Data Wrangling and Data Analysis Exploratory Data Analysis

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John Tukey (1915 – 2000)

- Data Scientist patient zero
- Inventor of:
 - The boxplot
 - The term "exploratory data analysis"
 - The Fast Fourier Transform
 - "Tukey's test"
 - The word "bit"
 - So, so much more (Wikipedia)

NO BODY OF DATA TELLS US ALL WE NEED TO KNOW ABOUT ITS OWN ANALYSIS.

IT ALWAYS TAKES INFORMATION AND INSIGHT GAINED FROM OTHER, PARALLEL BODIES TO LET US ANALYZE OUR BODY OF DATA AS WELL AS WE CAN.

- John Tukey 1970, EDA

imgflip.com

Where it fits in

- Data visualization principles
- The grammar of graphics
- Exploratory data analysis (EDA)
- Goal: know how to create and improve data visualizations, and know how to use them for exploration



Reading materials for this week

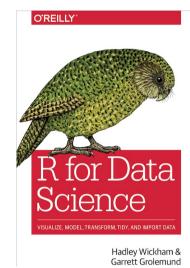
• R for Data Science (R4DS)

(Wickham & Grolemund 2017(ish))

- https://r4ds.had.co.nz
- Chapters 3,5, 7

Optional:

- Exploratory Data Analysis with R (Peng 2020)
- https://bookdown.org/rdpeng/exdata/



Exploratory Data Analysis with R



Roger D. Peng





Exploratory analysis

- Generate new insights
- Create new hypotheses
- Analysis may depend on data

Confirmatory analysis

- Test theory
- A priori hypothesis
- Analysis predefined
- Example: registered report



You work at Google and your colleague has implemented a new search algorithm for German searches. Your task is to check whether this algorithm performs better than the existing one.

• Confirmatory.

- Theory testing, hypothesis: new works better than old
- Analysis can be defined in advance: which outcome variables, how to sample from the population, which method?
- Full analysis script could be written before the data even exists



You have obtained access to your company's customer relations database. Your task is to find data-driven ways in which your company can improve customer retainment.

• Exploratory.

- Generate new insights at which touchpoint do customers drop out?
- Create new hypotheses there may be two types of customers who drop out
- Analysis cannot be defined in advance, task is to explore associations between features



- Both are necessary and complementary
- Typical scientific questions: exploratory -> confirmatory
- Then build on the tested theories to generate new insights: confirmatory -> exploratory?
- This lecture: exploratory analysis





Some good advice on data exploration



Tukey's approach to EDA

- Look at center and spread
- Find comparisons
- "Straightening and flattening":
- Use logarithms and other transforms
- Use models and residuals



hinges.

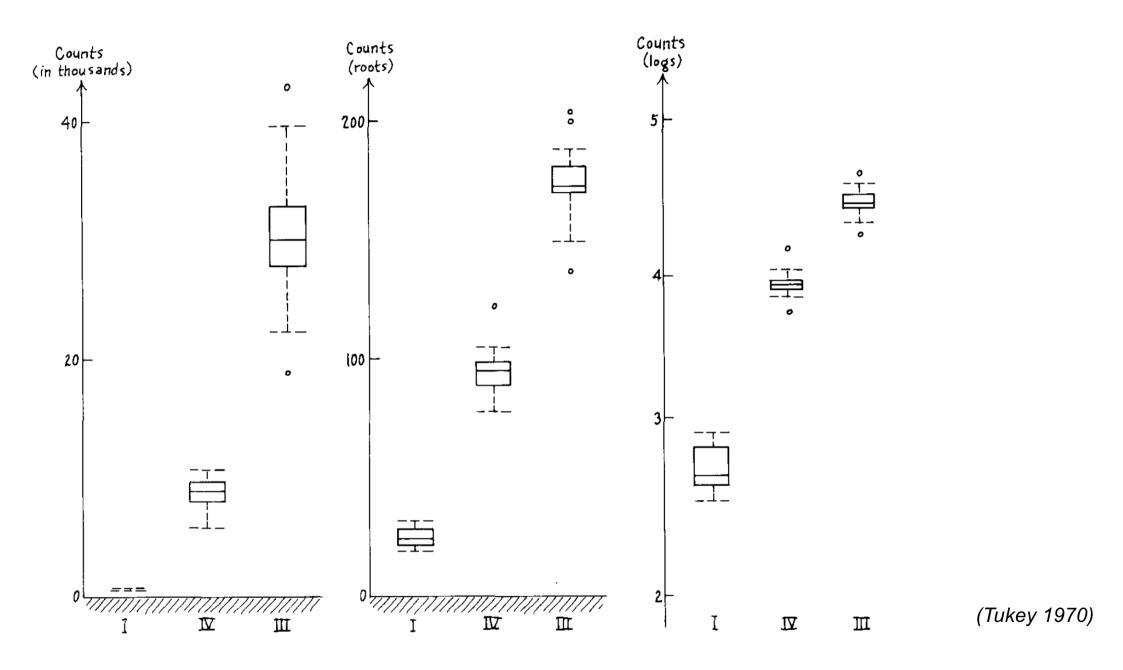
If we have 9 values in all, the 5th from either end will be the median, since $\frac{1}{2}(1+9) = 5$. Since $\frac{1}{2}(1+5) = 3$, the third from either end will be a hinge. If we have 13 values, the 7th will be the median—and the 4th from each end a hinge. In folded form, a particular set of 13 values appears as follows:

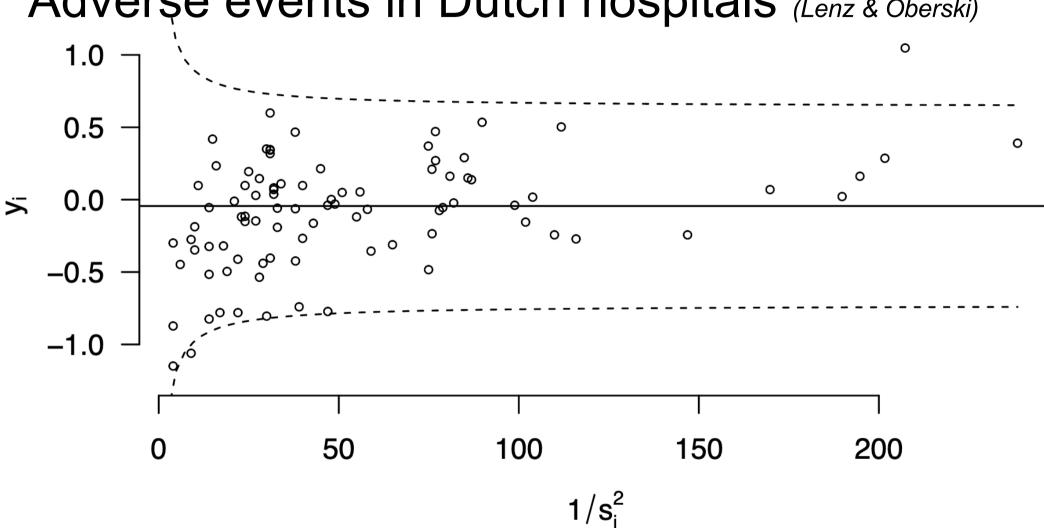
The five summary numbers are, in order, -3.2, 0.1, 1.5, 3.0, and 9.8, one at each folding point.

