## Penguins

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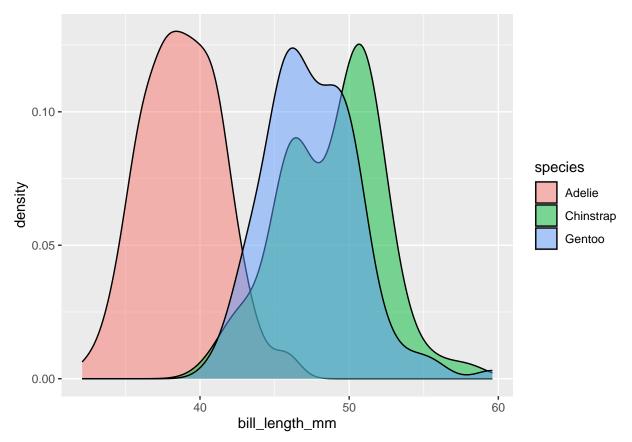
2024-02-09

## Data

- hella penguins yo!
- pic: # Single trait distribution

```
penguins %>%
ggplot() +
geom_density(mapping = aes(x = bill_length_mm, fill=species), alpha=0.5)
```

## Warning: Removed 2 rows containing non-finite values (`stat\_density()`).



# Relationship between multiple traits

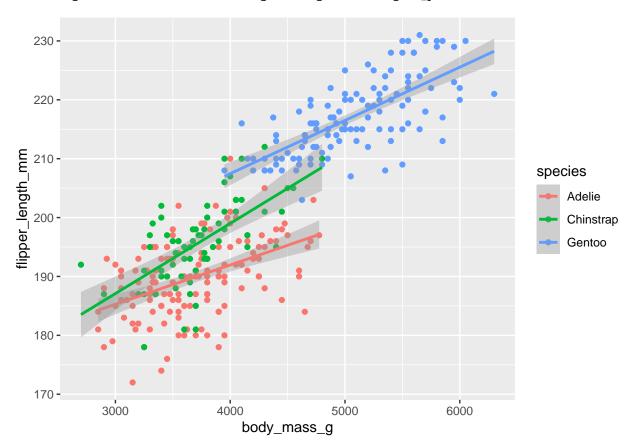
```
penguins %>%
ggplot(mapping = aes(x = body_mass_g, y=flipper_length_mm, color=species)) +
geom_point() +
```

## geom\_smooth(method="lm")

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

## Warning: Removed 2 rows containing non-finite values (`stat\_smooth()`).

## Warning: Removed 2 rows containing missing values (`geom\_point()`).



# Sexual dimorphism

```
penguins %>%
ggplot(mapping = aes(x = bill_length_mm, y=bill_depth_mm, color=sex)) +
geom_point(size=1) +
facet_wrap(~species)
```

## Warning: Removed 2 rows containing missing values (`geom\_point()`).

