

# Data Visualisation with R

## Workshop Day 1

### Scales and color

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MONASH University

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Scales



# Diamonds data

The diamonds data is part of ggplot2 

```
glimpse(diamonds)
```

```
## Rows: 53,940
## Columns: 10
## $ carat    <dbl> 0.23, 0.21, 0.23, 0.29, 0.31, 0.24, 0.24, 0.26, 0.22, 0.23, 0...
## $ cut       <ord> Ideal, Premium, Good, Premium, Good, Very Good, Very Good, Ver...
## $ color     <ord> E, E, E, I, J, J, I, H, E, H, J, J, F, J, E, E, I, J, J, J, I, ...
## $ clarity   <ord> SI2, SI1, VS1, VS2, SI2, VVS2, VVS1, SI1, VS2, VS1, SI1, VS1, ...
## $ depth     <dbl> 61.5, 59.8, 56.9, 62.4, 63.3, 62.8, 62.3, 61.9, 65.1, 59.4, 64...
## $ table     <dbl> 55, 61, 65, 58, 58, 57, 57, 55, 61, 61, 55, 56, 61, 54, 62, 58...
## $ price     <int> 326, 326, 327, 334, 335, 336, 336, 337, 337, 338, 339, 340, 34...
## $ x         <dbl> 3.95, 3.89, 4.05, 4.20, 4.34, 3.94, 3.95, 4.07, 3.87, 4.00, 4...
## $ y         <dbl> 3.98, 3.84, 4.07, 4.23, 4.35, 3.96, 3.98, 4.11, 3.78, 4.05, 4...
## $ z         <dbl> 2.43, 2.31, 2.31, 2.63, 2.75, 2.48, 2.47, 2.53, 2.49, 2.39, 2...
```

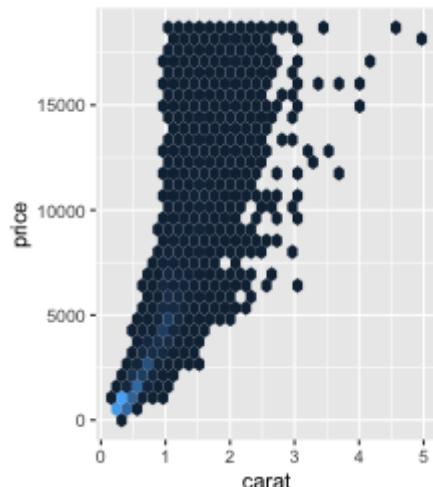
# Scales

- Scales control the mapping from **data to aesthetics**.

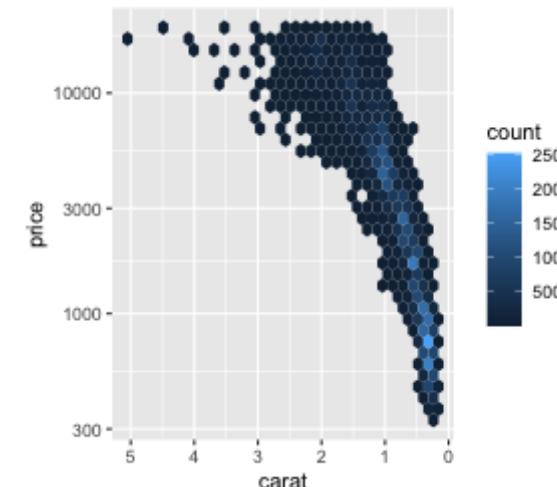
`scale_<aesthetic>_<type>`

```
g <- ggplot(diamonds, aes(carat, price) ) + geom_hex()
```

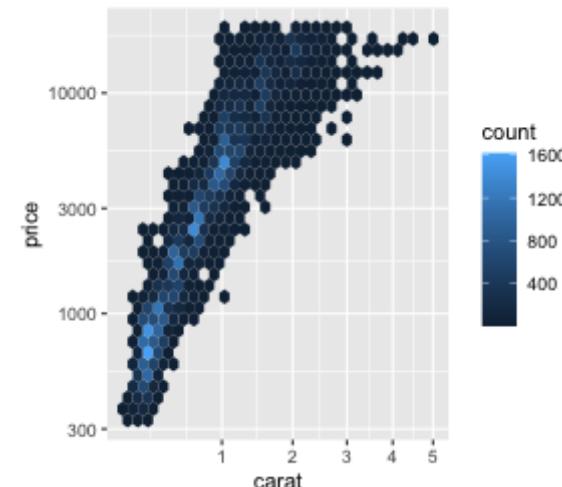
```
g + scale_y_continuous() +  
  scale_x_continuous()
```



```
g + scale_x_reverse() +  
  scale_y_continuous(trans="log10")
```



```
g + scale_y_log10() +  
  scale_x_sqrt()
```