We usually symbolize the 5 numbers (extremes, hinges, median) that make up a

5-number summary

by a simple summary scheme like this:





Adverse events in Dutch hospitals (Lenz & Oberski)

Peng's EDA checklist

- 1. Formulate your question
- 2. Read in your data
- 3. Check the packaging, (run str())
- 4. Look at the top and the bottom of your data
- 5. Check your "n"s
- 6. Validate with at least one external data source
- 7. Try the easy solution first
- 8. Challenge your solution
- 9. Follow up

Exploratory Data Analysis with R



Roger D. Peng



Example exploration of current corona cases in the Netherlands





Dataset: COVID-19 case counts in The Netherlands

CoronaWatchNL collects numbers on COVID-19 disease count cases in **The Netherlands**. The numbers are collected from various sources on a daily base, like RIVM (National Institute for Public Health and the Environment), LCPS (Landelijk Coördinatiecentrum Patiënten Spreiding), NICE (Nationale Intesive Care Evaluatie), and the National Corona Dashboard. This project standardizes, and publishes data and makes it **Findable**, **Accessible**, **Interoperable**, **and Reusable (FAIR)**. We aim to collect a complete time series and prepare a dataset for reproducible analysis and academic use.

Read data, check packaging

url_icu <"https://raw.githubusercontent.com/J535D165/CoronaWatchNL/master/dataic/data-nice/NICE_IC_long_latest.csv"</pre>

icu <- read_csv(url_icu)</pre>

> dim(icu) [1] 1935 3



Look at top and bottom

>	> icu %>% tail			
#	A tibble: 6 x 3			
	Datum	Туре	Aantal	
	<date></date>	<chr></chr>	<dbl></dbl>	
1	2020-09-28	Toename ontslag (ziekenhuis)	0	
2	2020-09-28	Totaal opnamen (IC)	144	
3	2020-09-28	Toename opnamen (IC)	7	
4	2020-09-28	Totaal ontslag (IC)	56	
5	2020-09-28	Cumulatief opnamen (IC)	3261	
6	2020-09-28	Toename ontslag (overleden)	0	



Look at top and bottom

>	> icu %>% head			
#	# A tibble: 6 x 3			
	Datum	Туре	Aantal	
	<date></date>	<chr></chr>	<dbl></dbl>	
1	2020-02-27	Totaal ingezette IC's	5	
2	2020-02-27	Cumulatief opnamen (IC)	7	
3	2020-02-27	Totaal ontslag (IC)	0	
4	2020-02-27	Cumulatief ontslag (ziekenhuis)	0	
5	2020-02-27	Cumulatief ontslag (overleden)	0	
6	2020-02-27	Toename ontslag (overleden)	0	



Validate with external source > C' 介 https://www.coronatracker.com/country/netherlands/

Corona Tracker Travel Alert What is COVID-19 Home Netherlands Overview Share: **f** 🍤 5.6% Fatality 114,540 6,380 3 OF TOTAL CASES Confirmed Recovered Deaths +6 new deaths +2,914 new cases Critical Cases treated in ICU **Daily Cases Receiving Treatment** 144 0 0.1% of total cases 0.0% of total cases

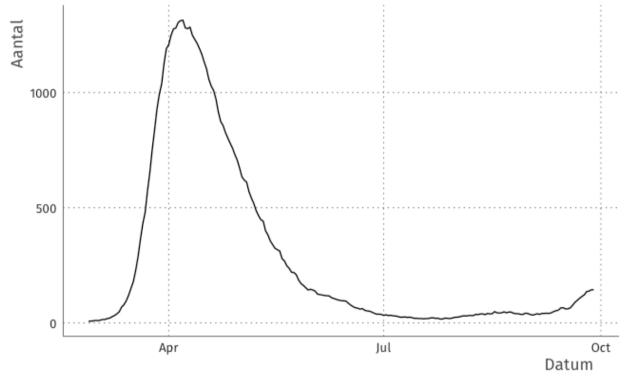
Check n's

> icu %>% group_by(Type) %>% summarize(n())

A tibble: 9×2 `n()` Type <int> <chr> 1 Cumulatief ontslag (overleden) 215 2 Cumulatief ontslag (ziekenhuis) 215 3 Cumulatief opnamen (IC) 215 4 Toename ontslag (overleden) 215 5 Toename ontslag (ziekenhuis) 215 6 Toename opnamen (IC) 215 7 Totaal ingezette IC's 215 8 Totaal ontslag (IC) 215 9 Totaal opnamen (IC) 215



```
icu %>%
  filter(Type == "Totaal opnamen (IC)") %>%
  ggplot(aes(Datum, Aantal)) + geom_line() +
  theme_fira()
```



Utrecht University

Understand what you're looking at

Bezetting IC units

		IC-units in Nederland		
		gem.	min.	max.
صغا رصغا رصغا	bedden	14	1	35
\$ °	intensivisten (in fte)	7	1	18
****	verpleegkundigen (in fte)	50	10	152

(84 ICUs)

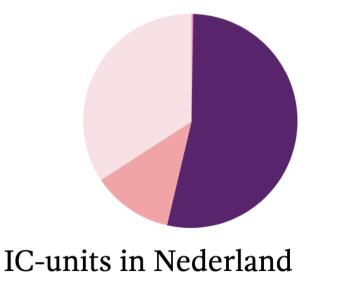
Back-of-the-envelope:

14×84 ≈ 1200 beds

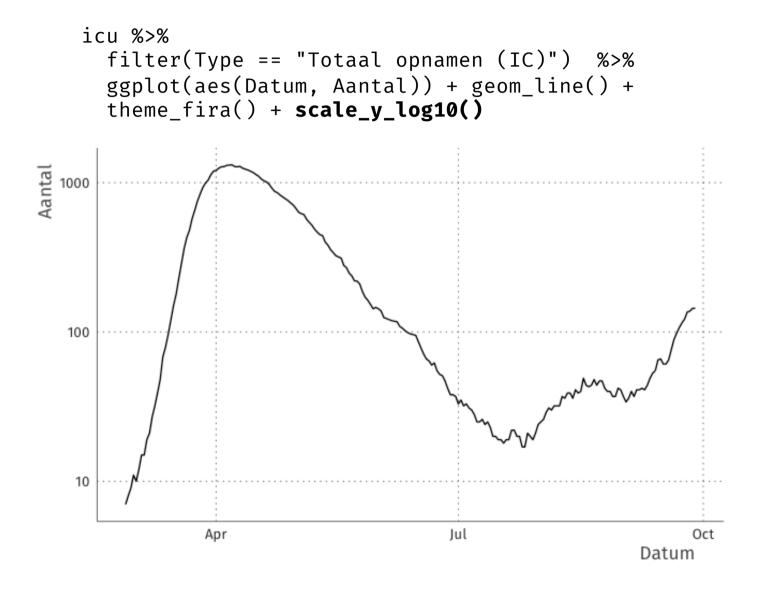
Geometric average: $\sqrt{1 \times 35} \approx 6$ beds

