TikZ annotations for ggplots with the ggtikz package

August 17, 2021

Contents

1 Prerequisites

1.1 LaTeX side

As the name implies, `ggtikz` requires `tikz`, which must be loaded in the document's preamble. Furthermore, the `calc` tikz library is required.

Thus, the preamble must contain:

```latex
\usepackage{tikz}
\usetikzlibrary{calc}
```

1.2   R side

The `tikzDevice` package is required to render plots and `ggtikz` annotations to the tikz format. We also have to make some base plots, using `ggplot2`.

Here, we set the graphics device to `tikz` – `ggtikz` does not work with any other graphics device!

```r
library(knitr)
library(ggplot2)
library(ggtikz)
opts_chunk$set(
    dev = "tikz",
    external = TRUE,
    fig.path = "example-vignette-figures/",
    fig.width = 3,
    fig.height = 3,
    fig.align = "center"
)
```

2   Basic usage with ggtikz()

For simple one-step annotations, the `ggtikz` helper function is available.

It accepts a ggplot object as its first argument. Further arguments are passed on to `ggtikzAnnotation` (see section 3).

```r
p <- ggplot(mtcars, aes(disp, mpg)) + geom_point()
ggtikz(p, "\\fill[red] (0.5,0.5) circle (2mm);", xy="plot")
```
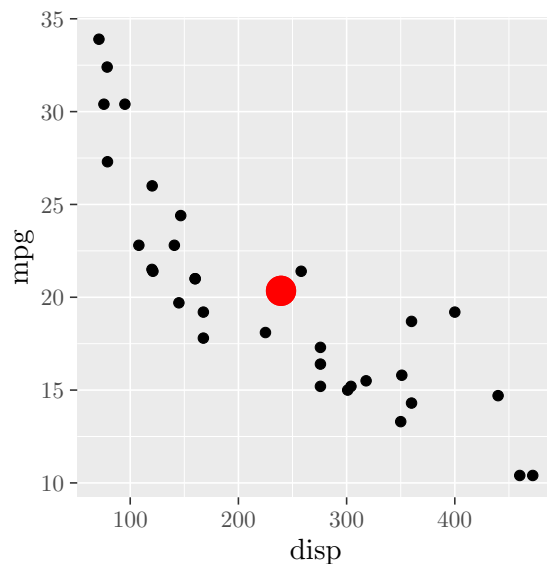
3 Advanced usage with canvases and annotations

With `ggtikz()`, only a single annotation can be added to a plot. If multiple annotations are needed, then we first need to create a `ggtikzCanvas()`, to which one or more `ggtikzAnnotation()` can be added.

3.1 Single-panel plots

Let's create a single-panel plot for annotation.

```
p <- ggplot(mtcars, aes(disp, mpg)) + geom_point()
```

We can then set up an annotation canvas and add tikz annotations. Note that first, we print the base plot to the device [1], and then the annotation canvas. The annotation canvas does not take care of drawing the annotated plot (the `ggtikz()` helper does handle this with the `draw = TRUE` parameter).

3.1.1 Annotation relative to the whole plot

```
canvas <- ggtikzCanvas(p)
annotation <- ggtikzAnnotation(
```
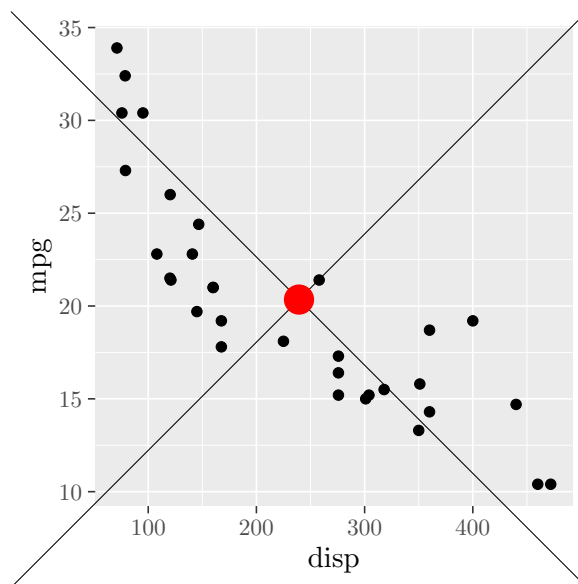
---

