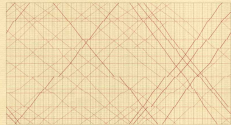


# data visualization

skimming deep waters



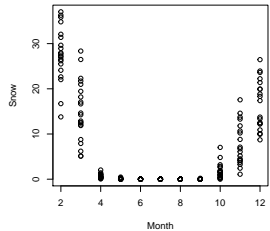
Flickr/Dejvy Gabburkinman



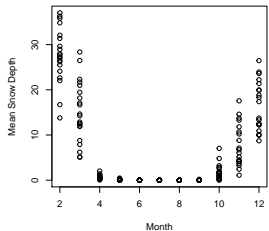
## The Visual Display of Quantitative Information

EDWARD R. TUFTE

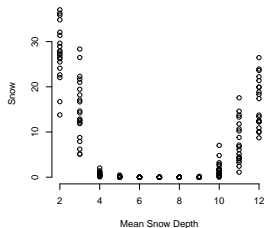
### Show the Data



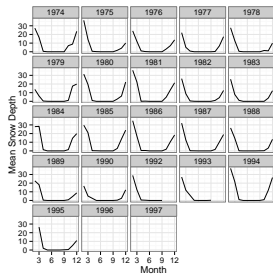
## Minimize Distraction



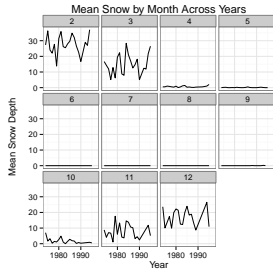
## Minimize Distraction



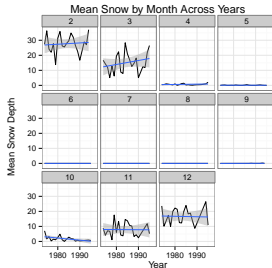
## Make Big Data Coherent



## Reveal Several Levels of Detail



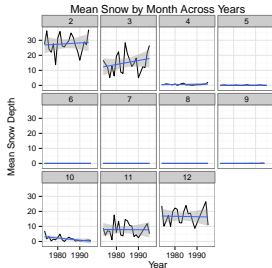
## Be Closely Integrated with Statistics



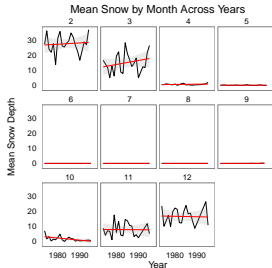
## The Data:Ink Ratio

1. Above all else show data.
2. Maximize the data-ink ratio.
3. Erase non-data-ink.
4. Erase redundant data-ink.
5. Revise and edit

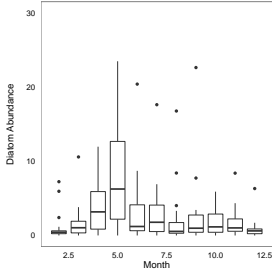
## Minimizing Ink



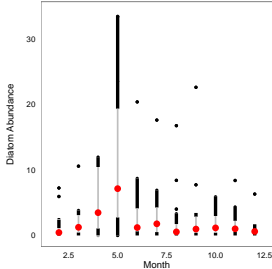
## Minimizing Ink



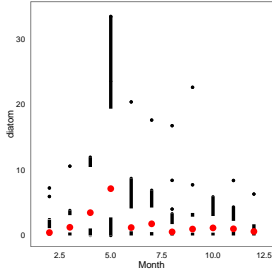
### Extra Ink in Boxes & Lines



### A Cleaner Boxplot



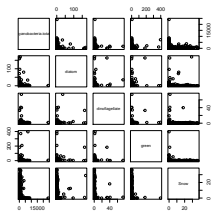
### Pure Tufte Boxplots



### Basic Plotting in R

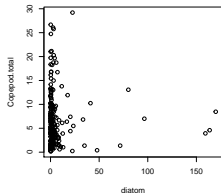
## Visualizing a Lot of the Data

```
pairs(plankton[, 14:18])
```



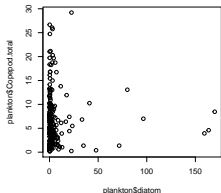
## A Basic Bivariate Plot

```
plot(Copepod.total ~  
      diatom, data = plankton)
```



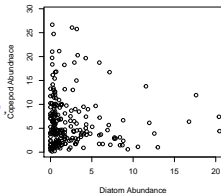
## A Basic Bivariate Plot

```
plot(plankton$diatom,  
      plankton$Copepod.total)
```



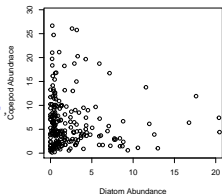
## Adding Axis Labels

```
plot(Copepod.total ~  
      diatom, data = plankton,  
      xlab = "Diatom Abundance",  
      ylab = "Copepod Abundance",  
      xlim = c(0, 20))
```



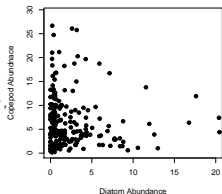
## Adding Axis Limits

```
plot(Copepod.total ~  
  diatom, data = plankton,  
  xlab = "Diatom Abundance",  
  ylab = "Copepod Abundance",  
  xlim = c(0, 20))
```



