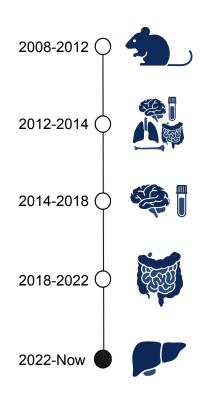
Communicating Biology Through Data Visualization

Rachel Edgar

University Health Network - University of Toronto Postdoctoral Fellow - MacParland and Bader labs

About Me: Rachel Edgar



Somatic mutations in the cerebellum of harlequin mice

Meta-analysis of Human DNA Methylation Data

Investigating the use of blood as surrogate for brain tissue in epigenetics

Multiomics Of The Human Gut Toward Insight Into Childhood Inflammatory Bowel

Single-cell gene expression of human liver

University of Western Ontario

BSc - Kathleen Hill

University of British Columbia

MSc - Paul Pavlidis

University of British Columbia

Research Assistant - Michael Kobor

University of Cambridge - EMBL-EBI

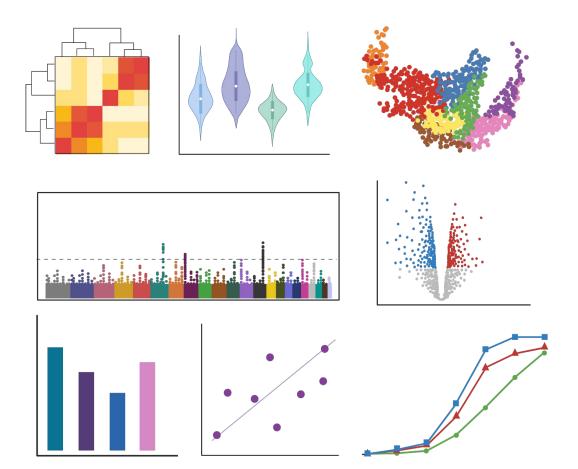
PhD - Daniel Zerbino

University Health Network

Postdoc - Sonya MacParland and Gary Bader

Above all else show the data.

- Edward R. Tufte The Visual Display of Quantitative Information



Data visualizations are powerful tools

Data is often incomprehensible in its raw form

- Visualization allows you to:
 - **Troubleshoot** your analysis
- **Understand** data for yourself

- **Generate** hypothesis
- - **Communicate** your new understanding to others
- 3456 2877 2523

3775

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3486

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4171

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4126

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2440

3608

3813

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3171

148

3453

2363

3782

2774

2783

3631

3493

2784

2002

4181

4135

3756

4134

2001

2443

3615

3820

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47.21551

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- 71751 69321
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-82.31692

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-107.193

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-93.6515

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-82.26806

-123 660

-110.3961

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-120.94772021-10-15 2021-09-22 2021-08-27 2021-08-27

2021-11-14

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2021-10-30

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2021-07-24

2021-07-18

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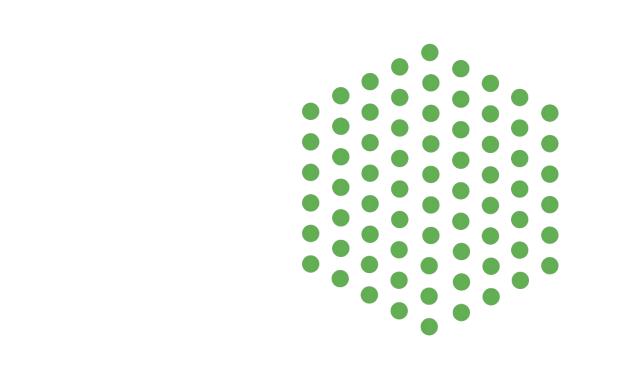
We create figures so that **humans** understand our data

