# cranvas: Interactive Statistical Graphics in R via Qt

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Introduction to cranvas

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# Appetizer

- the name (CRAN + canvas  $\Rightarrow$  cranvas)
- demo
  - old problems (with new look) in cranvas
  - NRC rankings of statistics departments

## Looking back

• standalone software packages: GGobi [Swayne et al., 2003], MANET [Unwin et al., 1996] and Mondrian [Theus, 2002], etc



- R itself has poor support for interactivity (ancient *locate()*? *getGraphicsEvent()*? no kidding! there is a **fun** package [Xie et al., 2011], and also **sudoku**)
- R packages: **iplots** [Urbanek and Wichtrey, 2011], **rggobi**, **playwith**, **rgl** (as I showed just now)...

# **Overview of cranvas**

- next generation of GGobi (and more)
  - interactive statistical graphics in R
  - focus on speed, flexibility and elegance
- infrastructures (details later):
  - painting (drawing): **qtbase** & **qtpaint** (connect R with Qt)
  - data structure: plumbr & objectSignals (mutaframes, reference classes, signals)

- aesthetics: **scales** (and other **ggplot2**'s offspring)

- all available at http://github.com/ggobi (to be released on Bioconductor & CRAN)
  - refer to the wiki for installation under Linux & Mac
  - Windows version not available at the moment
- usage
  - create a data object: mf <- qdata(df)</pre>
  - make a series of plots based on it: qhist(x1, data = mf);
    qscatter(x2, x3, data = mf)
- cranvas is still under active development

- milestone 1.0: get basic plots working (almost done)
- milestone 2.0 scheduled in November (after R 2.14.0): new types of plots, speed, documentation

#### Data structure: augmented data

- > library(cranvas)
- > data(flea, package = "tourr")
- > head(flea) # first 6 rows of flea

	tars1	tars2	head	aede1	aede2	aede3	species
1	191	131	53	150	15	104	Concinna
2	185	134	50	147	13	105	Concinna
3	200	137	52	144	14	102	Concinna
4	173	127	50	144	16	97	Concinna

5 171 118 49 153 13 106 Concinna 6 160 118 47 140 15 99 Concinna > str(flea)

'data.frame': 74 obs. of 7 variables: 185 200 173 171 160 188 186 174 163 \$ tars1 : int 191 134 137 127 118 118 134 129 131 115 \$ tars2 : int 131 53 50 52 50 49 47 54 51 52 47 ... \$ head : int : int 150 147 144 144 153 140 151 143 144 142 \$ aede1 : int 15 13 14 16 13 15 14 14 14 15 ... \$ aede2 104 105 102 97 106 99 98 110 116 95 \$ aede3 : int species: Factor w/ 3 levels "Concinna ", "Heikert. ",... \$

- > qflea <- qdata(flea, color = species)</pre>
- > head(qflea) # what is the difference?

	tars1	tars2	head	aede1	aede2	aede3	species	.brushed
1	191	131	53	150	15	104	Concinna	FALSE
2	185	134	50	147	13	105	Concinna	FALSE
3	200	137	52	144	14	102	Concinna	FALSE
4	173	127	50	144	16	97	Concinna	FALSE
5	171	118	49	153	13	106	Concinna	FALSE
6	160	118	47	140	15	99	Concinna	FALSE
	.visible .color		.borde	r .siz	e			
1	TRUE #F8766D		#F8766	D	4			
2	TF	RUE #F8	3766D	#F8766	D	4		

- 3
   TRUE
   #F8766D
   #F8766D
   4

   4
   TRUE
   #F8766D
   #F8766D
   4

   5
   TRUE
   #F8766D
   #F8766D
   4

   6
   TRUE
   #F8766D
   #F8766D
   4
- the augmented data contains information about each observation (row)
  - .brushed: logical vector whether a row is brushed or not
  - .visible: similar visible or not?
  - color, .border, .size aesthetics of graphical elements

#### Data structure: mutaframe

- what we created by *qdata()* was a mutaframe (using **plumbr**)
- a mutaframe is like a data frame, but is mutable (technically, it is an environment)
- > str(qflea, max.level = 1)

Classes 'mutaframe', 'environment' <environment: 0x589cb2
- attr(\*, "col.names")= chr [1:12] "tars1" "tars2" "head
- attr(\*, "row.names")= chr [1:74] "1" "2" "3" "4" ...</pre>

