

Class 5: Data Viz with ggplot

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Today we are exploring the **gggplot** package and how to make nice figures in R.

There are lots of ways to make figures and plot in R. These include:

- so called “base” R
- and add on packages like **ggplot2**

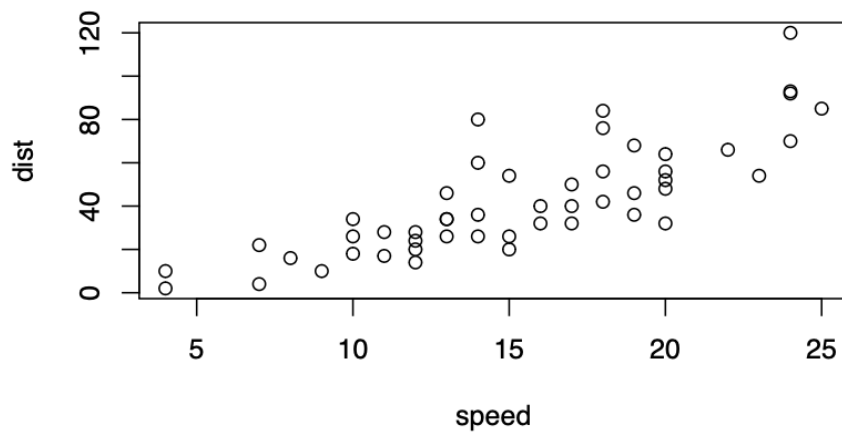
Here is a simple “base” R plot.

```
head(cars)
```

	speed	dist
1	4	2
2	4	10
3	7	4
4	7	22
5	8	16
6	9	10

We can simply pass this to the `plot()` function.

```
plot(cars)
```



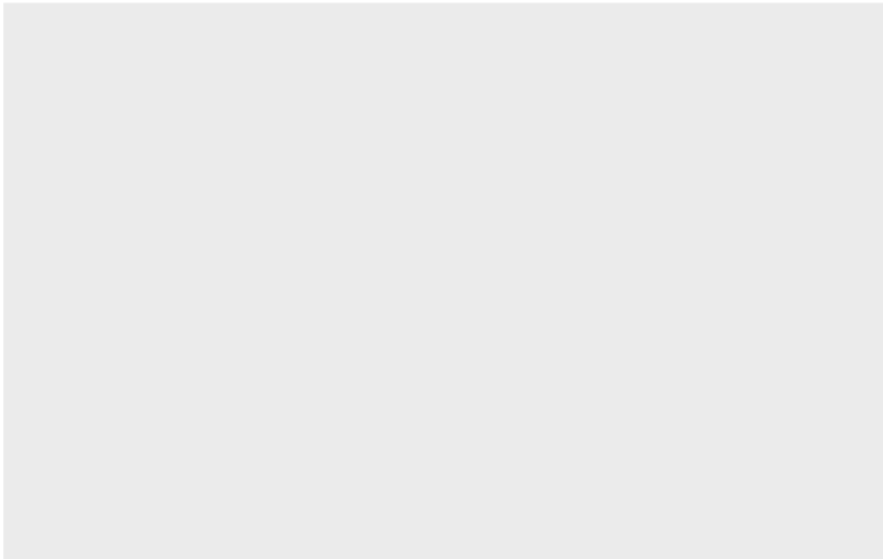
Key-point: Base R is quick but not so nice looking in some folks eyes.

Let's see how we can plot this with **ggplot2**...

1st I need to install this add-on package. For this we use the `install.packages()` function - **WE DO THIS IN THE CONSOLE, NOT our report**. This is a one time only deal.

2nd We need to load the package with `library()` function every time we want to use it.

