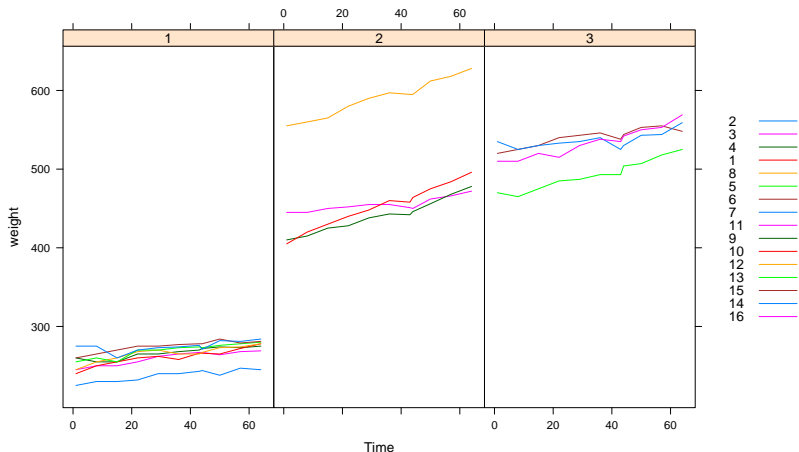
## Conditional scatterplots reveal difference between treatments

```
> xyplot(weight ~ Time | Diet, BodyWeight,  
+ groups = Rat, type = "l", layout = c(3,  
+ 1))
```



## Legends with more than a few items are very confusing

```
> xyplot(weight ~ Time | Diet, BodyWeight,  
+       groups = Rat, type = "l", layout = c(3,  
+       1), auto.key = list(space = "right",  
+       points = FALSE, lines = TRUE))
```



# Outline

The lattice system

Adding direct labels using the latticedl package

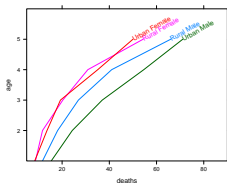


## How to plot direct labels in R?

- ▶ Lattice + latticedl: `direct.label(xyplot(y~x,data,groups=z),method=f)`
- ▶ Positions of direct labels can be specified as a function of the data:

```
f <- function(d,...){  
  # d is a data frame with columns x,y,groups of the data points  
  #... analyze the points and return the label positions:  
  return(data.frame(x=a,y=b,groups=c))  
}
```

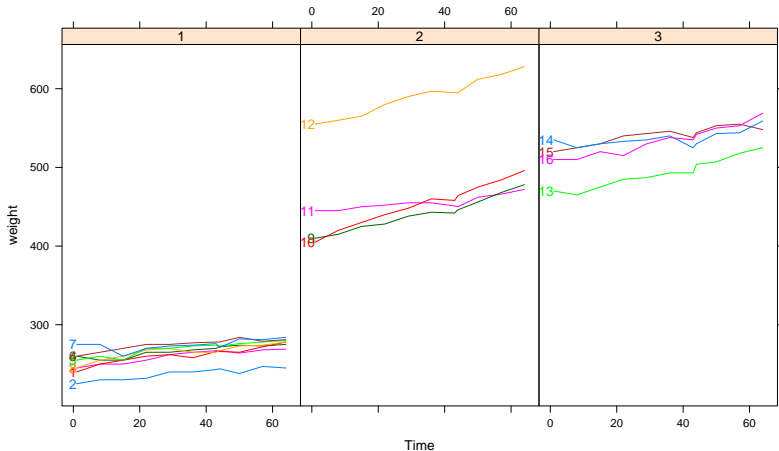
	groups	x	y	hjust	vjust	rot
1	Rural Male	66.0	5	0	0.5	30
2	Rural Female	54.3	5	0	0.5	30
3	Urban Male	71.1	5	0	0.5	30
4	Urban Female	50.0	5	0	0.5	30



- ▶ `latticedl` does the labeling for you, keeping track of the correct colors.
- ▶ Common plot types have default direct labeling methods.