Understand what you're looking at

Aantal opnamen per opnametype



		IC-units in Nederland	
	Туре	aantal	%
	Alle opnamen	73.979	100
	Medisch	39.654	53,6
	Spoed chirurgie	8.632	11,7
	Geplande chirurgie	25.499	34,5
	Overige	194	0,3

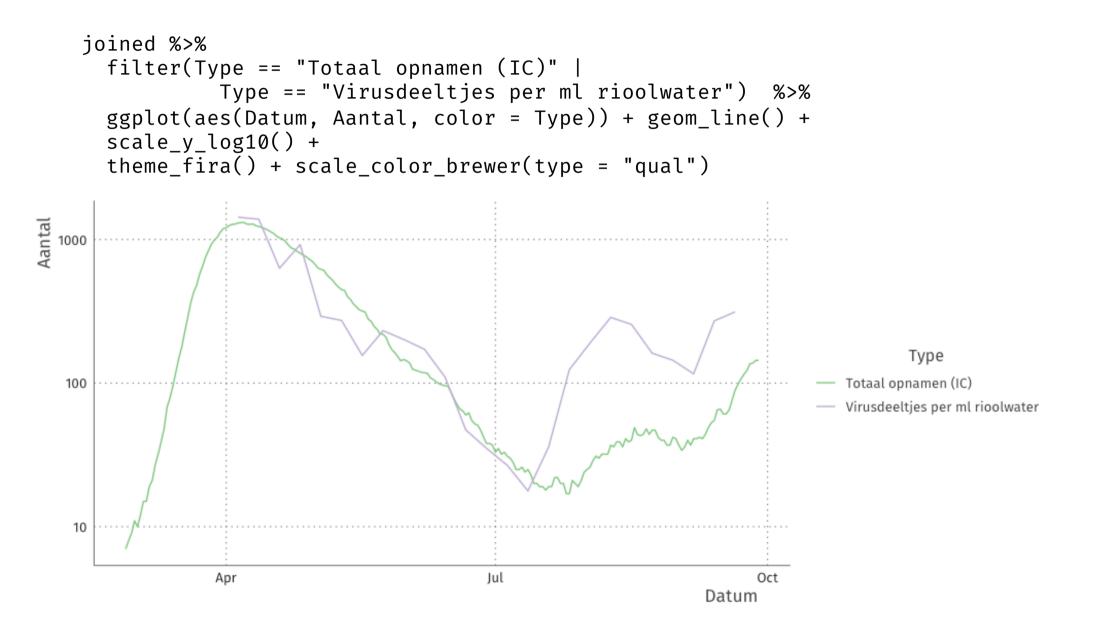


```
url_sewage <-
"https://raw.githubusercontent.com/J535D165/CoronaWatchNL/master/data-
dashboard/data-sewage/RIVM_NL_sewage_counts.csv"</pre>
```