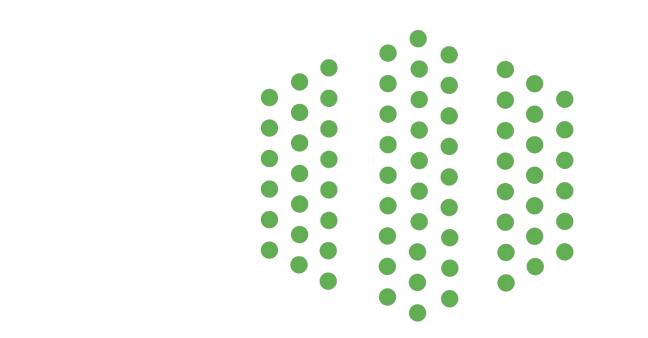
Gestalt principles of grouping describe human perception

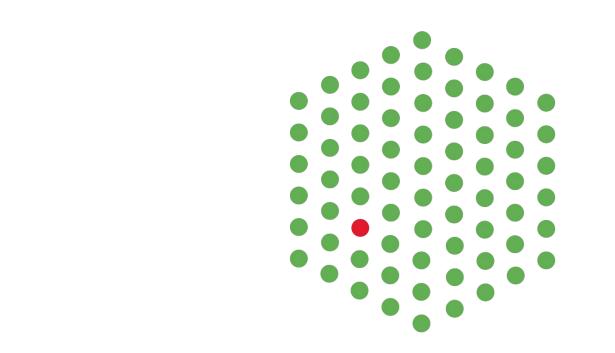
How do humans perceive?

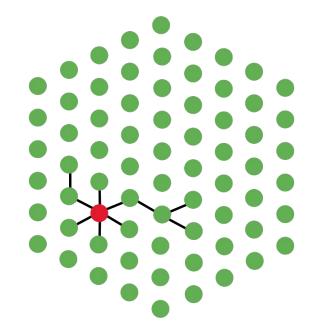
Humans naturally perceive objects as organized patterns and objects - law of *Prägnanz*

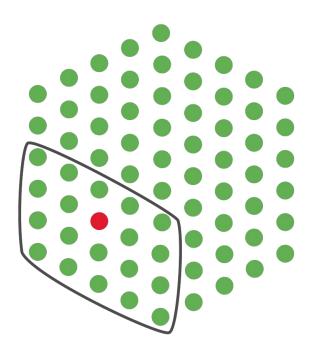
Similarity Proximity Connection Enclosure