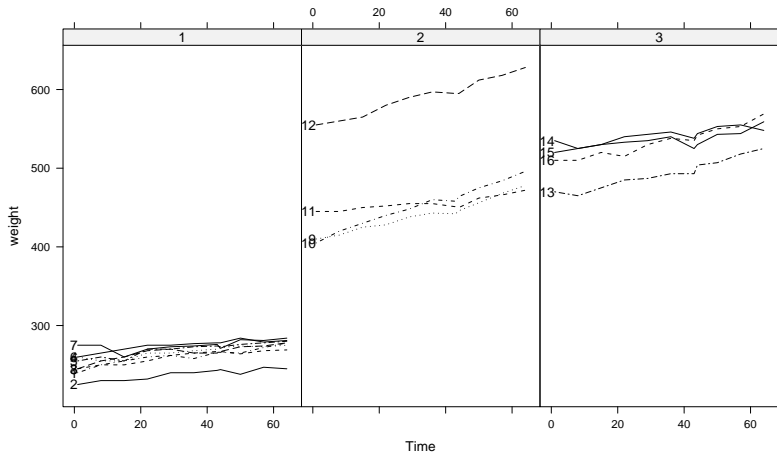
## Easy fix for confusing legend: direct labels

```
> library(latticedl)
> long <- xyplot(weight ~ Time | Diet, BodyWeight,
+   groups = Rat, type = "l", layout = c(3,
+   1))
> direct.label(long)
```



## Even works in black and white

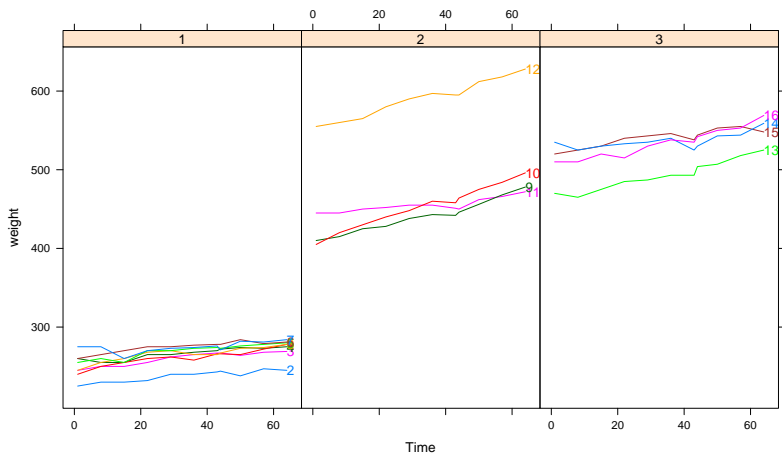
```
> longbw <- update(long, par.settings = standard.theme(color = FALSE))  
> direct.label(longbw)
```





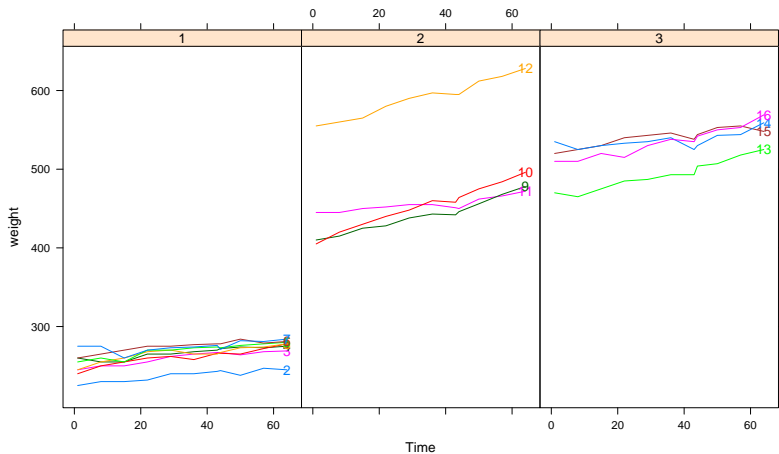
## Change label positions with the method argument

```
> direct.label(long, method = last.points)
```



