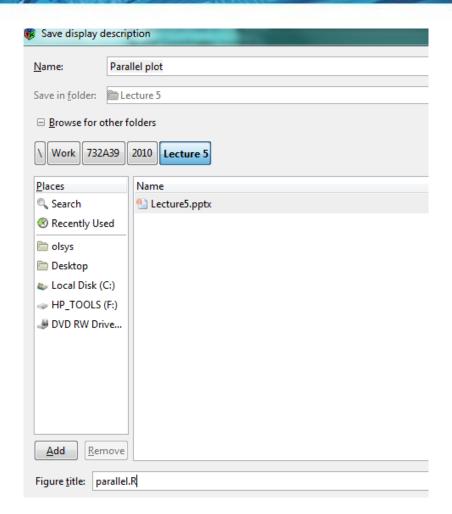


Publication quality graphics

 Use Tools-> Save Display description

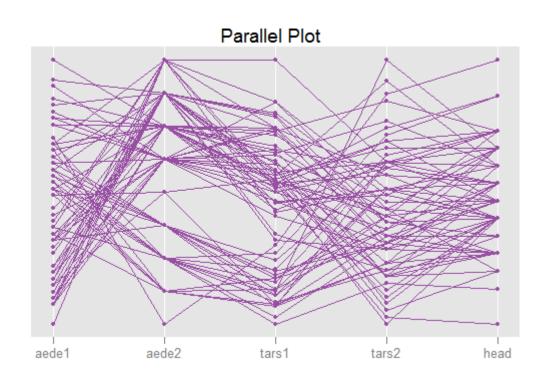


Publication quality graphics

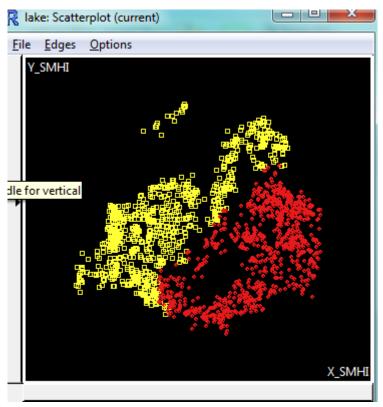
• In R:

```
library(DescribeDisplay);
d<-dd_load("parallel.R");
ggplot(d);</pre>
```

Using R plot, you can further enhance it in Inkscape



Seeing R data using GGobi



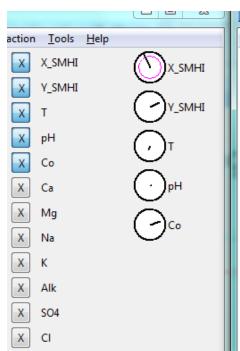
```
> lake <- read.csv("lakesurvey.csv");
> lake
     X SMHI Y SMHI
                                7.49 0.262
                                                    0.23
                               10.60 0.477
                                                    0.24
     654353 162104 5.4 7.46
                                            0.278
                                                    0.32
                               13.50 0.742
     654646 164746 5.5 7.52
                               15.40 0.872
                                            0.240
                                                    0.44
     654804 159298
                   3.8 7.60
                               13.70 0.720
                                            0.410
                                                    0.25
                               16.20 0.733
                    4.0 7.67
                                            0.512
                                                    0.39
                    2.7 6.77
                               16.70 0.379
                                            0.224
                                                    0.92
                                9.18 0.471
                                            0.173
                    5.8 7.32
                                                    0.21
                                            0.066
```

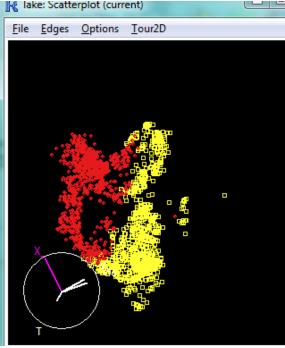
```
lake<-read.csv("lakesurvey.csv")
library(rggobi);
gg<-ggobi(lake);
glyph_color(gg[1])<-c(rep(1,800), rep(2,869))
glyph_type(gg[1])<-c(rep(5,800), rep(6,869))</pre>
```