# scale

scales	Description
scale_alpha, scale_alpha_continuous, scale_alpha_binned, scale_alpha_discrete, scale_alpha_ordinal, scale_alpha_datetime, scale_alpha_date	Alpha transparency scales
scale_x_binned, scale_y_binned	Positional scales for binning continuous data (x & y)
scale_colour_brewer, scale_fill_brewer, scale_colour_distiller, scale_fill_distiller, scale_colour_fermenter, scale_fill_fermenter, scale_color_brewer, scale_color_distiller, scale_color_fermenter	Sequential, diverging and qualitative colour scales from colorbrewer.org
scale_colour_continuous, scale_fill_continuous, scale_colour_binned, scale_fill_binned, scale_color_continuous, scale_color_binned	Continuous and binned colour scales

Previous

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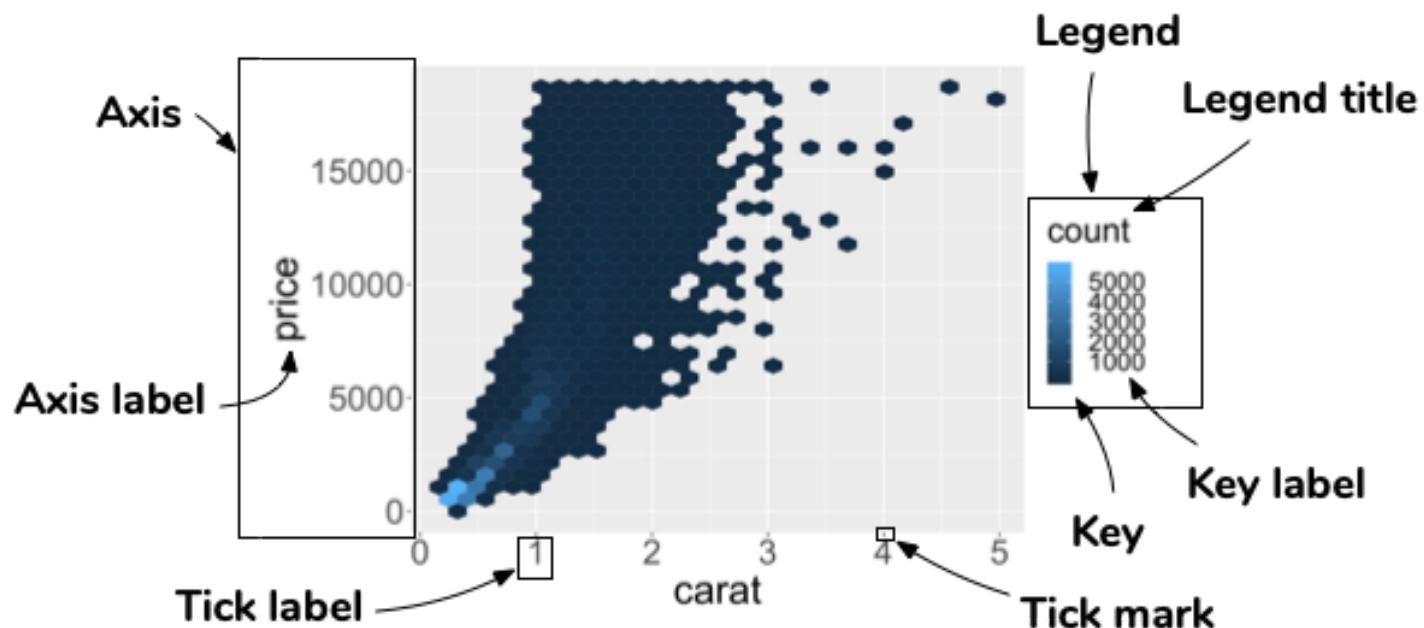
4