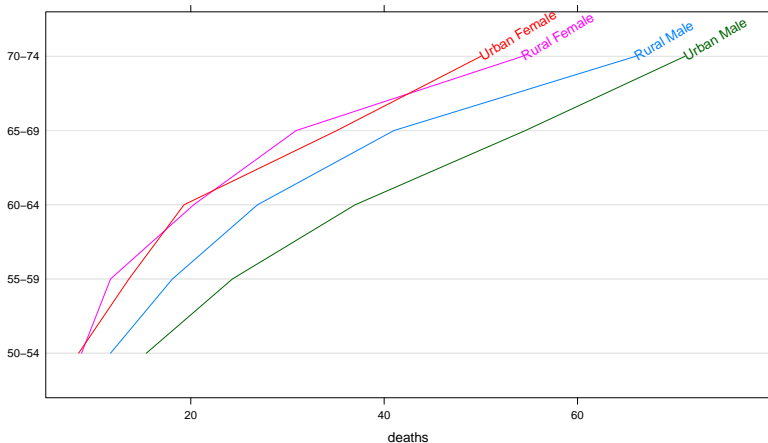
## Make your own positioning function using dl.indep

```
> direct.label(long, method = dl.indep(d[which.max(d$x),  
+   ]))
```



You can change text parameters (same as `grid::grid.text`)

```
> direct.label(dots2, method = list("last.points",  
+   rot = 30))
```



## Load some data on car fuel efficiency

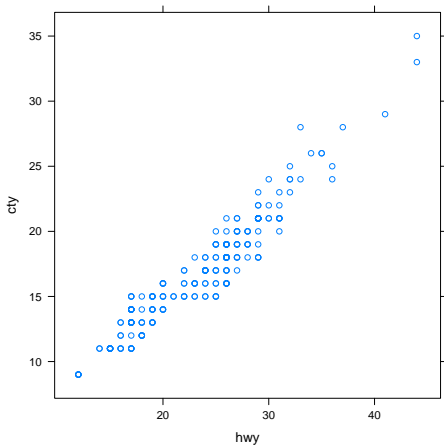
```
> data(mpg, package = "ggplot2")  
> head(mpg)
```

	manufacturer	model	displ	year	cyl	trans	drv	cty
1	audi	a4	1.8	1999	4	auto(l5)	f	18
2	audi	a4	1.8	1999	4	manual(m5)	f	21
3	audi	a4	2.0	2008	4	manual(m6)	f	20
4	audi	a4	2.0	2008	4	auto(av)	f	21
5	audi	a4	2.8	1999	6	auto(l5)	f	16
6	audi	a4	2.8	1999	6	manual(m5)	f	18

	hwy	fl	class
1	29	p	compact
2	29	p	compact
3	31	p	compact
4	30	p	compact
5	26	p	compact
6	26	p	compact

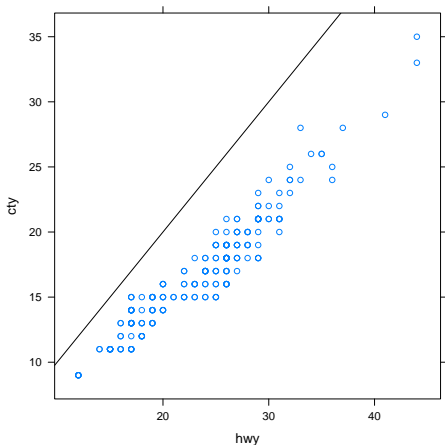
## Plot city versus highway fuel efficiency

```
> xyplot(cty ~ hwy, mpg, aspect = 1)
```



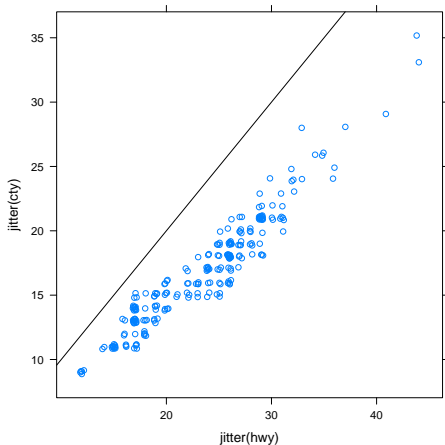
## Add a reference line $x=y$

```
> panel.xyref <- function(...) {  
+   panel.xyplot(...)  
+   panel.abline(0, 1)  
+ }  
> xyplot(cty ~ hwy, mpg, aspect = 1, panel = panel.xyref)
```