sewage <- read_csv(url_sewage)</pre>

joined <- bind_rows(sewage, icu)</pre>

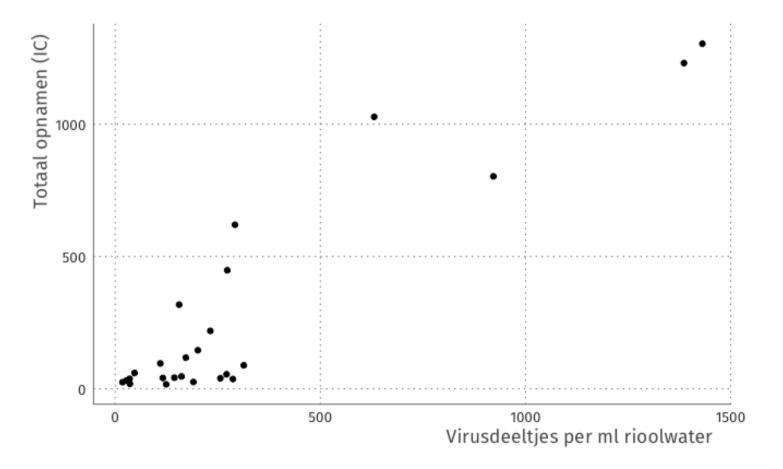


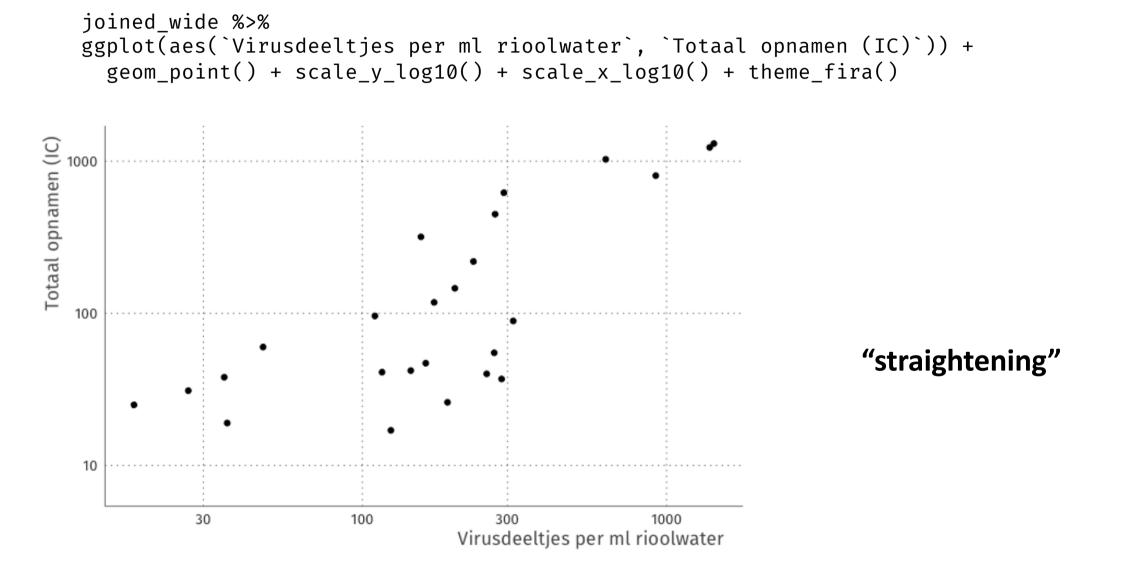


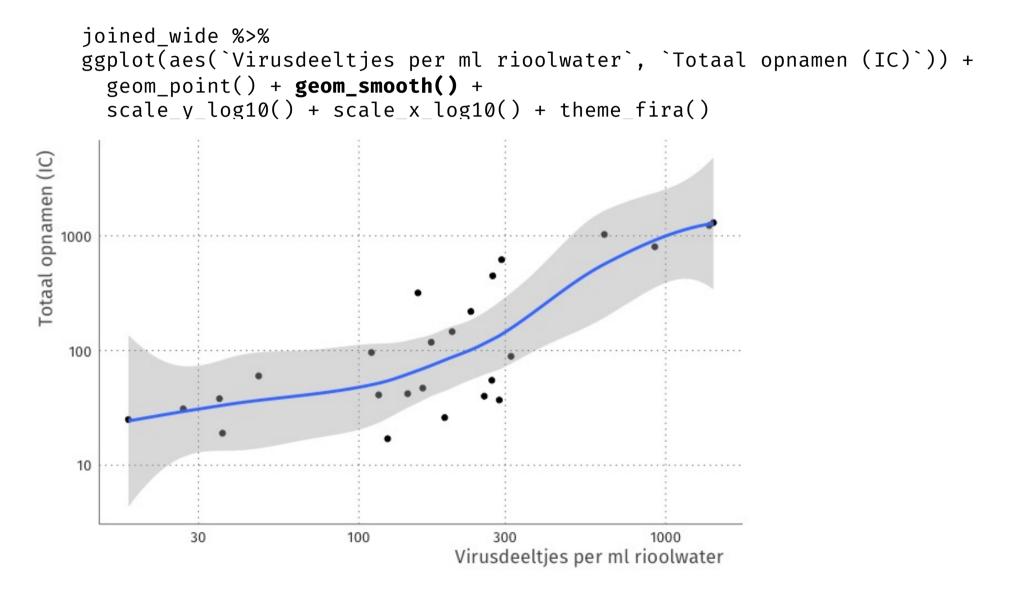
```
joined_wide <- joined %>%
filter(Type == "Totaal opnamen (IC)" |
        Type == "Virusdeeltjes per ml rioolwater") %>%
pivot_wider(names_from = Type, values_from = Aantal)
```

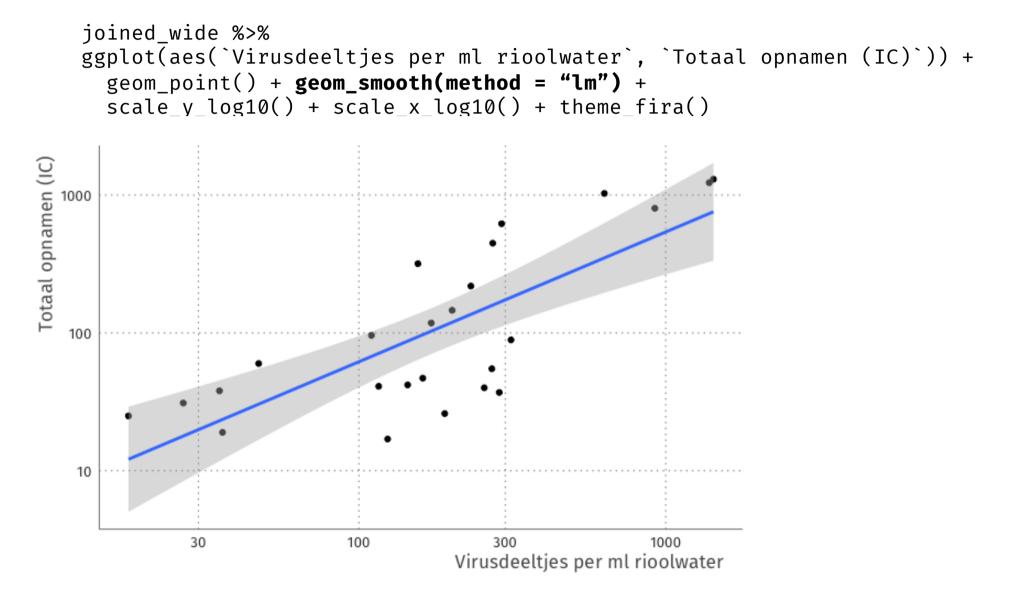


```
joined_wide %>%
ggplot(aes(`Virusdeeltjes per ml rioolwater`, `Totaal opnamen (IC)`)) +
  geom_point() + theme_fira()
```