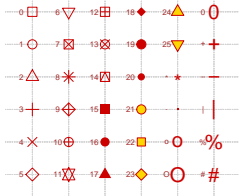
## More Point Shapes

```
plot(Copepod.total ~  
  diatom, data = plankton,  
  xlab = "Diatom Abundance",  
  ylab = "Copepod Abundance",  
  xlim = c(0, 20),  
  pch = 19)
```



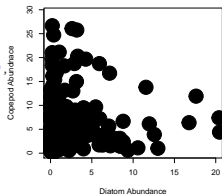
## More Point Shapes

plot symbols : points (... pch = \*, cex = 3)



## cex for Size

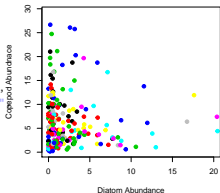
```
plot(Copepod.total ~  
  diatom, data = plankton,  
  xlab = "Diatom Abundance",  
  ylab = "Copepod Abundance",  
  xlim = c(0, 20),  
  pch = 19, cex = 4)
```



See also `cex.axis`,  
`cex.lab`, and more.

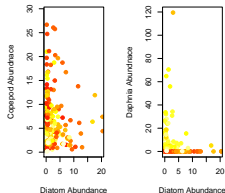
## Add a Little Color

```
plot(Copepod.total ~  
      diatom, data = plankton,  
      xlab = "Diatom Abundance",  
      ylab = "Copepod Abundance",  
      xlim = c(0, 20),  
      pch = 19, col = Month)
```



## Panels with Par and Mfrow

```
par(mfrow = c(1, 2))
```



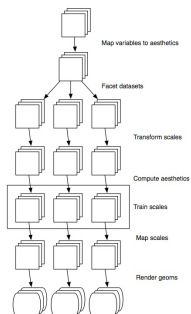
## Lots of Other Functions that For Plots

```
?matplot  
?lines  
?axis  
?title  
?legend  
?points  
?segments
```

So...Explore! Plot with the data, try different par settings, or use some of these functions!

## ggplot2

or how I learned to stop worrying and love <http://had.co.nz/ggplot2> & <http://stackoverflow.com/>



## Start with nothing...

```
p <- ggplot(data = plankton, mapping = aes(x = Month,
  y = Copepod.total))
```

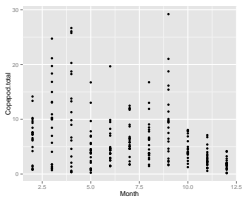
```
p
```

```
## Error: No layers in plot
```

There is no layout specified here for the data.

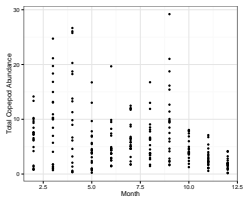
## Add a Layer

```
p <- p + geom_point()
p
```



## Format with Theme

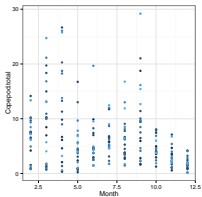
```
p <- p + ylab("Total Copepod Abundance") + theme_bw()
p
```





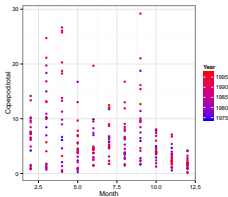
## Map a Variable to Color

```
p2 <- ggplot(data = plankton, aes(x = Month, y = Copepod.total,
  color = Year)) + geom_point() + theme_bw()
p2
```



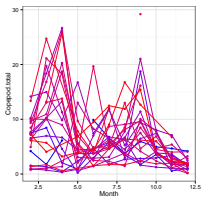
## Set Your Own Scale

```
p2 <- p2 + scale_color_gradient(low = "blue", high = "red")
p2
```



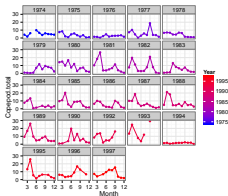
## And Maybe Add Another Layer

```
p2 <- p2 + geom_line(aes(group = Year))
p2
```



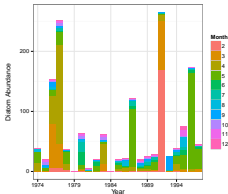
## Facet for Easier Visualization

```
p2 <- p2 + facet_wrap(~Year) + scale_x_continuous(breaks = c(3,
  6, 9, 12))
p2
```



## This All Can Lead to Interesting Visualizations

```
qplot(factor(Year), diatom, geom = "bar", fill = factor(Month),
      data = plankton) + theme_bw() + xlab("Year") +
  ylab("Diatom Abundance") + scale_fill_discrete(name = "Month") +
  scale_x_discrete(breaks = seq(1974, 1997, 5))
```



## Lots of Layers to Add to ggplot2 Objects

```
?theme
?labs
?xlim
?facet_grid
?scale_x_log10
?geom_histogram
?geom_ribbon
?geom_linerange
?geom_freqpoly
```

So...Explore!

Also, see <http://had.co.nz/ggplot2> for some examples