- attr(\*, "changed")=Formal class 'Signal' [package "obj - attr(\*, "Brush")=Formal class 'BRUSH' [package "cranva - attr(\*, "Link")=Classes 'mutalist', 'environment' <env ..- attr(\*, "envNames")= chr "linkid" ..- attr(\*, "userNames")= chr "linkid"
- attr(\*, "Scales")=Formal class 'Scales\_meta' [package
- attr(\*, "Generator") = chr "d38bbe46dae5fa45758f3609f5c
- modification to a mutaframe is always "global", which enables us to pass and modify a mutaframe anywhere in functions
  - application to tour (projections always changing, and the plot needs to know this)

- > a <- 10 # ordinary R object</pre>
- > b <- a
- > b <- 5
- > a # certainly a is not modified

#### [1] 10

- > ## but mutaframes are different
- > x <- qflea # assign qflea to x</pre>
- > qflea\$tars1[1] # first element of tars1

#### [1] 191

> x\$tars1[1] # same element in x

#### [1] 191

- > x\$tars1[1] <- 300 # modify it
- > x\$tars1[1]

#### [1] 300

> qflea\$tars1[1] # original copy also modified!

#### [1] 300

#### Data structure: listeners

- how does interaction work?
  - interaction does not modify the plot directly
  - instead, it modifies data and triggers listeners on data, which do the job of updating the plot
  - it makes developers' life a lot easier
- what is a listener?
  - it is a function attached to a mutaframe

- (roughly speaking) whenever the mutaframe is modified, the function is executed
- which part of the mutaframe is modified?
  - we should not abuse the listeners, hence comes the concept of the "data pipeline"
  - .brushed changed? update the brush layer!
  - .visible changed? update the main plot layer!
  - original data changed? go back to the beginning and recalculate everything, then update all the layers!
- each time we create a plot, a listener is attached to the mutaframe

 so the .brushed column can update several plots at the same time

#### Data structure: a summary

- a data object is sitting behind the scene
- interactions change the data
- listeners on the data update the plot(s)

#### Interaction: brushing or selection

- mouse interactions
  - left click and move the brush; right click to resize the brush; middle button to toggle between brush and selection
  - brush mode and identify mode (use ? to toggle)
- selection mode
  - AND, OR, XOR, NOT, COMPLEMENT (use the initial letters)
- related keyboard interactions

- Delete makes elements invisible; F5: all visible

# Interaction: linking

- one-to-one linking
  - brush one observation in this plot, highlight the same observation in other plots
- categorical linking
  - ..., all observations in the same category are brushed too
- kNN linking

- ..., k nearest neighbors are brushed

# Interaction: misc

- + and changes the alpha transparency
- PageDown and PageUp navigates through brush history

• ...

# Examples: plots in cranvas now, and what's new

- histogram (and spine plot): change binwidth with  $\uparrow$  and  $\downarrow$ ; shift bins with  $\leftarrow$  and  $\rightarrow$
- density plot: similar to histogram
- bar plot: application to missing value plot
- scatter plot: change size of points with  $\uparrow$  and  $\downarrow$

- boxplot: also show small boxplots for brushed observations inside original boxplots (not actually new)
- mosaic plot: dynamically change variables in the plot and layout
- parallel coordinates plot: rearrange order of variables (not new); show data ranges
- map: also cartograms
- time series plot: many, many new features (as a result of GSoC project)

#### **Examples: flea and states demographics**

- flea beetles
  - tour
  - categorical linking by species
- US states demographics
  - selection mode
  - kNN linking

# Limitations

- suffers from Qt flaws and bugs
  - can be slow (mainly in scatter plots) due to changes in BSP tree algorithm and cache failure; I was told it WAS extremely fast
  - mysterious clipping (aggressive: points clipped into halves; imprecise: no clipping even when points reach beyond boundary)
- Qt is big (in size) and installation of **qtbase** / **qtpaint** under Windows may be difficult

#### **Future plans**

- automatic and *interactive* legend
- conditioning (faceting) like Trellis
- more types of plots (e.g. hexagons)
- a lot to learn from **ggplot2** and **iplots** (ix?)

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#### Looking at the acknowledgement list



## **Questions & Comments?**

- Testing and bug reports (https://github.com/ggobi/ cranvas/issues) are welcome
- Thanks!

# References

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September 23, 2011

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