[1]no explicit calls to `tikz()` and `dev.off()` are needed, because knitr opens and closes the device automatically.

```
    "
    \\draw (0,0) -- (1,1);
    \\draw (0,1) -- (1,0);
    \\fill[red] (0.5,0.5) circle (2mm);
    ",
    xy = "plot"
)
p                    # first draw the plot
canvas + annotation  # then draw the annotations
```
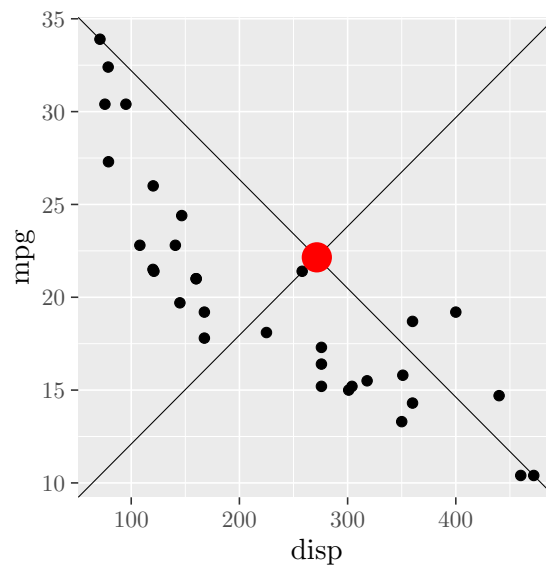

3.1.2 Annotation relative to the panel

```
canvas <- ggtikzCanvas(p)
annotation <- ggtikzAnnotation(
    "
    \\draw (0,0) -- (1,1);
    \\draw (0,1) -- (1,0);
    \\fill[red] (0.5,0.5) circle (2mm);
    ",
    xy = "panel",
    panelx = 1, panely = 1
)
p
canvas + annotation
```

3.1.3 Annotation relative to data coordinates

In addition to unitless tikz coordinates, you can also use absolute lengths, such as the 1 cm in the example below.

```
canvas <- ggtikzCanvas(p)
annotation <- ggtikzAnnotation(
    "
    \\draw[thick,red] (100,20) -| (400,15);
    \\draw[<-] (153,24) -- ++(30:1cm) node[at end, anchor=south] {Interesting!};
    ",
    xy = "data",
    panelx = 1, panely = 1
)
p
canvas + annotation
```
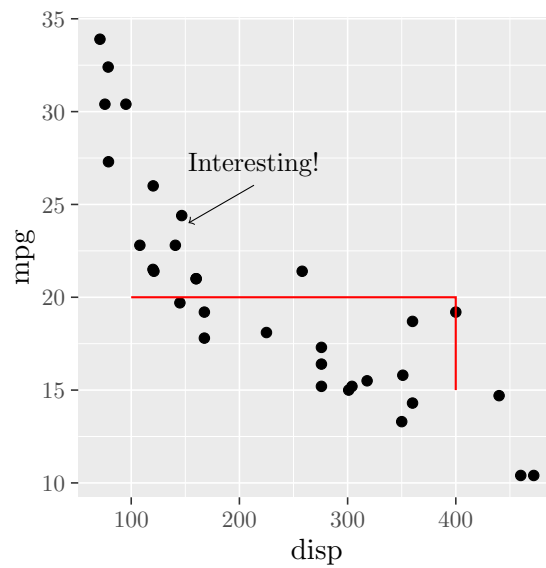
3.1.4 Mixing panel and data references