Seeing R using Ggobi

Tours

```
> d <- displays(gg)[[1]]</p>
> pmode(d) <- "2D Tour"
> ggobi display get tour projection(d)
               [,1]
                              [,2]
 [1,] -6.884923e-06 1.365615e-05
 [2,]
       1.247204e-05
                     7.038482e-06
 [3,] -1.091227e-02 -1.702719e-02
      2.080103e-02 -3.868532e-03
 [4,]
 [5,]
      2.632767e-04 1.008248e-04
 [6,]
       0.000000e+00
                     0.000000e+00
 [7,]
       0.000000e+00
                     0.000000e+00
       0.000000e+00
 [8,]
                     0.000000e+00
 [9,]
       0.000000e+00
                     0.000000e+00
       0.000000e+00
                     0.000000e+00
[10,]
       0.000000e+00
                     0.000000e+00
[11,]
[12,]
       0.000000e+00
                     0.000000e+00
[13,]
       0.000000e+00
                     0.000000e+00
       0.000000e+00
                      0.000000e+00
[14,]
```



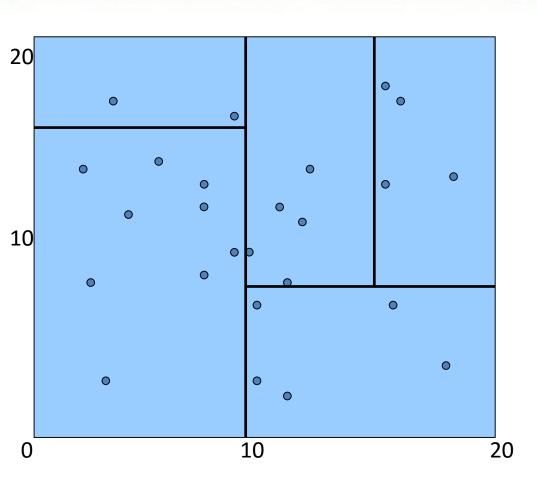


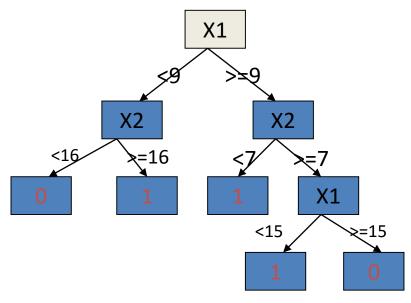
Visualizing and numerical methods

Data mining methods, aims:

- Classification (X,Y given)
 - Linear Discriminant analysis
 - Decision trees
 - Neural networks
 - **—** ...

- Clustering (X given)
 - Hierarchical clustering
 - Kohonen maps
 - **—** ...





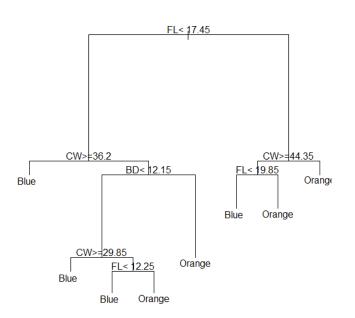
```
+ > crabs<-read.csv("australian-crabs.csv")</p>
> crabs
   species
           sex index
                      FL
                           RW
                                CL
                                         BD
1
     Blue
           Male
                    1 8.1 6.7 16.1 19.0
                                        7.0
     Blue Male
                    2 8.8 7.7 18.1 20.8 7.4
3
     Blue Male
                    3 9.2 7.8 19.0 22.4 7.7
4
     Blue Male
                    4 9.6 7.9 20.1 23.1
                                        8.2
5
    Blue Male
                    5 9.8 8.0 20.3 23.0 8.2
6
     Blue Male
                    6 10.8 9.0 23.0 26.5 9.8
     Blue
           Male
                    7 11.1 9.9 23.8 27.1 9.8
```

```
crabs<-read.csv("australian-crabs.csv");
library(rpart);
tree<-rpart(species~FL+RW+CL+CW+BD, data=crabs);</pre>
```

```
> librarv(rpart)
> tree<-rpart(species~FL+RW+CL+CW+BD, data=crabs)
> tree
n = 200
node), split, n, loss, yval, (yprob)
      * denotes terminal node
 1) root 200 100 Blue (0.50000000 0.50000000)
   2) FL< 17.45 135 47 Blue (0.65185185 0.34814815)</p>
     4) CW>=36.2 40 4 Blue (0.90000000 0.10000000) *
     5) CW< 36.2 95 43 Blue (0.54736842 0.45263158)
      10) BD< 12.15 62 13 Blue (0.79032258 0.20967742)
        20) CW>=29.85 21 0 Blue (1.00000000 0.00000000) *
        21) CW< 29.85 41 13 Blue (0.68292683 0.31707317)
          42) FL< 12.25 34 6 Blue (0.82352941 0.17647059) *
         43) FL>=12.25 7 0 Orange (0.00000000 1.00000000) *
      11) BD>=12.15 33 3 Orange (0.09090909 0.90909091) *
   3) FL>=17.45 65 12 Orange (0.18461538 0.81538462)
     6) CW>=44.35 33 12 Orange (0.36363636 0.63636364)
     12) FL< 19.85 11 0 Blue (1.00000000 0.00000000) *
     13) FL>=19.85 22 1 Orange (0.04545455 0.95454545) *
     7) CW< 44.35 32 0 Orange (0.00000000 1.00000000) *
```