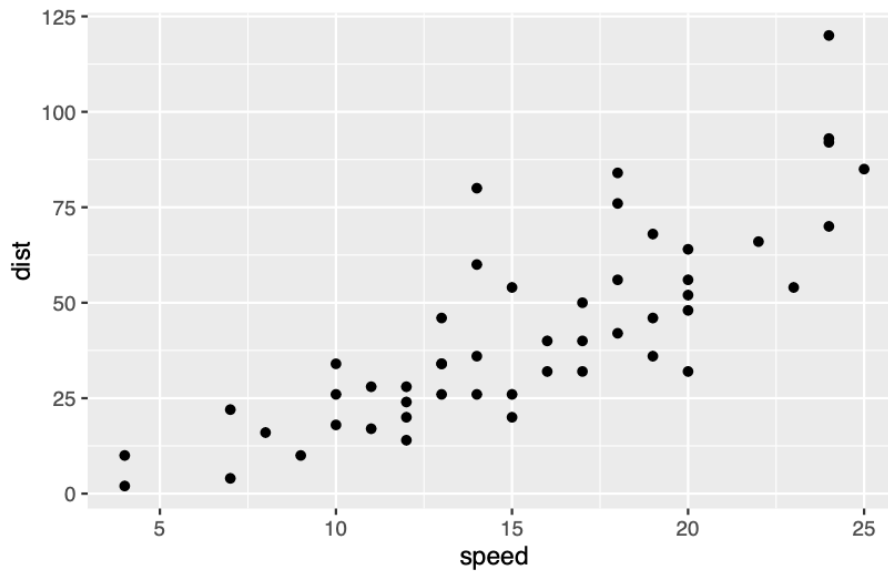
```
library(ggplot2)
ggplot(cars)
```



Every ggplot is composed of at least 3 layers:

- **data** (i.e a data.frame with the things you want to plot),
- aesthetics **aes()** that map the columns of data to your plot features (i.e. aesthetics)
- geoms like **geom_point()** that sort how the plot appears

```
ggplot(cars) +  
  aes(x=speed, y=dist) +  
  geom_point()
```



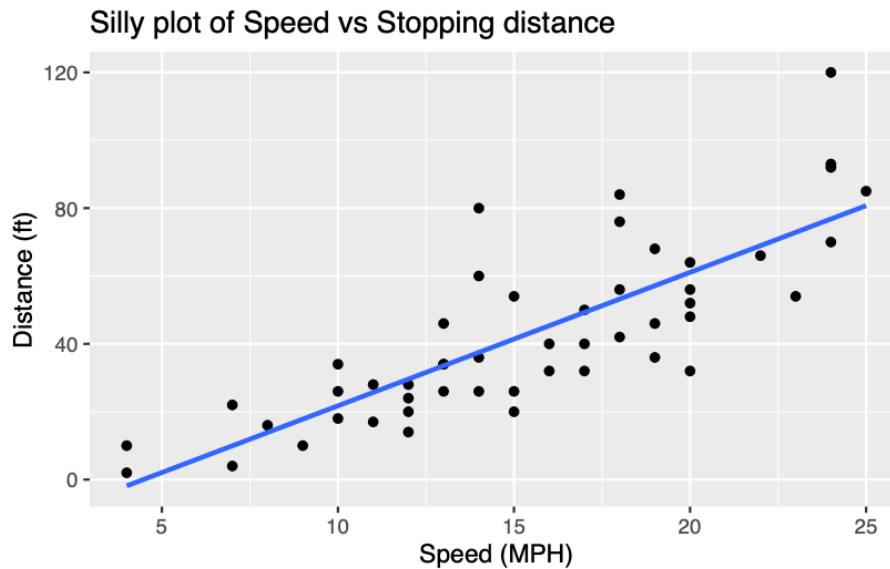
Key point: For simple “canned” graphs base R is quicker but as things get more custom and elaborate then ggplot wins out...