The reference frames for x and y coordinates can be separately assigned as `data` or `panel`. However, note that the `plot` reference frame must be given for both x and y directions (with the `xy` argument), and cannot be mixed!

```
canvas <- ggtikzCanvas(p)
annotation <- ggtikzAnnotation(
   "\\fill[red] (0.5,30) circle (2mm);",
   x = "panel", y = "data",
   panelx = 1, panely = 1
)
p
canvas + annotation
```
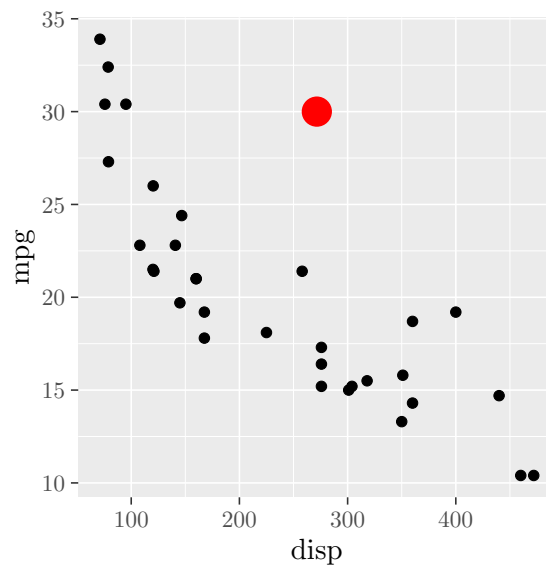
3.1.5 Turning off clipping

It is possible to turn off clipping for annotations, in order to draw outside of the plot area.
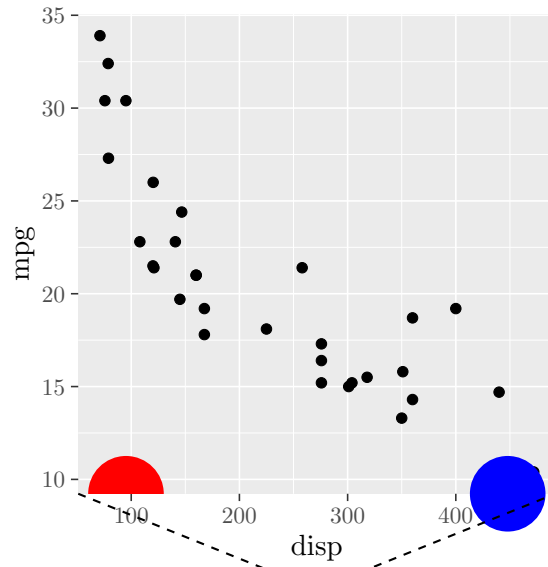
```r
canvas <- ggtikzCanvas(p)
annotation_clip <- ggtikzAnnotation(
    "\\fill[red] (0.1,0) circle (5mm);",
    xy = "panel",
    panelx = 1, panely = 1
)


annotation_unclip <- ggtikzAnnotation(
    "\\fill[blue] (0.9,0) circle (5mm);",
    xy = "panel",
    panelx = 1, panely = 1,
    clip = "off"
)


annotation_unclip2 <- ggtikzAnnotation(
    "\\draw[thick, dashed] (0,0) -- (0.5,-0.2) -- (1,0);",
    xy = "panel",
    panelx = 1, panely = 1,
    clip = "off"
)
p
canvas + annotation_clip + annotation_unclip + annotation_unclip2
```
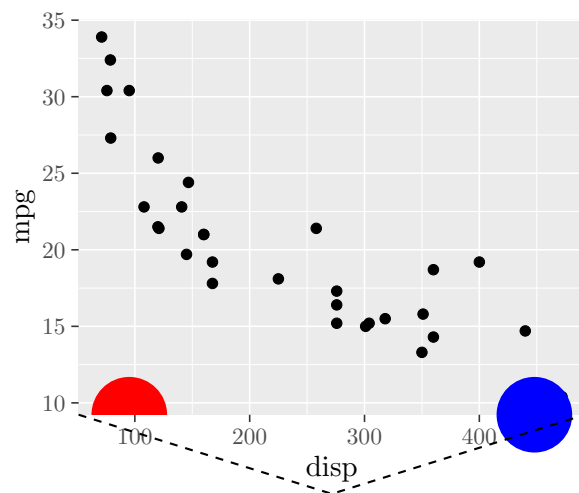
However, note that the surrounding plot area is not automatically unclipped to accommodate for the annotations. This can be alleviated manually by increasing the plot margins.

```
p + theme(plot.margin = margin(t=0.5, b = 1, unit = "cm"))
canvas + annotation_clip + annotation_unclip + annotation_unclip2
```