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Next

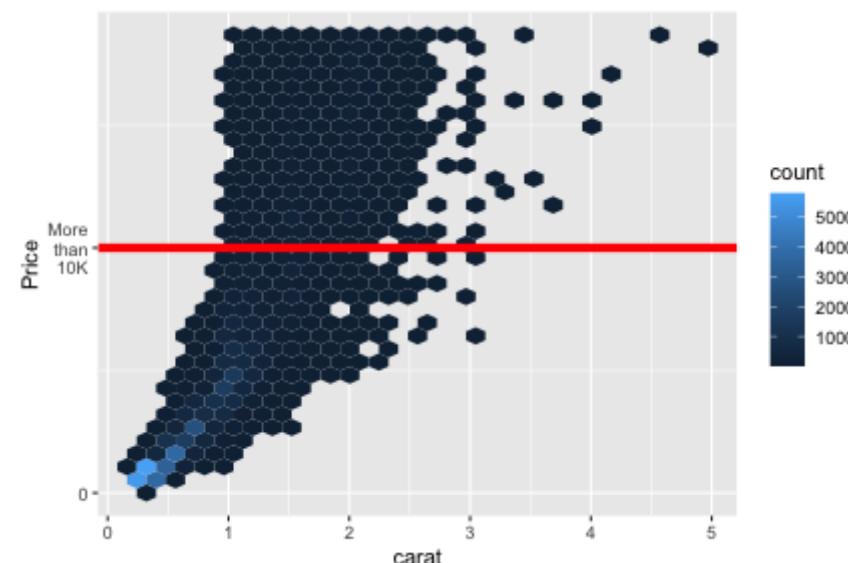
# Guide: an axis or a legend

- The scale creates a **guide**: an **axis** or **legend**.
- So to modify these you generally use **scale\_\*** or other handy functions (guides, labs, xlab, ylab and so on).



# Modify axis

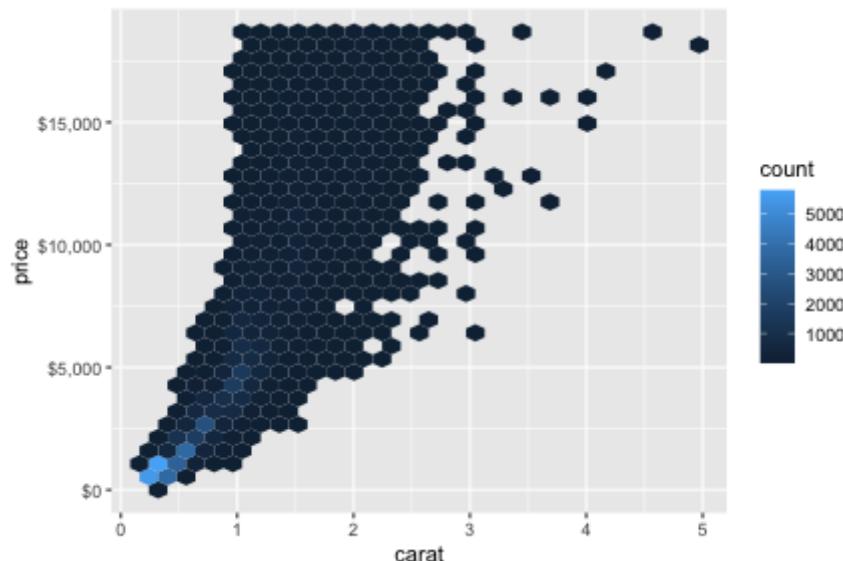
```
g +  
  scale_y_continuous(name = "Price",  
                     breaks = c(0, 10000),  
                     labels = c("0", "More\n than\n 10K")) +  
  geom_hline(yintercept = 10000, color = "red", size = 2)
```



# Nicer formatting functions in scales

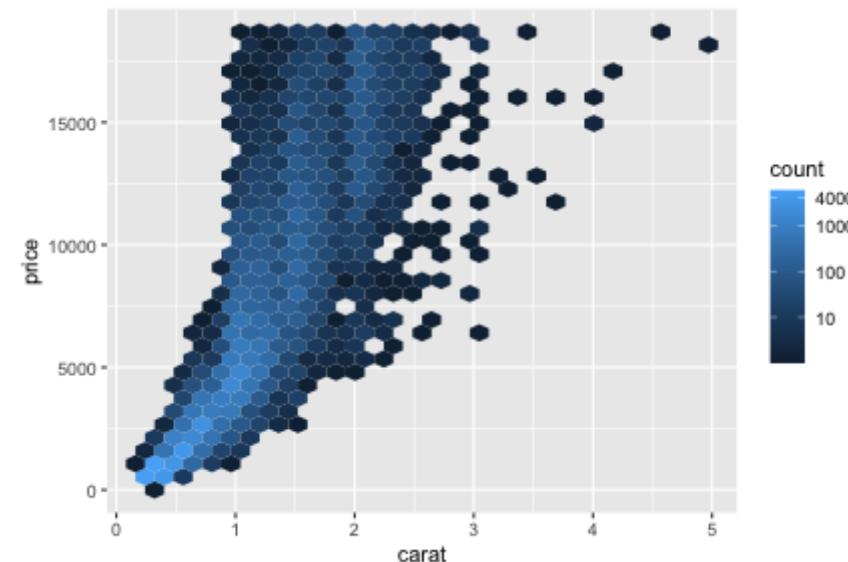


```
g +  
  scale_y_continuous(  
    label = scales::dollar_format()  
)
```