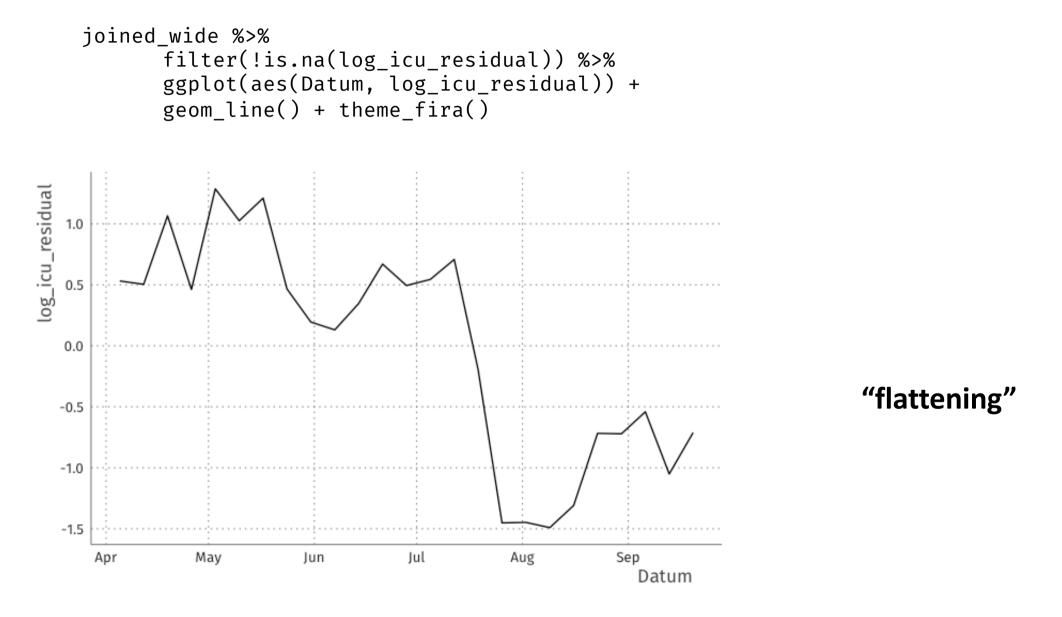


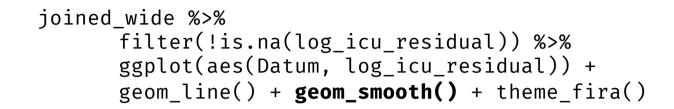


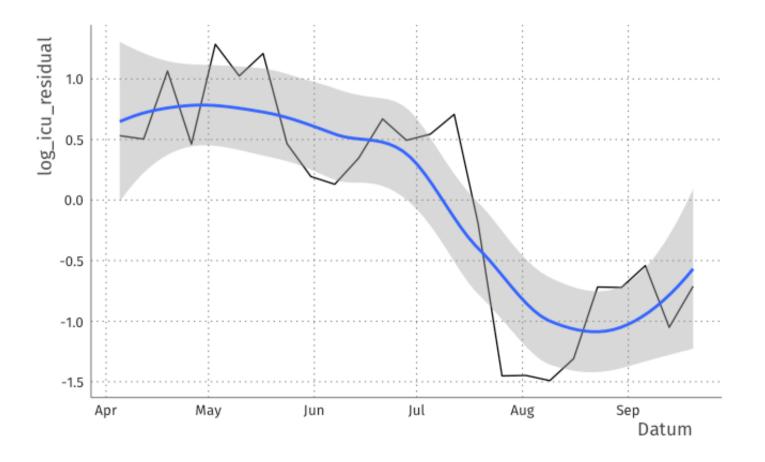
Looking at **residuals**

```
joined_wide <- joined_wide %>%
  mutate(
    log_icu = log(`Totaal opnamen (IC)` + 1),
    log_sewage = log(`Virusdeeltjes per ml rioolwater` + 1))
fit_lm <- lm(log_icu ~ log_sewage, data = joined_wide)
joined_wide <- joined_wide %>%
  mutate(log_icu_predicted = predict(fit_lm, newdata = .),
    log_icu_residual = log_icu - log_icu_predicted)
```



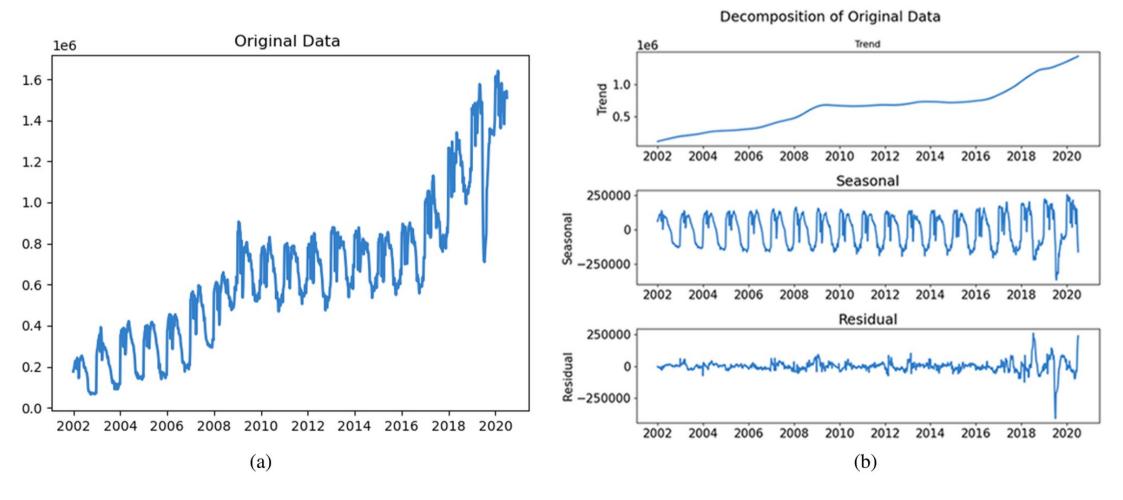








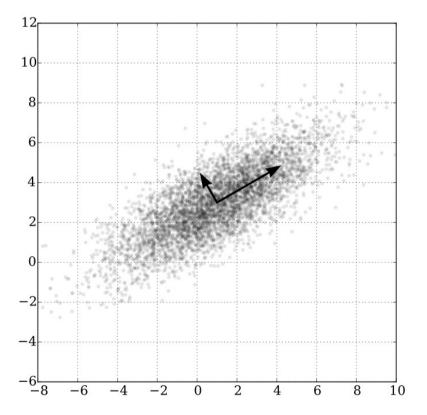
"Flattening" in time series analysis



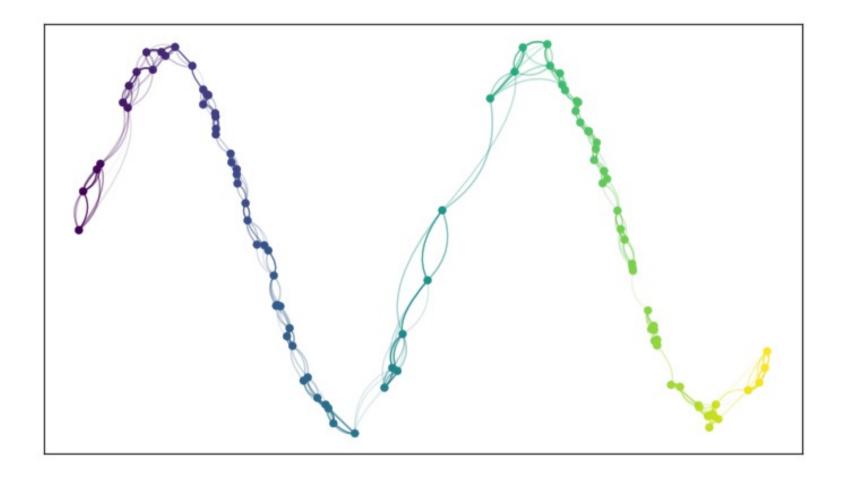
High-dimensional "flattening"

Principal component analysis (PCA)

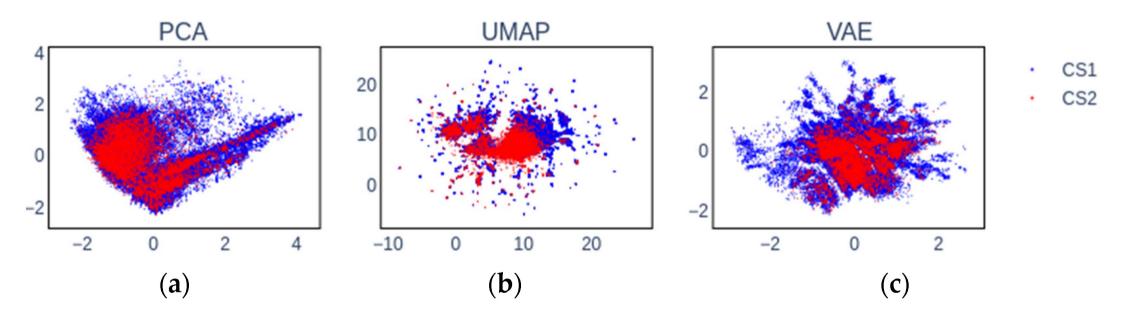
- Reduce dimensions, see which points end up close together
- Works for linearly related features
- Terrible for nonlinear relations