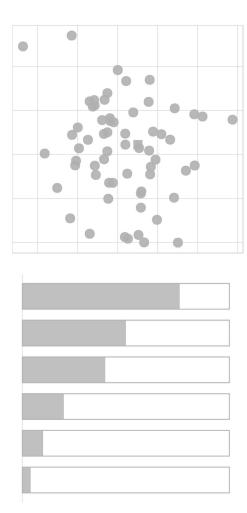


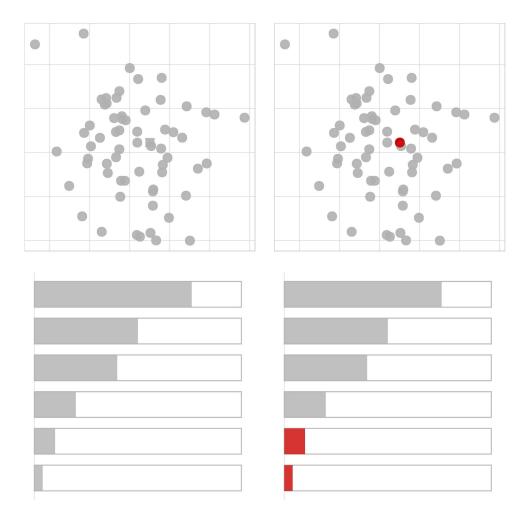


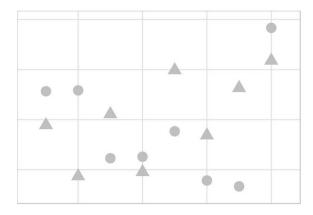


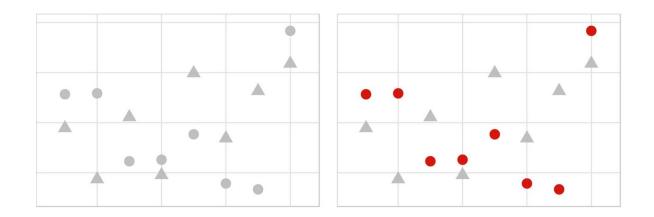


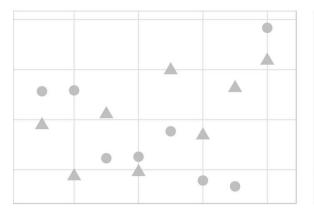


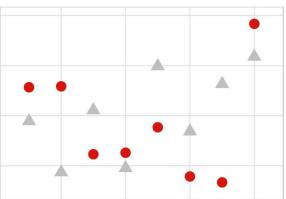


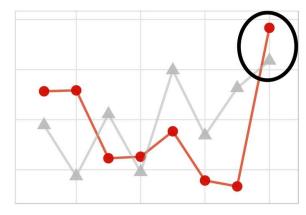








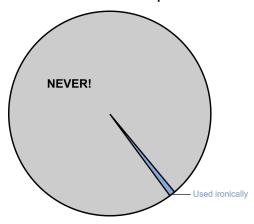




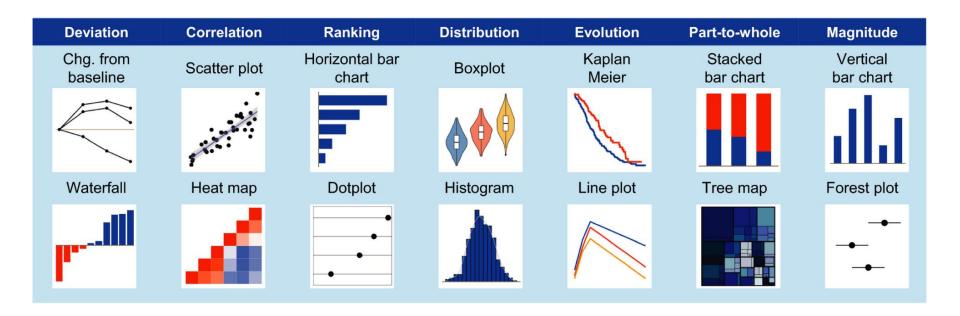
Effective Visual Communication

- 1. Have a defined purpose
- 2. Show the data clearly
 - a. No pie charts
- 3. Make the message obvious

When is it OK to use a pie chart?



What is the right type of plot for my data?



Graphs can be just for you or for communication

Analysis graphs

- Just for you
- See patterns
- Help design next questions or decide on better visualizations

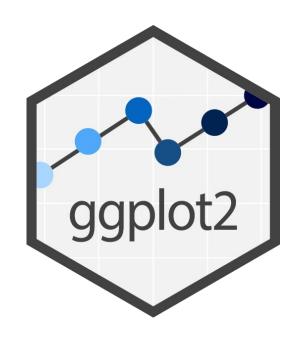
Communication graphs

- Make you point as clearly (and simply) as possible
- More is **not** better
- These will be refined versions of analysis graphs

How to get inspired?

- It can help to draw the plot you are imaging on paper
- Not imagining anything? Shamelessly steal and copy!
 - Try to recreate plots you see (in papers etc) for a dataset you have
 - I often google image search a description of what I am imaging to see similar examples and then try to remake these

Now you have a plot in mind how do you make it?



The grammar of graphics - ggplot2

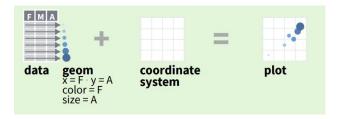


data: a data frame: quantitative, categorical; local or database query **aes**thetic mapping of variables into visual properties: size, color, x, y **geom**etric objects ("geom"): points, lines, areas, arrows, ...