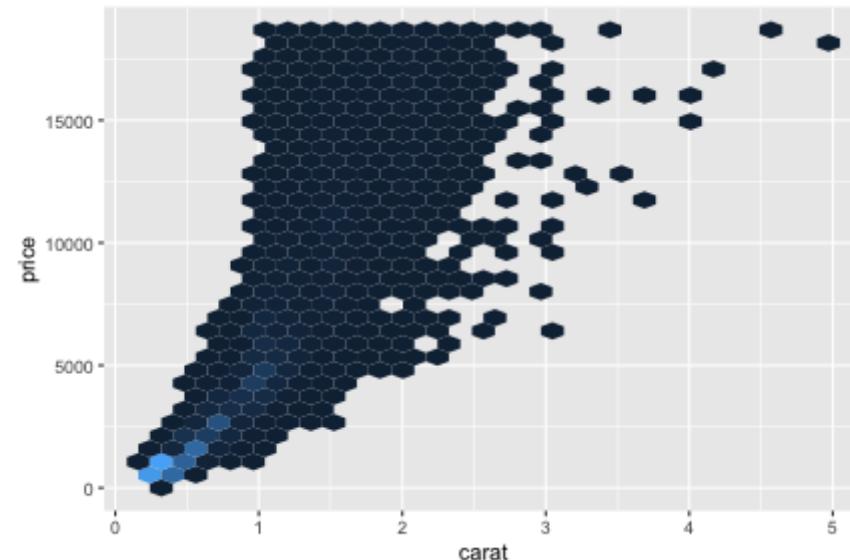
# Modifying legend

```
g +  
  scale_fill_continuous(  
    breaks = c(0, 10, 100, 1000, 4000),  
    trans = "log10"  
)
```



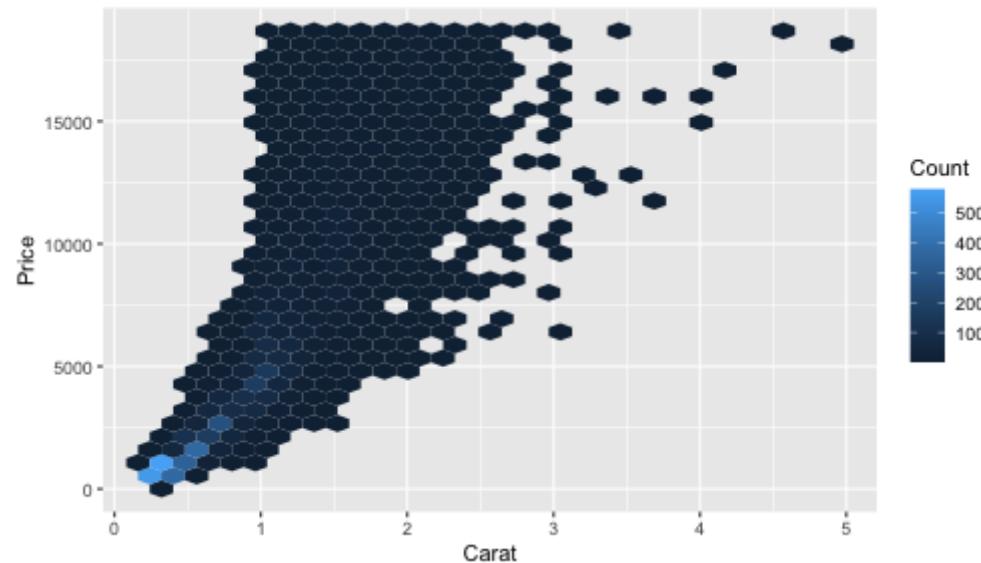
# Removing legend

```
g +  
  scale_fill_continuous(  
    guide = "none"  
)
```



# Alternative control of guides

```
g +  
  ylab("Price") + # Changes the y axis label  
  labs(x = "Carat", # Changes the x axis label  
        fill = "Count") # Changes the legend name
```



```
g + guides(fill = "none") # remove the legend
```