Modern high-dimensional "flattening"

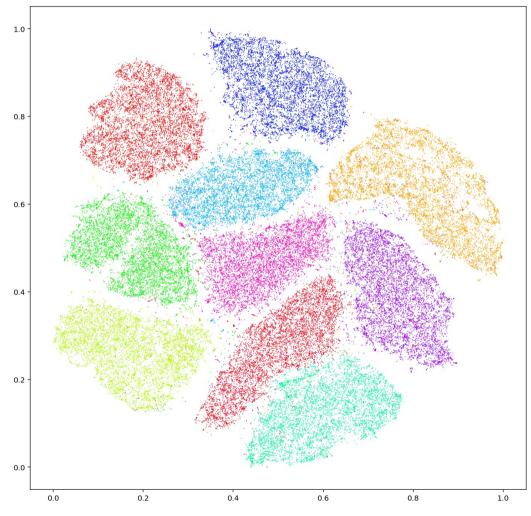
t-SNE

(Van der Maaten & Hinton 2008)

- MNIST dataset (right)
- Minimize a Cauchy distance

$$q_{ij} = rac{(1+\|\mathbf{y}_i-\mathbf{y}_j\|^2)^{-1}}{\sum_k \sum_{l
eq k} (1+\|\mathbf{y}_k-\mathbf{y}_l\|^2)^{-1}}$$

- Large distances are meaningless
- Short distances highly informative



Modern high-dimensional "flattening"

UMAP

(McInnes & Healy 2018)

- Single-cell dataset (right)
- Force-directed graph layout
- Faster than t-SNE
- Can deal with very high-D data





Using some standard plots for EDA



Visual EDA: two questions

- Variation: How are the features distributed?
 - Univariate
 - Specific: among people, among time, among flights
 - General: among the unit of measurement, among examples
 - In tabular data, generally one example/unit per row
- Association: What type of covariation occurs in the data?
 - Which features covary? When A is high, is B high?
 - Multidimensional, multivariate.



Visual EDA: two questions

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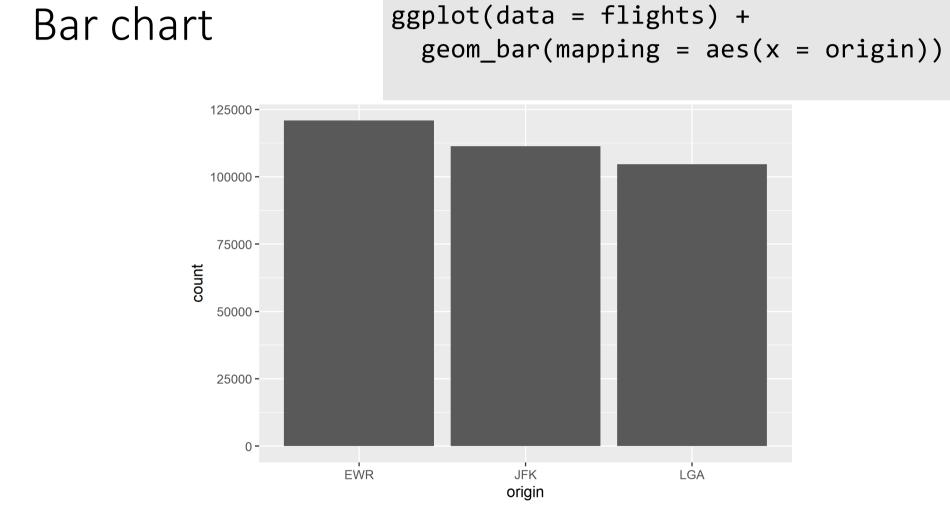
<pre>library(nycflights13)</pre>								
flights								
#> # A tibble: 336,776 x 19								
#>	year	month	day	dep_time	<pre>sched_dep_time</pre>	dep_delay	arr_time	sched_arr_time
#>	<int></int>	<int></int>	<int></int>	<int></int>	<int></int>	<dbl></dbl>	<int></int>	<int></int>
#> 1	2013	1	1	517	515	2	830	819
#> 2	2013	1	1	533	529	4	850	830
<i>#> 3</i>	2013	1	1	542	540	2	923	850
<i>#></i> 4	2013	1	1	544	545	-1	1004	1022
#> 5	2013	1	1	554	600	-6	812	837
#> 6	2013	1	1	554	558	-4	740	728
<pre>#> # with 3.368e+05 more rows, and 11 more variables: arr_delay <dbl>,</dbl></pre>								
<pre>#> # carrier <chr>, flight <int>, tailnum <chr>, origin <chr>, dest <chr>,</chr></chr></chr></int></chr></pre>								
<pre>#> # air_time <dbl>, distance <dbl>, hour <dbl>, minute <dbl>, time_hour <dttm></dttm></dbl></dbl></dbl></dbl></pre>								



Bar chart

- X position: feature of interest
- Y position: count of occurrences (statistical transformation)
- Geom: bars/rectangle
- For categorical features e.g., flight origin







Bar chart

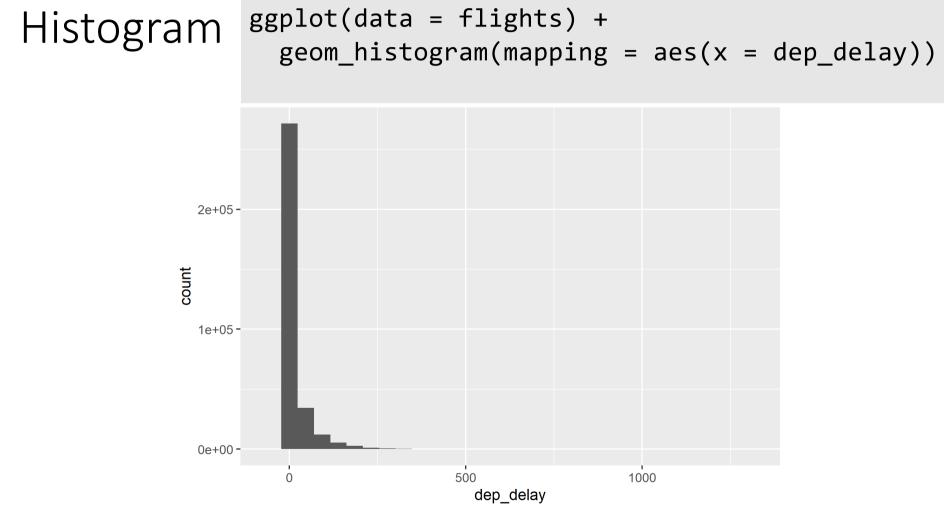
- Fewest flights depart from LGA
- Almost 125000 flights from EWR