Let’s add more layers to our ggplot

Add a line showing the relationship between x and y Add a title Add custom axis labels “Speed (MPH)” and “Distance (ft)” Chnage the theme...

```
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point() +
  geom_smooth(method="lm", se=F) +
  labs(title="Silly plot of Speed vs Stopping distance") +
  labs(x="Speed (MPH)", y="Distance (ft)")
```

`geom_smooth()` using formula = 'y ~ x'



```
theme_bw()
```

```
<theme> List of 144
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  ..@ linewidth   : num 0.5
  ..@ linetype    : num 1
  ..@ lineend     : chr "butt"
  ..@ linejoin    : chr "round"
  ..@ arrow       : logi FALSE
  ..@ arrow.fill  : chr "black"
  ..@ inherit.blank: logi TRUE
 $ rect      : <ggplot2::element_rect>
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  ..@ colour      : chr "black"
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  ..@ linetype    : num 1
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  ..@ inherit.blank: logi TRUE
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  ..@ family      : chr ""
  ..@ face        : chr "plain"
```

```

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$ point                                     : <ggplot2::element_point>
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..@ size        : num 1.5
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$ geom                                     : <ggplot2::element_geom>
..@ ink         : chr "black"

```

```

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..@ borderwidth: num 0.5
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..@ bordertype : int 1
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..@ pointsize  : num 1.5
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..@ fill       : NULL
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$ margins      : <ggplot2::margin> num [1:4] 5.5 5.5 5.5 5.5
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..@ angle      : NULL

```

```

..@ lineheight : NULL
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..@ fontwidth         : num NA
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```



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..@ margin    : <ggplot2::margin> num [1:4] 2.2 0 0 0
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..@ vjust     : num 0
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$ axis.text.x.bottom : NULL
$ axis.text.y : <ggplot2::element_text>
..@ family    : NULL
..@ face      : NULL
..@ italic    : chr NA

```

```

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..@ fontweight : num NA
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..@ vjust : NULL
..@ angle : NULL
..@ lineheight : NULL
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..@ size : NULL
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..@ vjust : NULL
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..@ lineheight : NULL
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..@ debug : NULL
..@ inherit.blank: logi TRUE

```

```

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$ axis.ticks.x.top                           : NULL
$ axis.ticks.x.bottom                         : NULL
$ axis.ticks.y                               : NULL
$ axis.ticks.y.left                          : NULL
$ axis.ticks.y.right                         : NULL
$ axis.ticks.theta                           : NULL
$ axis.ticks.r                               : NULL
$ axis.minor.ticks.x.top                     : NULL
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$ axis.minor.ticks.length.y                  : NULL
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$ axis.minor.ticks.length.y.right            : NULL
$ axis.minor.ticks.length.theta              : NULL
$ axis.minor.ticks.length.r                  : NULL
$ axis.line                                  : <ggplot2::element_blank>
$ axis.line.x                               : NULL

```

```

$ axis.line.x.top           : NULL
$ axis.line.x.bottom       : NULL
$ axis.line.y              : NULL
$ axis.line.y.left         : NULL
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$ legend.key.spacing.y     : NULL
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  ..@ lineheight            : NULL

```

```

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$ legend.justification.left : NULL
$ legend.justification.right : NULL
$ legend.justification.inside : NULL
[list output truncated]
@ complete: logi TRUE
@ validate: logi TRUE

```

Going further

Read some expression data

```

url <- "https://bioboot.github.io/bimm143_S20/class-material/up_down_expression.txt"
genes <- read.delim(url)
head(genes)

```

	Gene	Condition1	Condition2	State
1	A4GNT	-3.6808610	-3.4401355	unchanging
2	AAAS	4.5479580	4.3864126	unchanging
3	AASDH	3.7190695	3.4787276	unchanging
4	AATF	5.0784720	5.0151916	unchanging
5	AATK	0.4711421	0.5598642	unchanging
6	AB015752.4	-3.6808610	-3.5921390	unchanging

Q1. How many genes are in this wee dataset?

```
nrow(genes)
```

```
[1] 5196
```

Q2. How many “up” regulated genes are there?