The grammar of graphics - ggplot2



data: a data frame: quantitative, categorical; local or database query **aes**thetic mapping of variables into visual properties: size, color, x, y **geom**etric objects ("geom"): points, lines, areas, arrows, ...



The grammar of graphics - ggplot2

Aesthetics **aes**() make data visible:

x, y: variables

colour: colours the lines of geometries

fill: fill geometries

group: groups based on the data

shape: defines the shape (point, triangles)

linetype: defines the type of line (solid, dashed)

size: define sizes of elements

alpha: changes the transparency

The grammar of graphics - basic structure

The grammar of graphics - basic structure

```
ggplot (data = <DATA>) +

<GEOM_FUNCTION> (mapping = aes(<MAPPINGS>),

stat = <STAT>, position = <POSITION>) +

<COORDINATE_FUNCTION> +

<FACET_FUNCTION> +

<SCALE_FUNCTION> +

<THEME_FUNCTION>
```

Example data: Penguins!

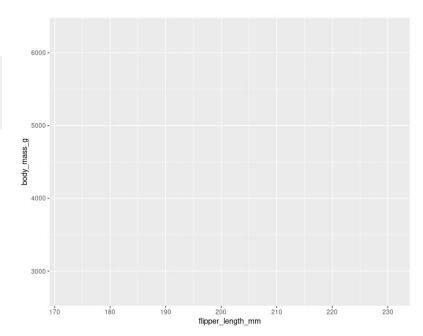
species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g	sex	year
Adelie	Torgersen	39.1	18.7	181	3750	male	2007
Adelie	Torgersen	41.8	19.4	198	4450	male	2008
Adelie	Biscoe	37.8	18.3	174	3400	female	2007
Gentoo	Biscoe	43.5	14.2	220	4700	female	2008
Chinstrap	Dream	45.4	18.7	188	3525	female	2007



```
ggplot(
  data = penguins
)
```

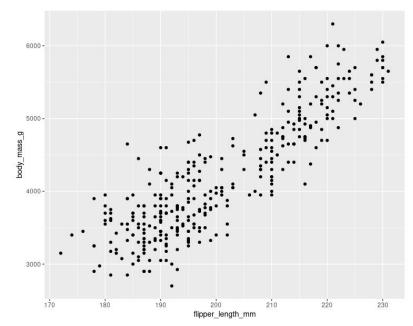
species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g	sex	year
Adelie	Torgersen	39.1	18.7	181	3750	male	2007
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Gentoo	Biscoe	43.5	14.2	220	4700	female	2008
Chinstrap	Dream	45.4	18.7	188	3525	female	2007

```
ggplot(
  data = penguins,
  mapping = aes(x = flipper_length_mm, y = body_mass_g)
)
```

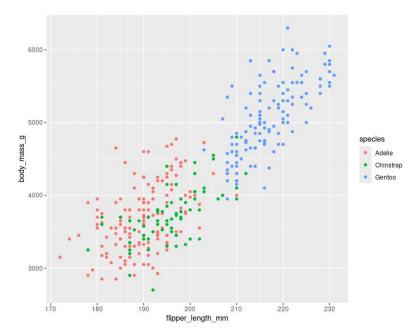


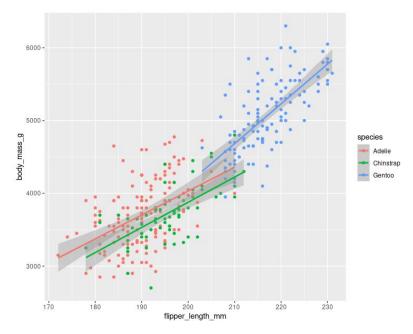
species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g	sex	year
Adelie	Torgersen	39.1	18.7	181	3750	male	2007
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Gentoo	Biscoe	43.5	14.2	220	4700	female	2008
Chinstrap	Dream	45.4	18.7	188	3525	female	2007

```
ggplot(
  data = penguins,
  mapping = aes(x = flipper_length_mm, y = body_mass_g)
) +
  geom_point()
```