# 4

# Color space

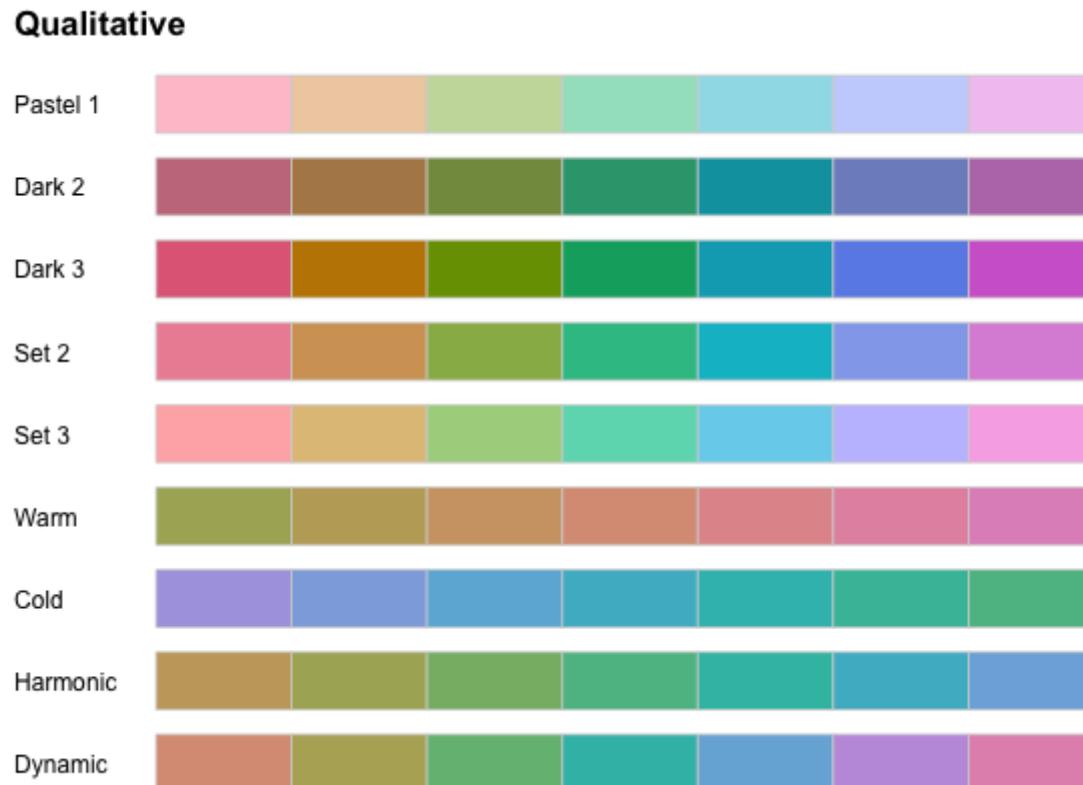
Zeileis, Fisher, Hornik, Ihaka, McWhite, Murrell, Stauffer, Wilke (2019). *colorspace: A Toolbox for Manipulating and Assessing Colors and Palettes*. *arXiv 1903.06490*

Zeileis, Hornik, Murrell (2009). Escaping RGBland: Selecting Colors for Statistical Graphics. *Computational Statistics & Data Analysis* 53(9) 3259-3270

# Qualitative palettes

designed for categorical variable with no particular ordering

```
colorspace::hcl_palettes("Qualitative", plot = "TRUE", n = 7)
```



# Sequential palettes

designed for ordered categorical variable or number going from low to high (or vice-versa)

```
colorspace::hcl_palettes("Sequential", plot = "TRUE", n = 7)
```



# Diverging palettes

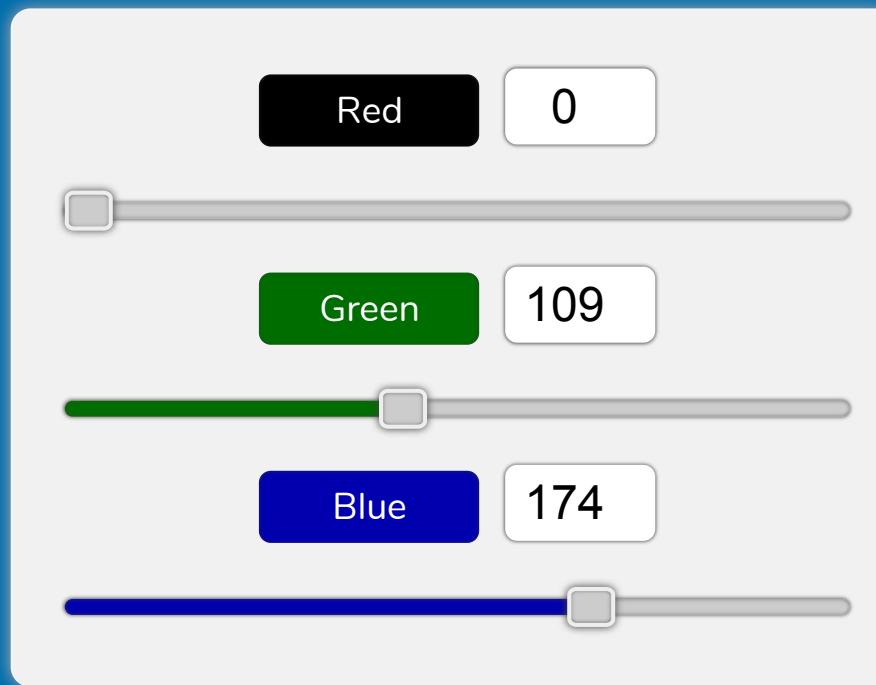
designed for ordered categorical variable or number going from low to high (or vice-versa) with a neutral value in between

```
colorspace::hcl_palettes("Diverging", plot = "TRUE", n = 7)
```