Histogram

- X position: feature of interest
- Y position: count of occurrences in bins (statistical transformation)
- Geom: bars/rectangle
- For continuous features, e.g., departure delay in minutes





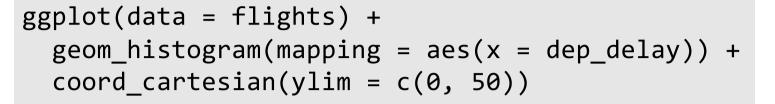


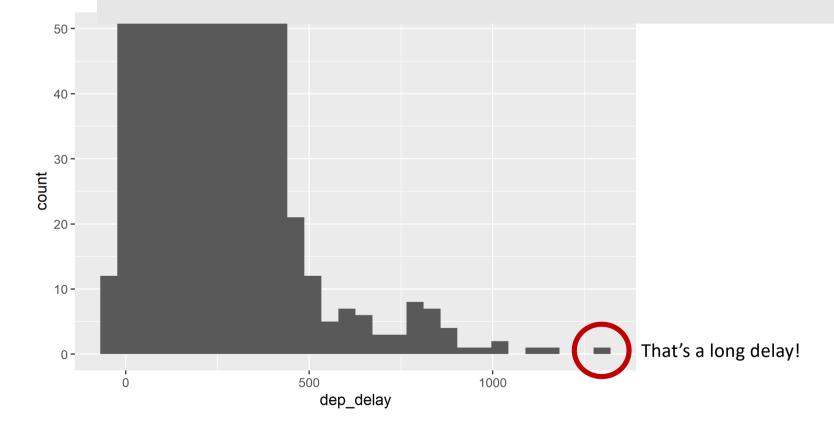
Histogram

 New question: why the strange distribution? What happens at the tail end?



Histogram (zoomed)







Histogram

- Most flights are approximately on time
- There are even some flights that depart early
- There are a few flights with more than 1000 minutes delay

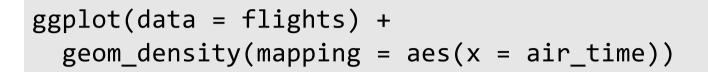


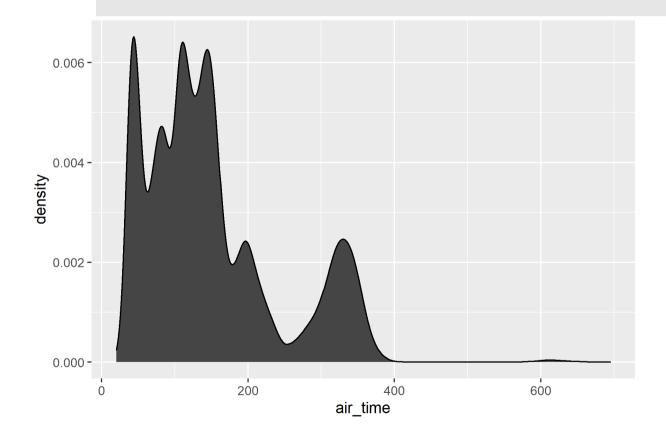
Density

- X position: feature of interest
- Y position: density (statistical transformation, smoothed histogram)
- Geom: polygon / line
- For continuous features, e.g., airtime in minutes



Density







Density

- Most flights from NY are under 200 minutes (3.3 hours)
- Few flights are between 200 and 250 minutes
- Quite some flights are between 250 and 400 minutes
- Some flights are over 600 minutes (weird bump? Remember!)
- Air times are not normally distributed!



Visual EDA: two questions

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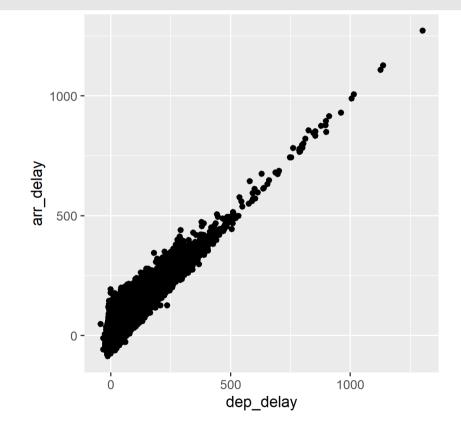
Scatter plot

- X position: feature A of interest (in regression, generally predictor)
- Y position: feature B of interest (in regression, outcome)
- Geom: points/dots
- For continuous features, e.g., departure delay in minutes and arrival delay in minutes



Scatter plot

ggplot(data = flights) +
 geom_point(aes(x = dep_delay, y = arr_delay))





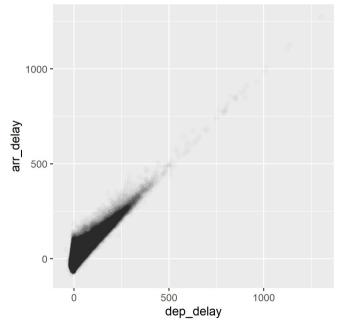
Scatter plot

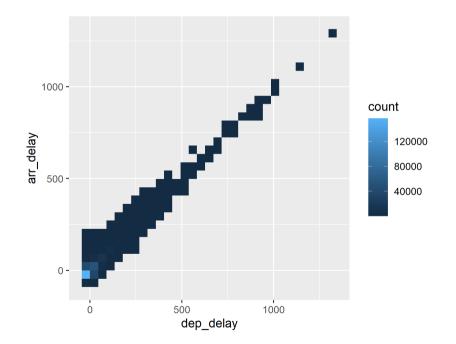
- Departure and arrival delay correlate strongly
- Arrival delay is generally higher than departure delay
- Only a few delays are above 500 minutes



Solutions to overplotting

• Transparency (alpha) or binning (geom_bin2d)

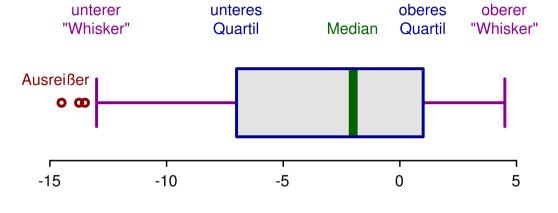






Box plot (Tukey)

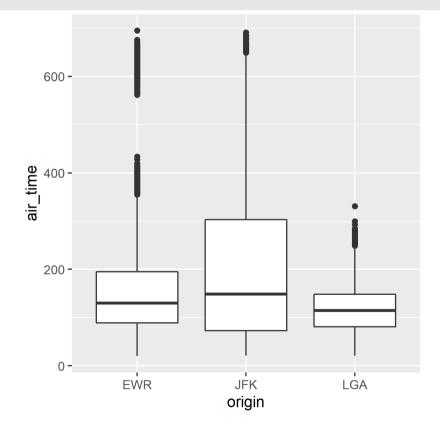
- X position: feature A of interest
- Y position: feature B of interest
- Geom: rectangles for box, lines (whiskers), points for outliers
- Statistical transformations:
- median, 25% and 75% percentile (inter quartile range, IQR), 1.5×IQR for "whiskers"
- Continuous vs. categorical, e.g., origin and airtime