3.2 Multi-panel plots: wrap

```
p_wrap <- p + facet_wrap(~cyl, scales="free", ncol=2)
```

3.2.1 Annotations in separate panels, relative to data or panel coordinates
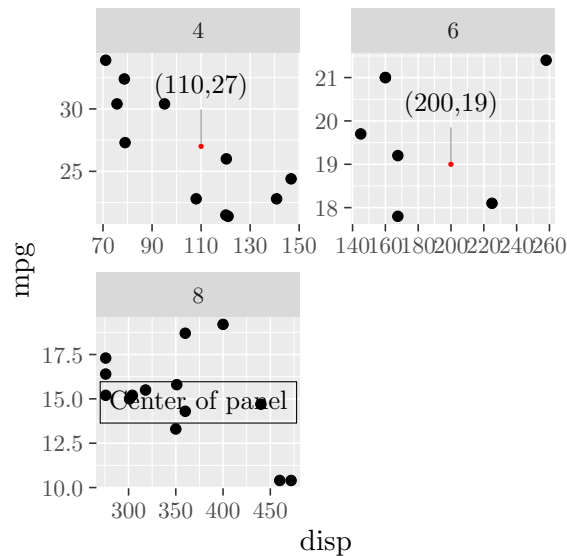
```
canvas <- ggtikzCanvas(p_wrap)

# Relative to data coordinates
annotation1 <- ggtikzAnnotation(
    "
    \\node[pin={90:(110,27)}, circle, fill=red,
        inner sep=0, outer sep=0, minimum size=2pt]
        at (110,27)
        {};
    ",
    xy = "data",
    panelx = 1, panely = 1
)

# Relative to data coordinates
annotation2 <- ggtikzAnnotation(
    "
    \\node[pin={90:(200,19)}, circle, fill=red,
    inner sep=0, outer sep=0, minimum size=2pt]
    at (200,19)
    {};
    ",
    xy = "data",
    panelx = 2, panely = 1
)

# Relative to panel coordinates
annotation3 <- ggtikzAnnotation(
    "
    \\node[draw, anchor=center] at (0.5, 0.5)
        {Center of panel};
    ",
    xy = "panel",
    panelx = 1, panely=2
)
```

```
p_wrap
canvas + annotation1 + annotation2 + annotation3
```


3.2.2 Annotations in separate panels, relative to data coordinates

```
canvas <- ggtikzCanvas(p_wrap)
annotation1 <- ggtikzAnnotation(
    "
    \\node[pin={90:(110,27)}, circle, fill=red,
        inner sep=0, outer sep=0, minimum size=2pt]
        at (110,27)
        {};
    ",
    xy = "data",
    panelx = 1, panely = 1
)
annotation2 <- ggtikzAnnotation(
    "
    \\node[pin={90:(200,19)}, circle, fill=red,
    inner sep=0, outer sep=0, minimum size=2pt]
    at (200,19)
    {};
    ",
    xy = "data",
    panelx = 2, panely = 1
)
```

```
p_wrap
canvas + annotation1
```
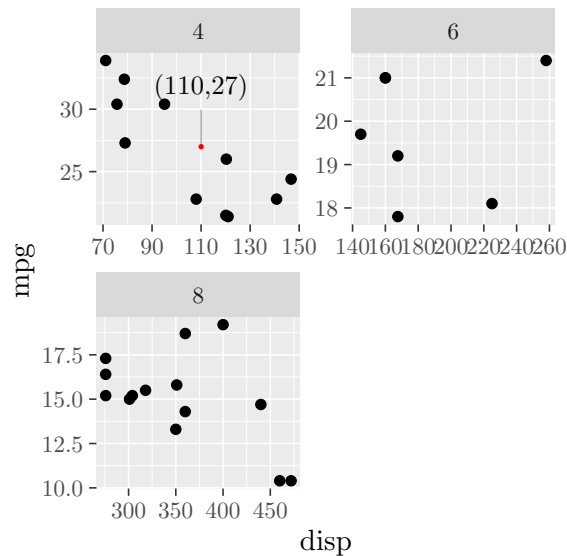