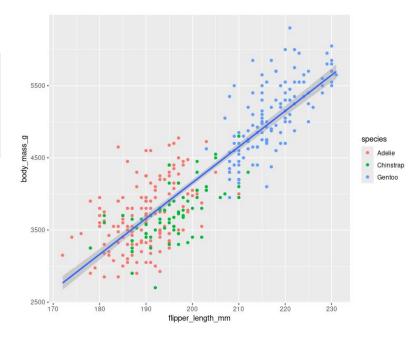


species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g	sex	year
Adelie	Torgersen	39.1	18.7	181	3750	male	2007
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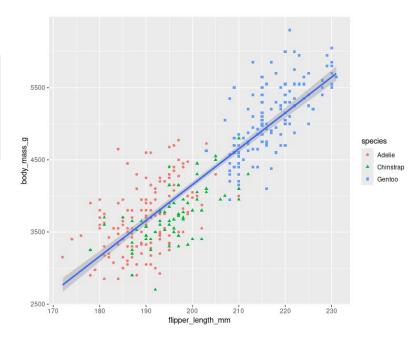




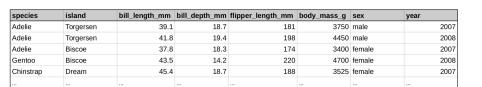
```
ggplot(
  data = penguins,
  mapping = aes(x = flipper_length_mm, y = body_mass_g)
) +
  geom_point(aes(color = species)) +
  geom_smooth(method = "lm")
```

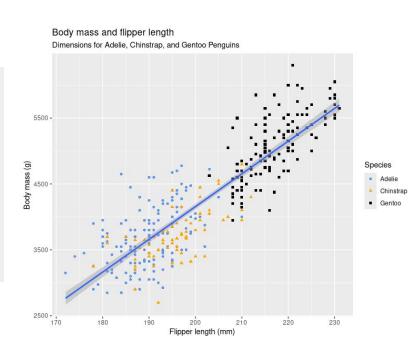


```
ggplot(
  data = penguins,
  mapping = aes(x = flipper_length_mm, y = body_mass_g)
) +
  geom_point(aes(color = species, shape = species)) +
  geom_smooth(method = "lm")
```



```
ggplot(
  data = penguins,
  mapping = aes(x = flipper_length_mm, y = body_mass_g)
) +
  geom_point(aes(color = species, shape = species)) +
  geom_smooth(method = "lm") +
  labs(
    title = "Body mass and flipper length",
    subtitle = "Dimensions for Adelie, Chinstrap, and Gentoo
Penguins",
    x = "Flipper length (mm)", y = "Body mass (g)",
    color = "Species", shape = "Species")+
scale_color_manual(values=c("cornflowerblue","orange","black"))
```





Workshop overview and Example Plots

redgar598.github.io/COSS_data_vis_R

How did set-up go?

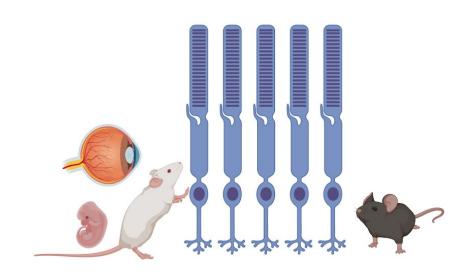


Data We Will Be Using in ggplot Introduction

Gene expression data from mouse photoreceptors

Samples from different developmental stages (E16,P2,P6,P10 and 4 weeks)

Two mouse lines, a wildtype (**wt**) and knockouts for rod cell specific transcription factor (**NrIKO**)



Plots Made in Workshop