Plotting the tree:

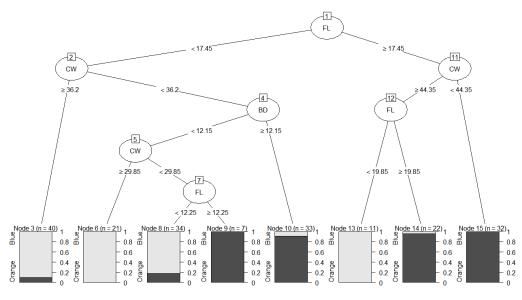
```
plot(tree);
text(tree);
```



• A better visualization:

Use library partykit

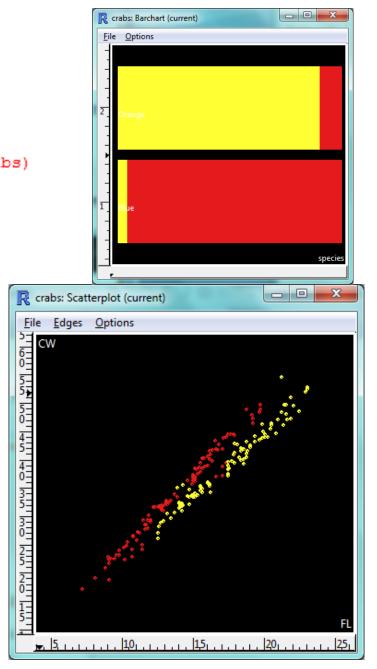
```
plot(as.party(tree))
```



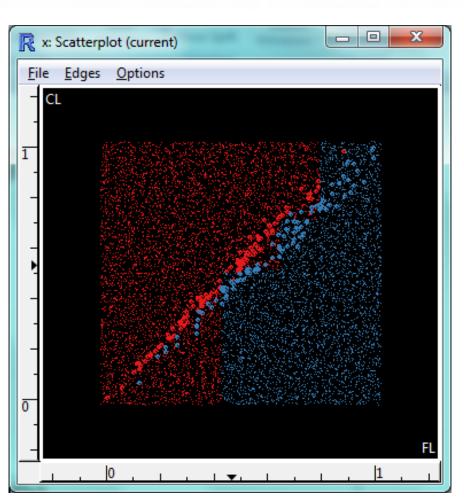
```
> probabilities
    Blue    Orange
1    0.82352941    0.1764706
2    0.82352941    0.1764706
3    0.82352941    0.1764706
4    0.82352941    0.1764706
5    0.82352941    0.1764706
6    0.82352941    0.1764706
7    0.82352941    0.1764706

crabspredict<-
    probabilities[1,]>probabilities[2,];
gg<-ggobi(crabs);
glyph_color(gg[1])<-crabspredict+1;</pre>
```

SEE 2D-tour also



Seeing boundaries



Use "classifly" for different methods (decision trees, neural networks, LDA)