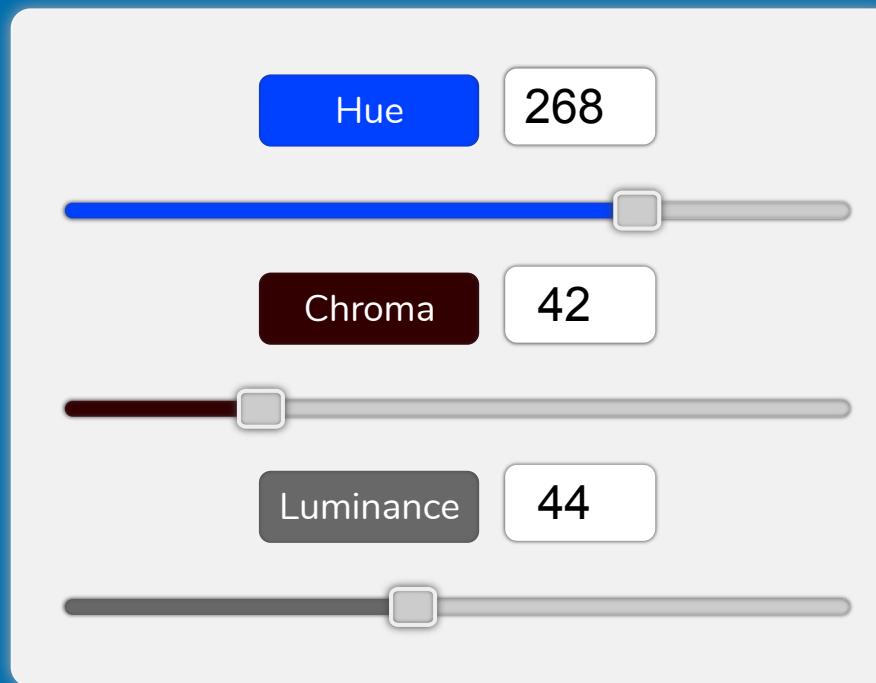
# RGB color space

made for screen projection



# HCL color space

made for human visual system



# colorspace



Interactively choose/create a palette using the HCL color space.

```
library(colorspace)  
hcl_wizard() # OR choose_palette()
```

LIVE DEMO

# hcl\_wizard

## Base Options

Type of palette  
Basic: Sequential (multi-hue)

Base color scheme  
Purple-Blue

Example  
Map

## Control Options

Reverse  
 Correct colors  
 Dark mode  
 Desaturated

**Vision**  
 Normal  
 Deutan  
 Protan  
 Tritan

## Color Settings

HUE 1: 300  
HUE 2: 200  
CHRON: 60  
CHRON: NA  
LUMIN.: 25  
LUMIN.: 95  
POWER: 0.7  
POWER: 1.3  
NUMBER: 7  
NUMBER: 40

Return to R

Example Plot Spectrum Color Plane Export Info

RAW GrADS Python matlab R Register

If you use R the preferred way to handle color palettes is to use [choose\\_palette\(\)](#) or [hclwizard\(\)](#) on your local machine. Both graphical user interfaces return an R color palette function. However, you can also use the function call below to use the current palette in your R scripts.

```
## Custom color palette
sequential_hcl(n = 7, h = c(300, 200), c = c(60, NA, 0), l = c(25, 95), power = c(0.7, 1.3), register = )
```