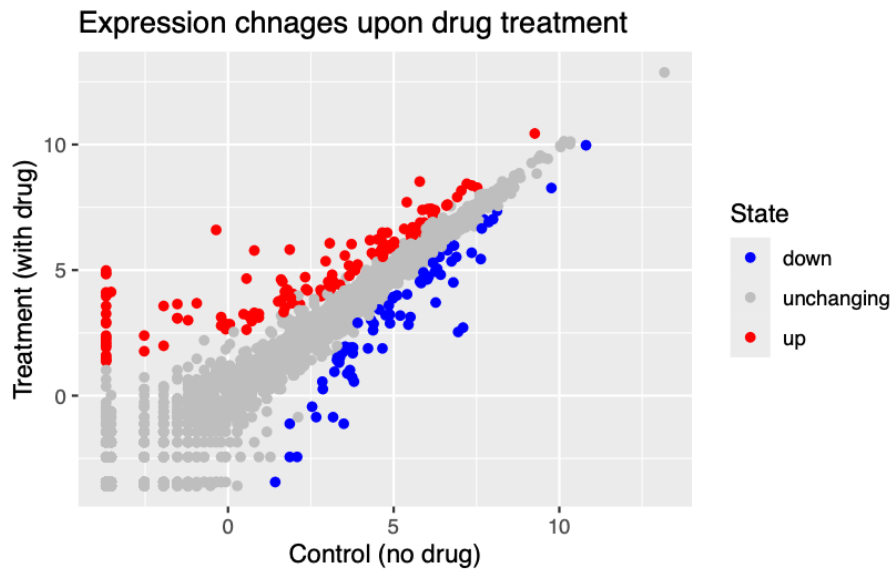
```
table(genes$State)
```

down	unchanging	up
72	4997	127

A useful function for counting up occurrences of things in a vector is the `table()` function.

Make a v1 figure

```
p<-ggplot(genes) +
  aes(x= Condition1,
      y= Condition2,
      col=State) +
  geom_point() +
  labs(title= "Expression chnages upon drug treatment",
       x="Control (no drug)",
       y="Treatment (with drug)")
p + scale_colour_manual(values = c("blue", "gray","red"))
```



```
##More PLOtting library(gapminder)
```

```
# File location online
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.ts"

gapminder <- read.delim(url)
```

Lets have a wee peak

```
head(gapminder, 3)
```

	country	continent	year	lifeExp	pop	gdpPercap
1	Afghanistan	Asia	1952	28.801	8425333	779.4453
2	Afghanistan	Asia	1957	30.332	9240934	820.8530
3	Afghanistan	Asia	1962	31.997	10267083	853.1007

Q4. How many different country values are in this dataset?

```
nrow(gapminder)
```

```
[1] 1704
```

```
length(table(gapminder$country))
```

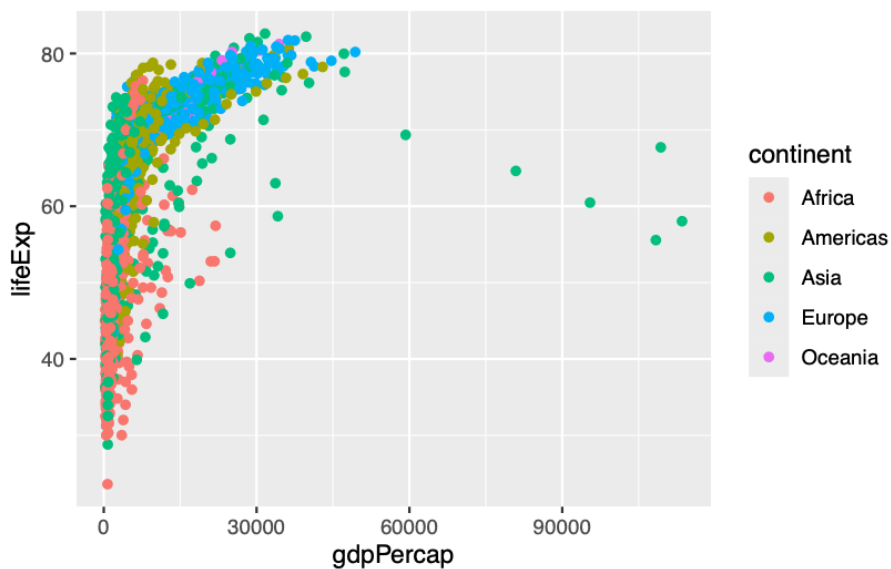
```
[1] 142
```

Q5. How many different continent values are in this dataset.