3.3 Multi-panel plots: grid

Annotations can also be made on individual panels of plots faceted with `facet_grid`.

```
p_grid <- p + facet_grid(gear~cyl, scales="free", as.table=FALSE)
```

```
canvas <- ggtikzCanvas(p_grid)
annot_grid1 <- ggtikzAnnotation(
    "\\node[fill=white, draw, text width=2cm] at (0.5,0.5)
        {panelx=1, panely=1};",
    xy = "panel",
    panelx = 1, panely = 1
)
annot_grid2 <- ggtikzAnnotation(
    "\\node[fill=white, draw, text width=2cm] at (0.5,0.5)
        {panelx=2, panely=3};",
    xy = "panel",
    panelx = 2, panely = 3
)
annot_grid3 <- ggtikzAnnotation(
    "
    \\draw[<-, blue] (90,15) -- ++(30:5mm)
        node [at end, anchor=south west] {(90,15)};
```

```
    ",
    xy = "data",
    panelx = 1, panely = 3
)
p_grid
canvas + annot_grid1 + annot_grid2 + annot_grid3
```
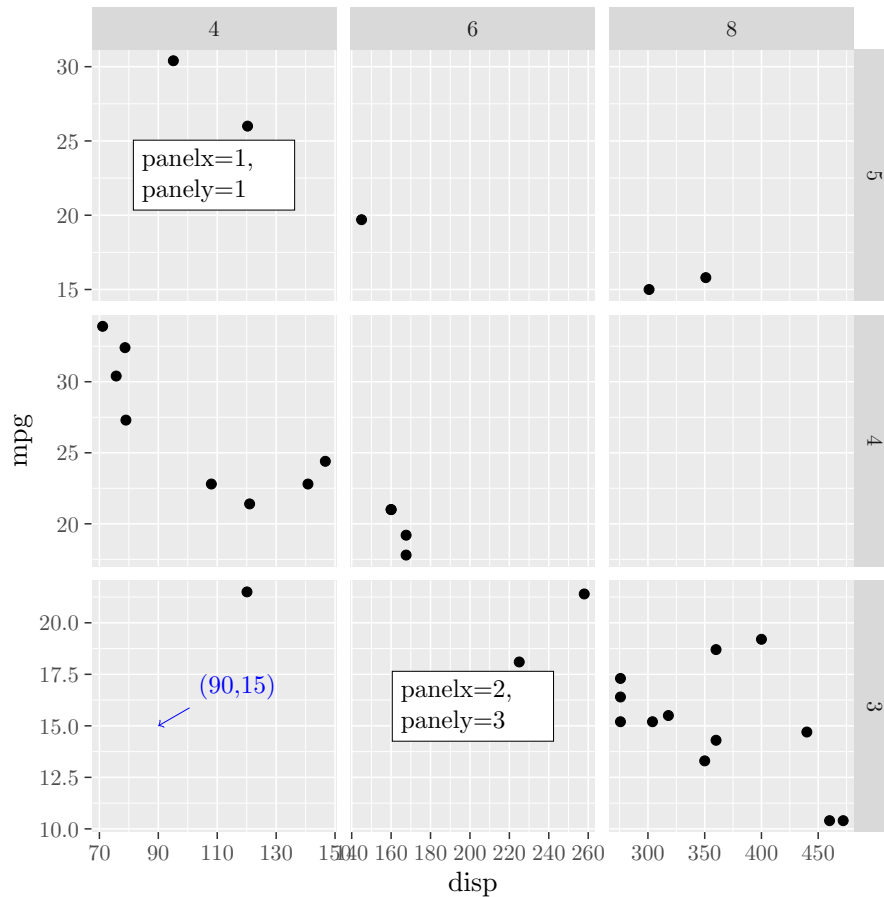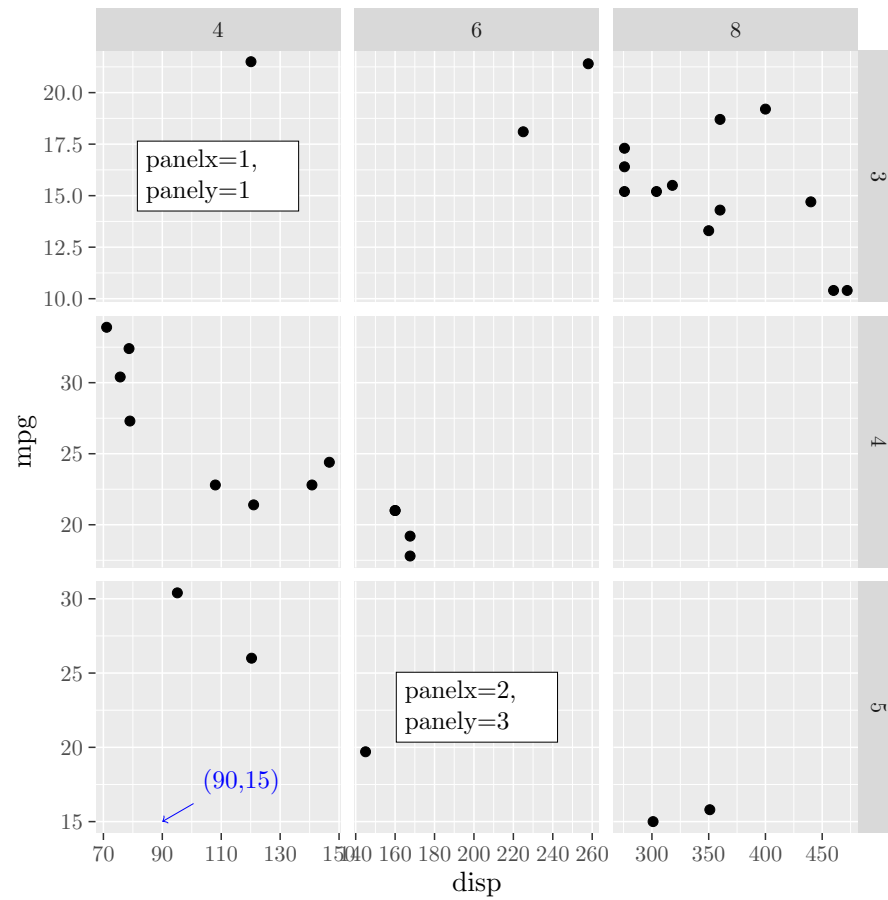

