R for cytometry - plotting

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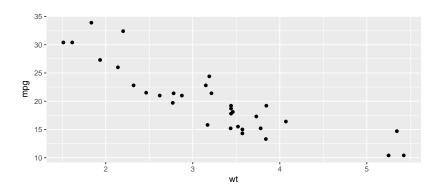
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plotting with R

- ▶ Plotting commands from different algorithms
- ▶ ggplot

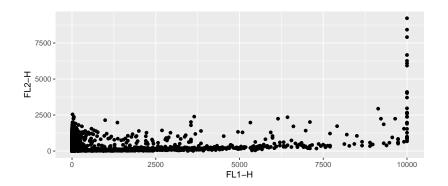
ggplot simple example

```
require(ggplot2)
ggplot(mtcars, aes(x = wt, y = mpg)) +
  geom_point()
```



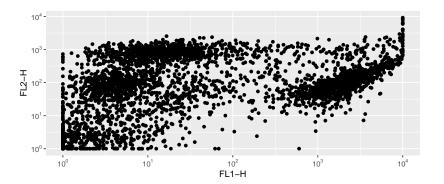
ggplot with flow data

```
require(flowCore)
data(GvHD)
frame <- GvHD[[1]]
df <- as.data.frame(frame@exprs)
ggplot(df, aes(x = `FL1-H`, y = `FL2-H`)) +
    geom_point()</pre>
```

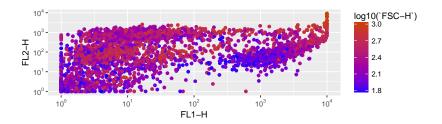


log scales

```
require(scales)
ggplot(df, aes(x = `FL1-H`, y = `FL2-H`)) +
  geom_point() +
  scale_x_log10(breaks = 10^(0:5),
   labels = trans_format("log10", math_format(10^.x))) +
  scale_y_log10(breaks = 10^(0:5),
   labels = trans_format("log10", math_format(10^.x)))
```



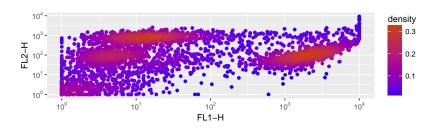
colors



density

```
require(MASS)
get_density <- function(x, y, n = 100){
  dens <- kde2d(x = x, y = y, n = n)
  ix <- findInterval(x, dens$x)
  iy <- findInterval(y, dens$y)
  dens$z[cbind(ix, iy)]
}</pre>
```

density colors

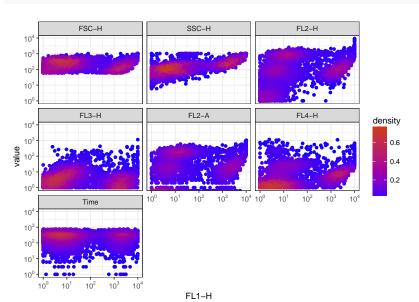


multiple plots at same time

```
require(reshape2)
require(dplyr)
df melt <- melt(df, id.vars = "FL1-H")</pre>
df melt <- mutate(group by(df melt, variable),</pre>
  density = get density(log10(`FL1-H` + 1),
                         log10(value + 1)))
p \leftarrow ggplot(df melt, aes(x = FL1-H), y = value,
                          color = density))+
  geom point() +
  scale x log10(breaks = 10^{\circ}(0:5),
    labels = trans_format("log10", math_format(10^.x))) +
  scale_y_log10(breaks = 10^(0:5),
    labels = trans_format("log10", math_format(10^.x))) +
  scale_color_gradient(name = "density",
    low = "#0000FF", high = "#CD4000") +
  facet_wrap(~variable) +
  theme bw()
```

multiple plots at same time

print(p)



further reading

Have a look at the following:

```
?geom_line
?geom_abline
?geom_smooth
?geom_linerange
?geom_boxplot
?geom_histogram
?geom_text
```

Also this tutorial.