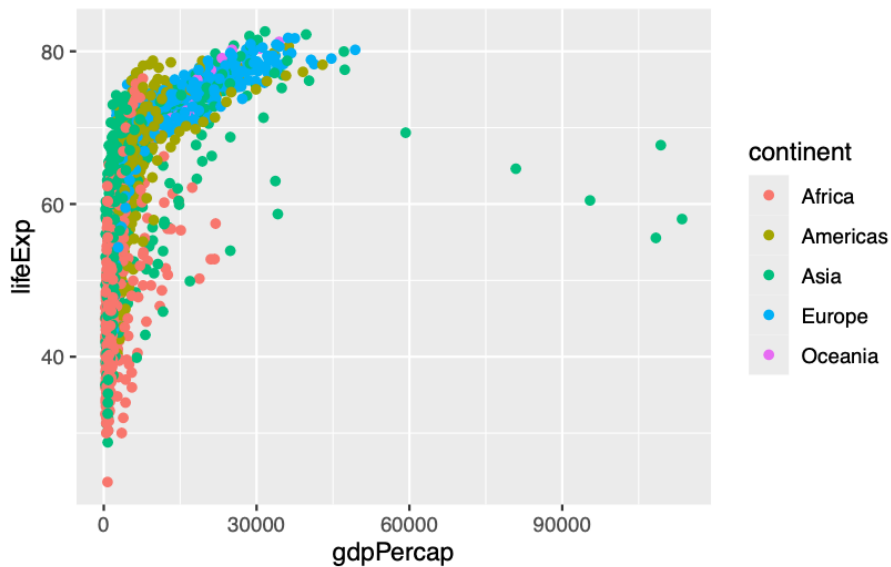
```
unique(gapminder$continent)
```

```
[1] "Asia"      "Europe"    "Africa"    "Americas"  "Oceania"
```

```
ggplot(gapminder) +  
  aes(gdpPercap, lifeExp, color=continent) +  
  geom_point()
```



```
ggplot(gapminder) +  
  aes(gdpPercap, lifeExp, color=continent, label=country) +  
  geom_point()
```

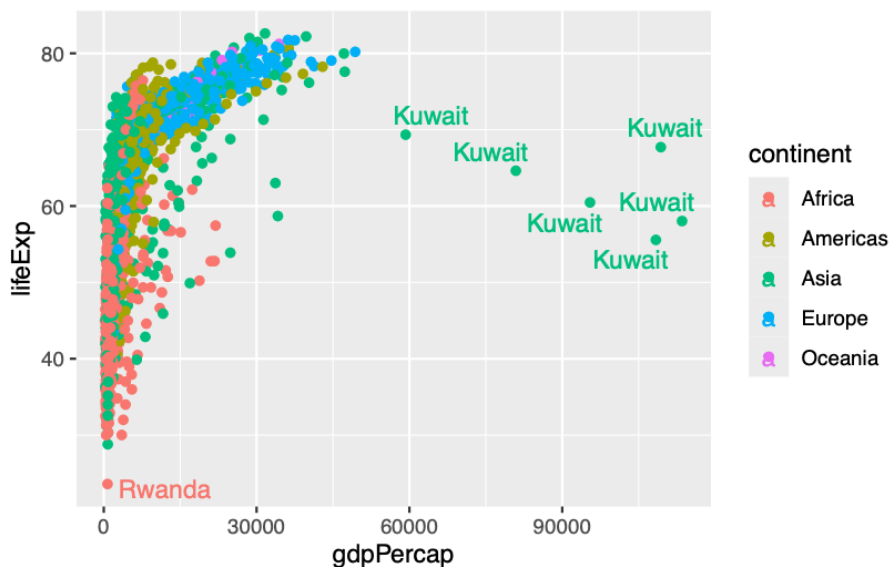



I can use the **ggrepel** package to make more sensible labels here.

```
library(ggrepel)

ggplot(gapminder) +
  aes(gdpPercap, lifeExp, color=continent, label=country) +
  geom_point() +
  geom_text_repel()
```