- 1. Close all Ggobi windows
- 2. Run:

```
library(classifly);
classifly(crabs, species~FL+CL+RW, rpart);
```

Clustering

 To find groups of object that are close to each other and separated from other groups

Hierarchical clustering:

- 1. Put each observation in a cluster
- 2. Merge 2 nearest clusters into 1 cluster
- 3. Repeat 2 until you get 1 cluster

Concepts:

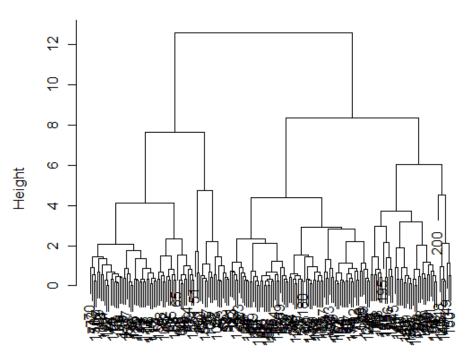
- Dendrogram
- Threshold
- → Use the given shreshold to define the amount of clusters.

Clustering

See dendrogram

x<-data.frame(crabs\$CL, crabs\$RW); crabs_dist<-dist(x); crabs_dend<-hclust(crabs_dist, method="average") plot(crabs_dend);</pre>

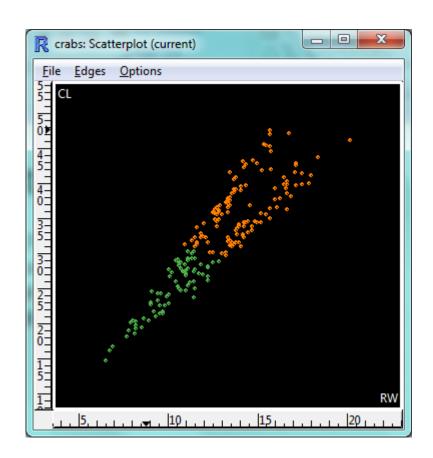
Cluster Dendrogram



crabs_dist hclust (*, "average")

Clustering

```
gd<-gg[1];
clust2=cutree(crabs_dend, k=2);
glyph_color(gd)[clust2==1]<-4;
glyph_color(gd)[clust2==2]<-5;</pre>
```



Dynamic plots

Interactive and dynamic plots

- Interactive= user can manipulate the plot to change it
- Dynamic= user is observing a moving picture
- Interactive:
 - All GGobi plots
 - Dynamic
 - Tours
 - Motion charts

Animation: good or bad?

Advantages

- Effective at attracting attention
- Same object can be used at another time point → visualizing one more dimension with same graphs (time)
- Easily perceived in peripheral vision → many features can be captured at one time point

Disadvantages

- Can be powerful force for destraction
- Unappropriate transformation (transition) → false conclusions
- Speed of the graphics may hide important details /make it too boring (unefficient)

Animation: recommendations

- Maintain valid data during transitions
 - Example: using splines
- Careful when using interpolations: use appropriate models
- Group similar transitions
- Minimize occlusion
- Use simple transistions
- If trajectory is stable, use slow-in, slow-down
- Make transition as long as needed but no longer.

- See http://www.gapminder.org
- Gapminder is
 - A database that stores important world statistics (by country)
 - Each dataset can be analysed online with Motion Chart
 - Offline version can be downloaded
- 1. Go to Gapminder
- 2. Click Data and look through available datasets
- 3. Choose set "Life expectancy (years)"
- 4. Run and analyse Motion Chart



- Comment available tools:
 - Axes
 - Lin/log
 - Time scale
 - Size indicator and bubble size
 - Color (how countries are marked)
 - Select countries
 - Identify (move mouse over a bubble)
 - Play
 - Speed of video
 - Trails
 - Chart/Map
 - Full screen
 - Opacity (visibility of the non-selected countries)
 - Zoom
 - How to use

- Life expectancy set
 - How the minimum and maximum life expectancies changed after 200 years
 - Which countries had in general the highest life expectancy? Lowest lifexpectancy? Highest income level? Lowest income level?
 - What happened after 1946 and why
 - Compare development of China and USA (mark this countries and use "trace")