Custom color palettes can also be registered to be able to call custom palettes by name. If the optional argument register = "Custom-Palette" is set (where "Custom-Palette" is the name of your new palette) the palette will be added to the list of available color palettes. Existing palettes can also be overruled. Note that the graphical interfaces ([choose\\_palette\(\)](#)) will not use custom palettes. These register-calls can also be added to your local `~/.Rprofile`.

```
## Register custom color palette
colorspace::sequential_hcl(n = 7, h = c(300, 200), c = c(60, NA, 0), l = c(25, 95),
power = c(0.7, 1.3), register = "Custom-Palette")
```

R colorspace 1.4.1

Choose your palette > Export > R > Copy the command

# Registering your own palette

```
library(colorspace)
# register your palette
sequential_hcl(n = 7,
               h = c(300, 200),
               c = c(60, 0),
               l = c(25, 95),
               power = c(2.1, 0.8),
               register = "my-set")
# now generate from your palette
sequential_hcl(n = 3,
               palette = "my-set")
## [1] "#6B0077" "#7C8393" "#F1F1F1"
```

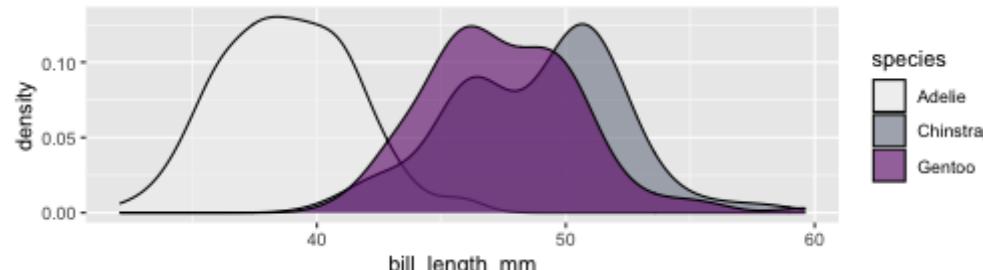
```
hcl_palettes(n = 5, palette = "my-set", plot = T)
```

## Sequential (multi-hue)

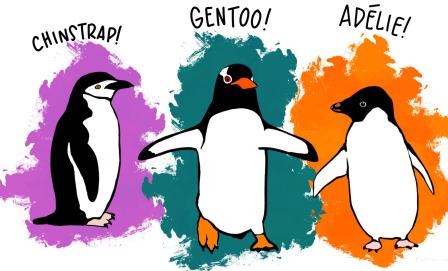


## Combining with ggplot:

```
ggplot(penguins,
       aes(bill_length_mm, fill = species)) +
  geom_density(alpha = 0.6) +
  # notice here you don't need to specify the n!
  scale_fill_discrete_sequential(palette = "my-set")
```



# Manually selecting colors

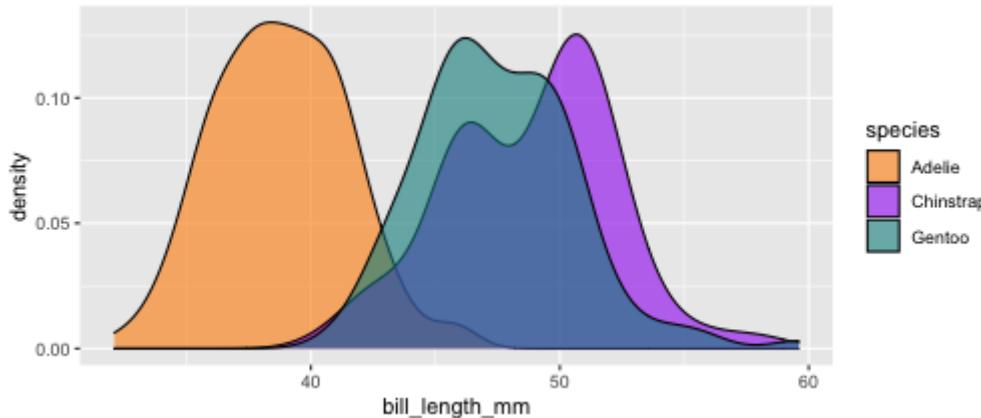


```
g <- ggplot(penguins,  
             aes(bill_length_mm, fill = species)) +  
  geom_density(alpha = 0.6) +  
  scale_fill_manual(  
    breaks = c("Adelie", "Chinstrap", "Gentoo"), # optional but makes it more robust  
    values = c("darkorange", "purple", "cyan4"))  
g
```