Warning: ggrepel: 1697 unlabeled data points (too many overlaps). Consider increasing max.overlaps



I want a separate panel per continent

```
ggplot(gapminder) +
  aes(gdpPercap, lifeExp, color=continent, label=country) +
  geom_point() +
  geom_text_repel() +
  facet_wrap(~continent)
```

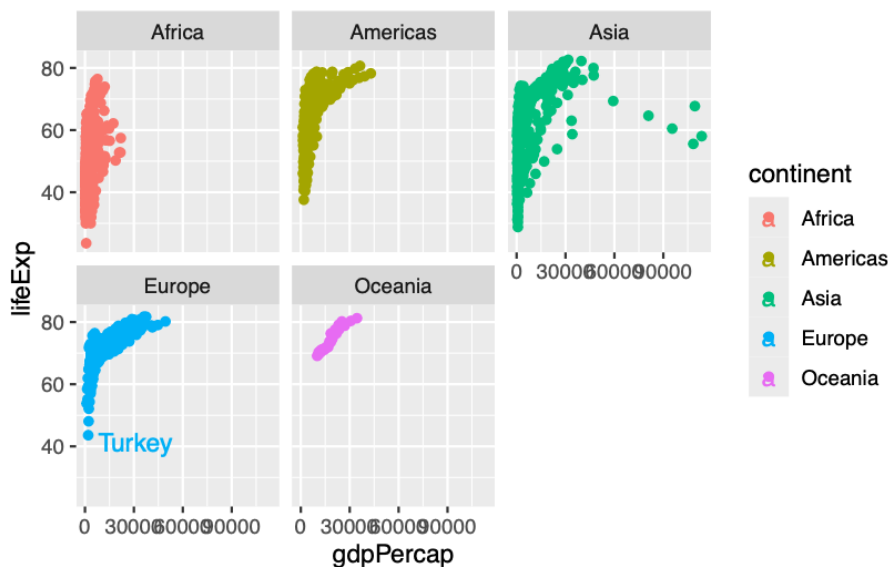
Warning: ggrepel: 624 unlabeled data points (too many overlaps). Consider increasing max.overlaps

Warning: ggrepel: 359 unlabeled data points (too many overlaps). Consider increasing max.overlaps

Warning: ggrepel: 300 unlabeled data points (too many overlaps). Consider increasing max.overlaps

Warning: ggrepel: 24 unlabeled data points (too many overlaps). Consider increasing max.overlaps

Warning: ggrepel: 396 unlabeled data points (too many overlaps). Consider increasing max.overlaps



ggplot2 offers several advantages over base R plots:

1. **Layered Grammar:** ggplot2 uses a consistent, layered approach for building plots, making it easier to add or modify elements (data, aesthetics, geoms) step by step [1], [2], [3], [5], [4].
2. **Publication Quality:** ggplot2 produces visually appealing, publication-quality figures with sensible defaults, which are harder to achieve with base R without extensive tweaking [1], [2], [3], [4].
3. **Customization:** Customizing complex plots is more straightforward in ggplot2. Adding legends, themes, or combining multiple plots is easier and less error-prone than in base R [1], [2], [3].
4. **Consistency:** The syntax and logic are consistent across different plot types, reducing the need to learn many separate functions as in base R [1], [2], [3], [5].
5. **Extensibility:** ggplot2 is part of a larger ecosystem, allowing integration with other packages for advanced graphics and analysis [1].

What do you think is the most important advantage for your own work?