3.4   Re-using annotations

Annotations can be re-used between plots and ggtikz canvases. However, be aware that panel position specifications rely on the *visual position* of the panels, and *not on the value of the facet variables.*

```
p_grid2 <- p + facet_grid(gear~cyl, scales="free", as.table=TRUE)
canvas2 <- ggtikzCanvas(p_grid2)
```
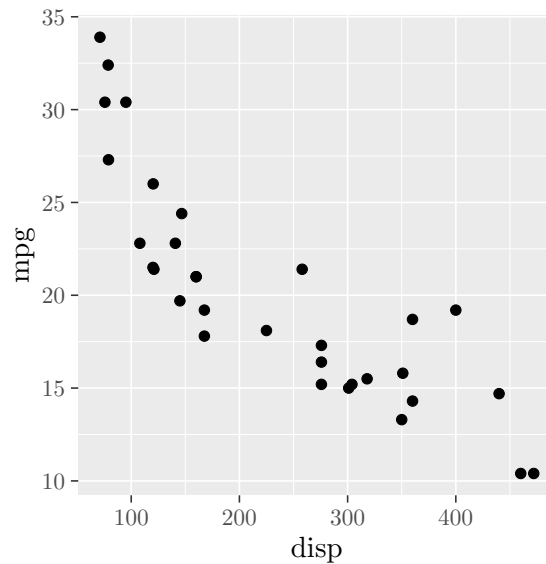
```
p_grid2
canvas2 + annot_grid1 + annot_grid2 + annot_grid3
```


It is also not possible to add annotations to a plot for which the requested panels are not available.

```
p
```

```
canvas <- ggtikzCanvas(p)
canvas + annot_grid2
```

```
## Error in get_annotation_valid.ggtikzCanvas(self, ggtikzAnnotation):
Annotation wants to be placed in panelx = 2, but the plot only has
1.
```