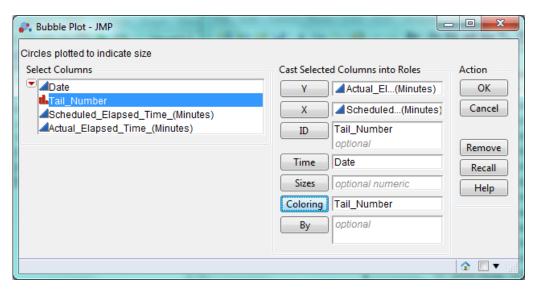
Motion Chart - JMP

- Excel file should contain
 - 1 column= observation name (country)
 - 1 column= time
 - 2 column= X och Y variables
 - More columns= Region, Bubble size

- SAS JMP:
 - − Graph → Bubble plot

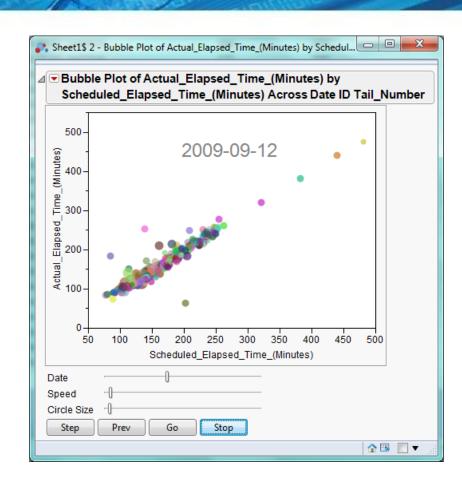
Example

Delays of the airlines



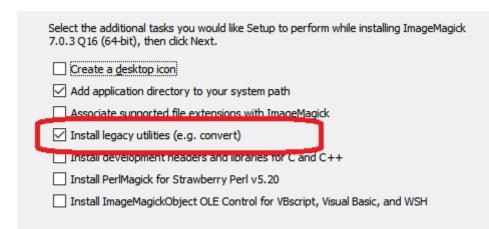
Motion Chart - JMP

- See which airlines delay or come earlier (deviations Y=X)
- Mark one or more bubbles and then use "Trail bubbles" or "Trail lines"
- Show legend
- Choose speed, pause



Dynamic from static

- Dynamic graphs can be created by combining several static graphs
- Install ImageMagick to make GIF animation
 - Mark option "Install legacy utilities"
- Download FFMPEG for video files
- Package animation



Animation

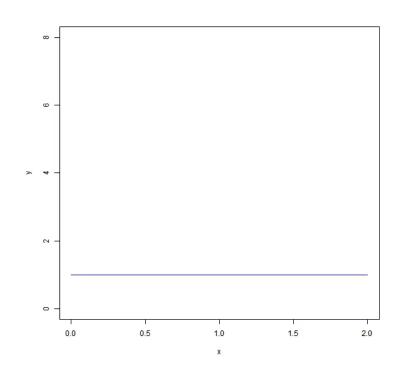
- saveGIF(code, movie.name, interval, ani.width, ani.height,...)
 - Code: the loop producing animation
 - Interval: time interval between frames
 - Ani.width, ani.height animation size
- saveVideo(code, video.name,...)
- saveLatex(...)
 - PDF animation

Dynamic from static

Example: Visualizing $y=x^a$, a=0...3

```
#specifying where ffmpeg is located
ani.options(ffmpeg="D:\\ffmpeg\\bin\\ffmpeg
.exe")
```

```
saveVideo({
    x<-seq(0,2,0.01)
    for (a in seq(0,3,by=0.01)) {
        y<-x^a
        plot(x,y, type="I", col="blue", ylim=c(0,8))
    }
},video.name="Z:\\732A39_732A98\\2016\\Le
cture 5\\Powers.mp4", interval=0.05,
    ani.width=600,ani.height=600
)</pre>
```



Reading home

- http://www.gapminder.org
- Chapter 8, p.302-326
- Heer, Jeffrey, and George Robertson. "Animated transitions in statistical data graphics." *IEEE* transactions on visualization and computer graphics 13.6 (2007): 1240-1247.