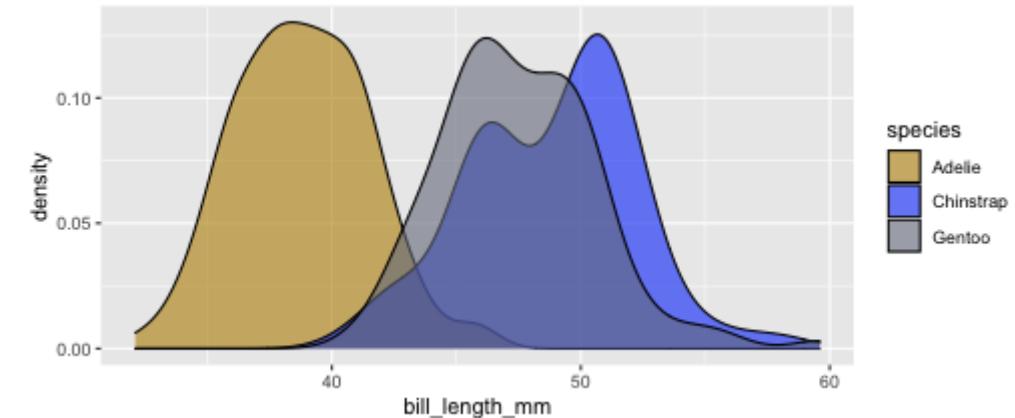
# Check that it's colour blind friendly!

```
cols <- c("darkorange", "purple", "cyan4")
```

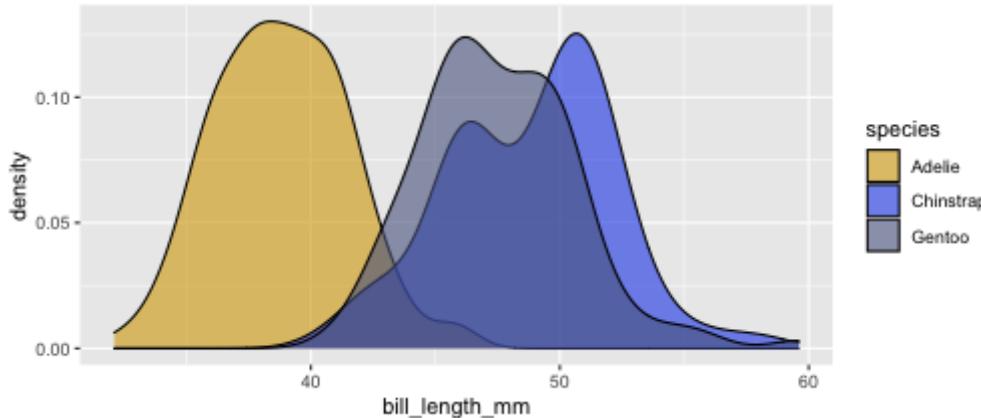
```
g # original
```



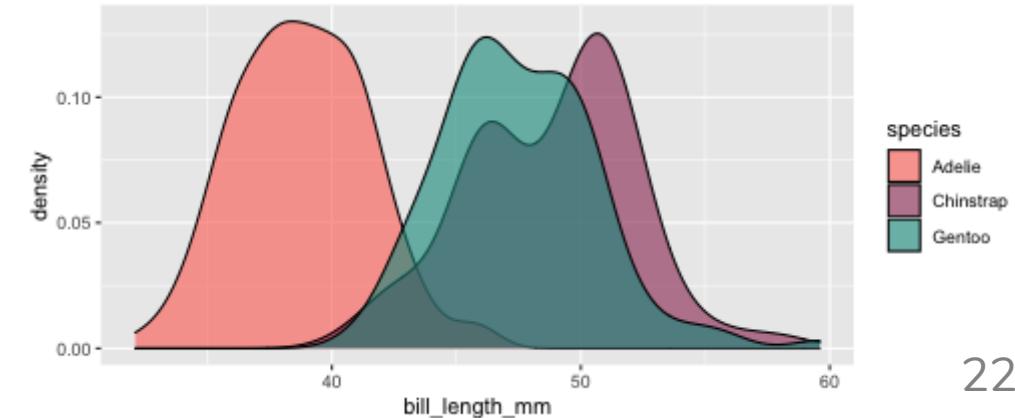
```
g + scale_fill_manual(values = protan(cols))
```



```
g + scale_fill_manual(values = deutan(cols))
```



```
g + scale_fill_manual(values = tritan(cols))
```





</> Open day1-exercise-03.Rmd

15:00

# Session Information

```
devtools::session_info()
```

```
## - Session info --
##   setting  value
##   version  R version 4.0.1 (2020-06-06)
##   os        macOS 10.16
##   system   x86_64, darwin17.0
##   ui        X11
##   language (EN)
##   collate  en_AU.UTF-8
##   ctype    en_AU.UTF-8
##   tz       Australia/Melbourne
##   date     2021-04-07
##
## - Packages --
##   package      * version    date lib

```

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