Box plot

ggplot(data = flights) + geom_boxplot(aes(x = origin, y = air_time))





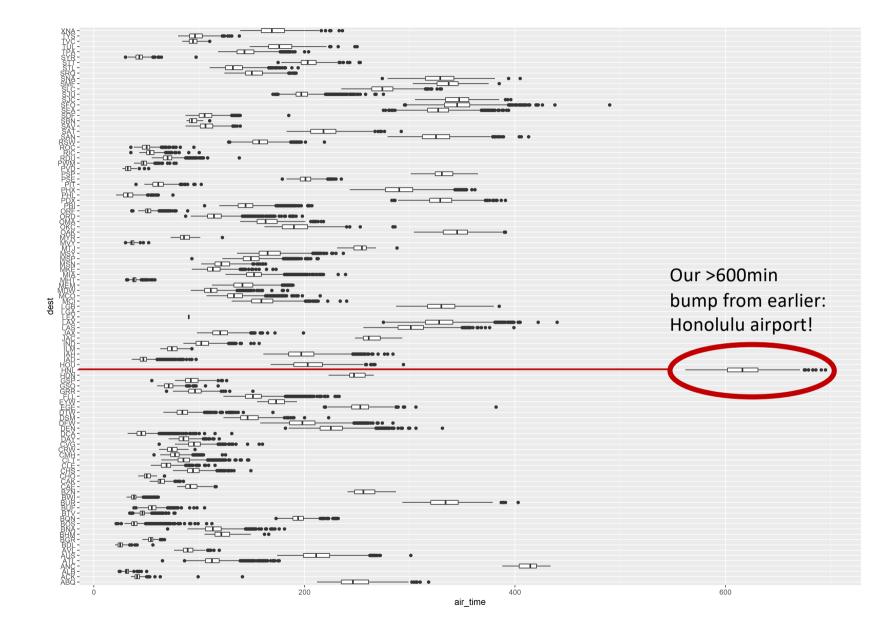
Box plot

- Flights from LGA do not have the >600 minute bump
- Flights from JFK take longest, on average (median)
- Flight times from JFK have largest IQR

• Let's look at destination rather than origin?

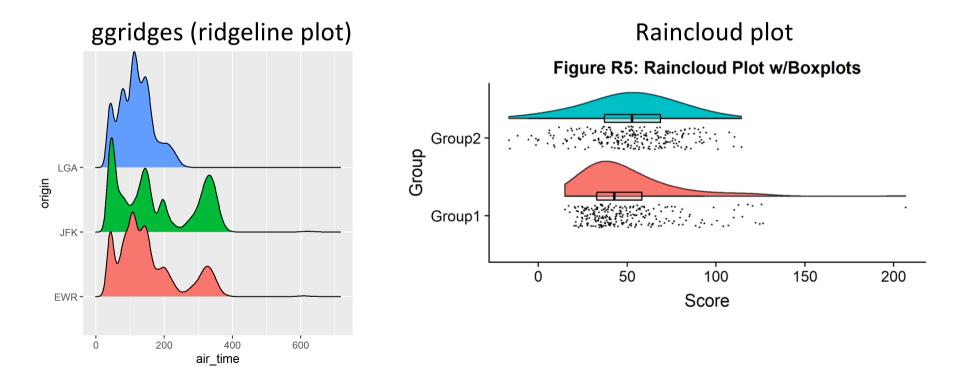


Box plot by destination





Many options available for categorical vs continuous





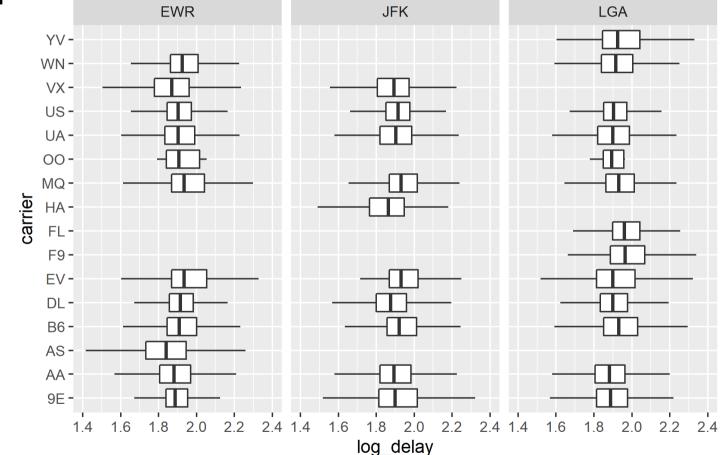
Facets

- Faceting a plot means creating a subplot per category
- Also called "small multiples" (tufte)
- Allows for another categorical variable in the visualisation
- Danger: clutter

```
+ facet_wrap(~feature)
```



Boxplot of (log) delay by carrier, faceted by origin





Facets

- Some carriers do not fly from some airports (missing data)
- Delays are quite similar across origin airports and carriers

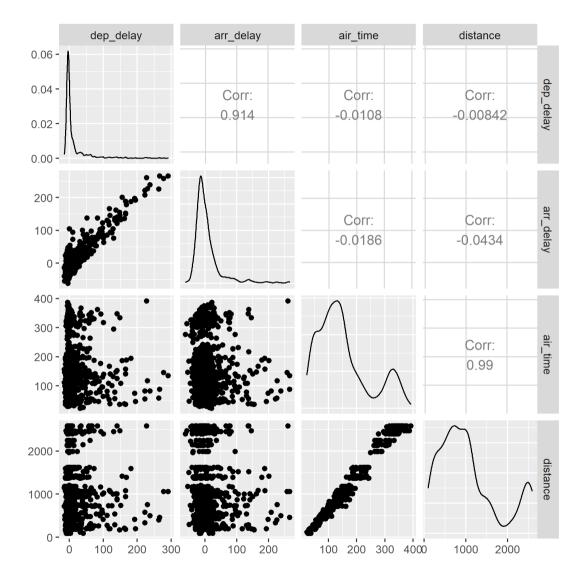


Pairs plot

- Pairs plot: a scatter plot for each variable
- Multivariate
- Difficult in ggplot: facet by variable (not a feature in the original dataset)
- Data frame needs to be
- Package available: GGally (ggpairs)









Pairs plot

- Air time and distance correlate strongly
- Departure and arrival delay correlate strongly
- Air time / distance do not correlate with delay
- In other words: distance cannot predict delays well



Conclusion

- Exploration is key to understanding things you did not know already
- EDA walks a fine line between seeing useful and less useful things (overfitting, if you like)
- Some useful principles are:
 - Peng's checklist
 - Understanding what you're seeing
 - Finding interesting comparisons
 - "Straightening and flattening" (using models and residuals)
 - Standard graphs to look at variation and association