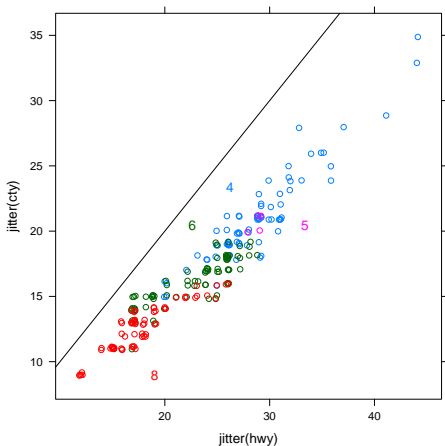
## Jitter the data to see all the points

```
> xyplot(jitter(cty) ~ jitter(hwy), mpg,  
+       aspect = 1, panel = panel.xyref)
```



## Group data by number of cylinders in the engine

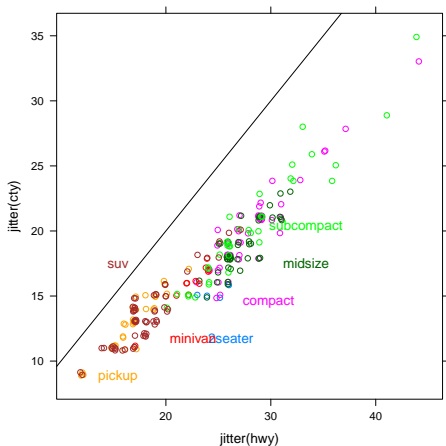
```
> direct.label(xyplot(jitter(cty) ~ jitter(hwy),  
+   mpg, aspect = 1, panel = panel.xyref,  
+   groups = factor(cyl)))
```





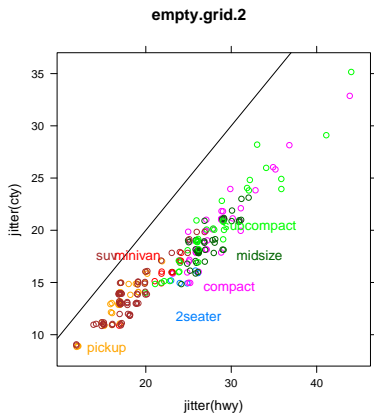
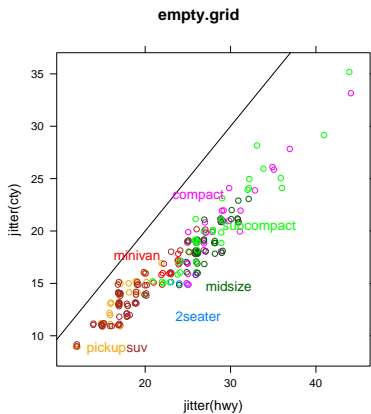
## Group data by car class

```
> direct.label(xyplot(jitter(cty) ~ jitter(hwy),  
+   mpg, aspect = 1, panel = panel.xyref,  
+   groups = class))
```



## Compare direct labeling methods

```
> compare.methods(c("empty.grid", "empty.grid.2"),  
+   xyplot, mpg, jitter(cty) ~ jitter(hwy),  
+   class, aspect = 1, panel = panel.xyref,  
+   horiz = TRUE)
```



## References for learning more about lattice and latticedl

- ▶ First load the libraries in R
  - ▶ `library(lattice)`
  - ▶ `library(latticeExtra)`
  - ▶ `library(latticedl)`
- ▶ Then you can look at the interactive help pages
  - [Overview](#) `?Lattice`
  - [Customizing plots](#) `?xyplot`
  - [Included panel functions](#) `?panel.functions`, `?llines`
  - [Multiple plots per page](#) `?plot.trellis`, `?c.trellis`
  - [Direct labeling](#) `?direct.label`
- ▶ Deepayan Sarkar (2008) *Lattice: Multivariate Data Visualization with R*, Springer.
- ▶ R code from the slides available on the web:  
<http://directlabels.r-forge.r-project.org>
- ▶ Email me directly: `toby.hocking AT inria.fr`