4   Transformed scales

TikZ coordinates can automatically be transformed to accommodate trans-
formed scales, such as log-transformed ones. This is activated by setting `transform = TRUE`
in the call to `ggtikzAnnotation`. With `transform = FALSE`, annotations made
with data coordinates are out of place in log plots, but the transformation cal-
culation can be done manually.

```
p_log <- ggplot(mtcars, aes(mpg, disp)) +
    geom_point() +
    scale_x_continuous(trans="log10")
```

```
canvas_log <- ggtikzCanvas(p_log)
# Untransformed coordinates: wrong position
annot_log <- ggtikzAnnotation(
    "\\fill[red] (1,100) circle (2mm);
    \\node[anchor=west, text=red] at (1, 100)
```

```
        {The circle is not at (1,100)!};
    ", xy = "data", transform = FALSE, panelx = 1, panely = 1
)

# Transformed coordinates: correct position
# The literal coordinate in the node text was wrapped in an \mbox
# LaTeX command to prevent automatic transformation -- it can't
# distinguish between coordinates which are supposed to be text,
# and actual coorinates.
annot_log2 <- ggtikzAnnotation(
    "\\fill[blue] (20,200) circle (2mm);
    \\node[anchor=south, text=blue] at (20, 200)
        {This circle is at (\\mbox{20,200})!};
    ", xy = "data", transform = TRUE, panelx = 1, panely = 1
)

# Untransformed coordinates, calculated manually by hand:
# correct position
annot_log3 <- ggtikzAnnotation(
    "\\fill[magenta] (1.477,400) circle (2mm);
    \\node[anchor=east, text=magenta] at (1.477, 400)
        {This circle is at (30,400)!};
    ", xy = "data", transform = FALSE, panelx = 1, panely = 1
)
p_log
canvas_log + annot_log + annot_log2 + annot_log3
```
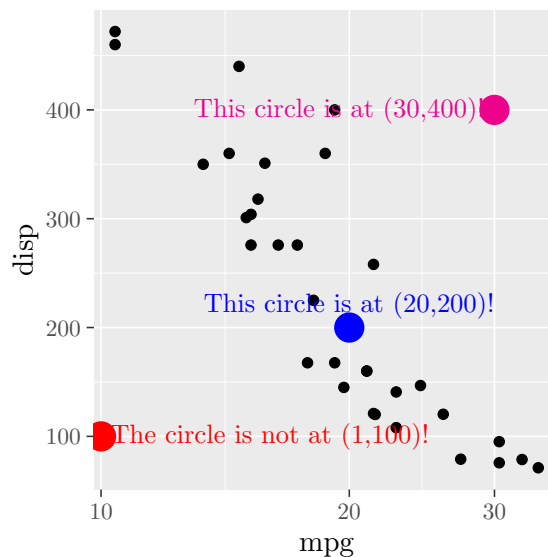
5   Using styles defined in the surrounding document

Annotations can access styles which are defined in the containing document before the relevant .tikz file is included, allowing you to re-use global styles. Note that by default, knitr sets the option `external` to `TRUE`. Therefore, tikz graphics are pre-compiled to pdf. In that case, the tikzDevice needs to know about these styles, or an error will occur during externalization.

```
\tikzset{loud/.style={
    draw=yellow,
    fill=red,
    text=blue
}}
```

```
p
```
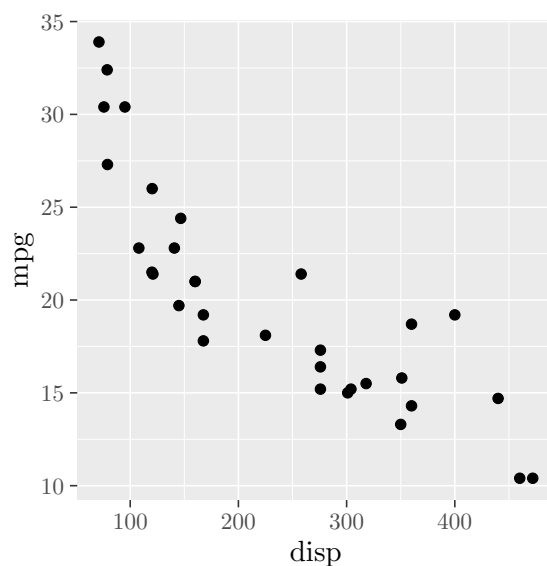


```
canvas <- ggtikzCanvas(p)
styled_annot <- ggtikzAnnotation(
    "\\node[loud] at (0.5,0.5) {Look at me!};",
    xy = "plot"
)
p
canvas + styled_annot
```

Look at me!