DS 116 - Data Visualization

Visualizing Uncertainty

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Visualizing uncertainty

There are many sources of uncertainty

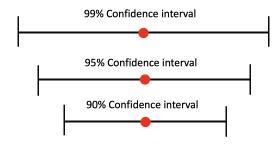
- Uncertainty of point estimates
- Uncertainty of distributions
- Uncertainty of predictions (curve fit) etc..

Section 1

Uncertainty of point estimates

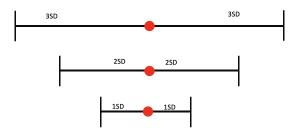
Uncertainty of point estimates

Error bars with standard errors (Confidence intervals)

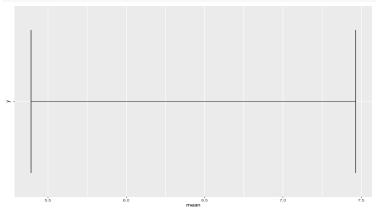


Uncertainty with standard deviations

Error bars with standard deviations



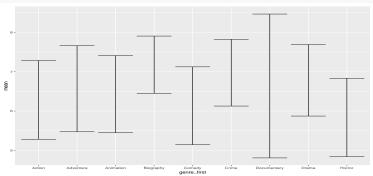
```
ggplot(summarised, mapping = aes(x = "", y = mean, ymax = mean+sd,
                                      ymin = mean-sd)) + geom_errorbar()
 7.0 -
nean
6.5
 6.0 -
 5.5
```



If you want to look at the error bars by groups - summarise by groups using dplyr

```
summarised <- movies_small %>% group_by(genre_first) %>%
 summarise(mean = mean(imdbRating), sd = sd(imdbRating))
summarised
## # A tibble: 9 x 3
  genre_first mean sd
##
## <chr> <dbl> <dbl>
## 1 Action 6.28 1.00
## 2 Adventure 6.56 1.09
## 3 Animation 6.43 0.980
## 4 Biography 7.18 0.728
## 5 Comedy 6.13 0.988
## 6 Crime 6.97 0.848
## 7 Documentary 6.64 1.83
## 8 Drama 6.78 0.908
## 9 Horror 5.84 0.995
```

• use the grouping variable as x aesthetics



There are few functions in R that can be used for generating the data for error bars

```
• +- 1 SD
```

```
smean.sdl(movies_small$imdbRating, mult = 1)
## Mean Lower Upper
## 6.428149 5.391696 7.464602

• +- 2sd

smean.sdl(movies_small$imdbRating, mult = 2)
## Mean Lower Upper
## 6.428149 4.355243 8.501055
```

Hmisc also has functions to construct Confidence Intervals

• 95%

```
smean.cl.normal(movies_small$imdbRating, conf.int = 0.95)
## Mean Lower Upper
## 6.428149 6.384806 6.471493
```

99% confidence interval

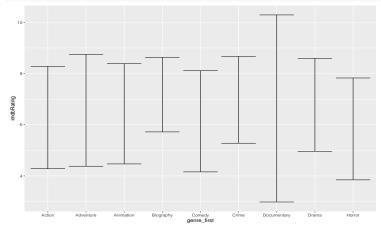
```
smean.cl.normal(movies_small$imdbRating, conf.int = 0.99)
## Mean Lower Upper
## 6.428149 6.371168 6.485130
```

- ggplot2 has a wrapper for Hmisc functions: mean_cl_normal, mean_sdl, mean_se
- To construct the error bars with the functions we will use stat_summary()
 layer

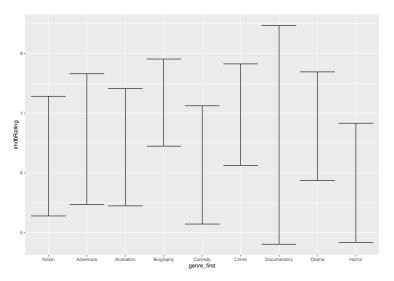
 $stat_summary()$ applies defined function on y by the given values of x. No need to summarise with dplyr and create new dataframe

Error bars for imdbRating by genre (+- 2 SD)

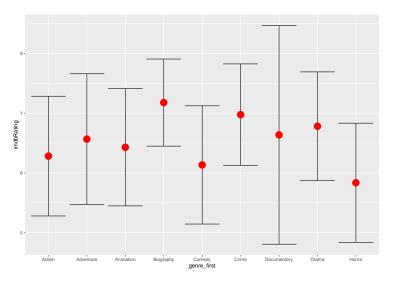
```
ggplot(movies_small, aes(x = genre_first, y = imdbRating)) +
   stat_summary(fun.data = mean_sdl, geom = "errorbar")
```



- By default, mult = 2.
- If you want to change this, use fun.args in stat_summary()



use stat_summary() again to add the mean as a red point to the error bar



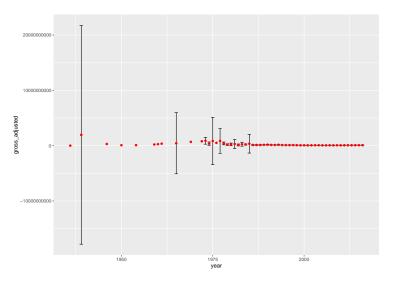
Section 2

Confidence intervals

We can visualize confidence intervals with error bars as well

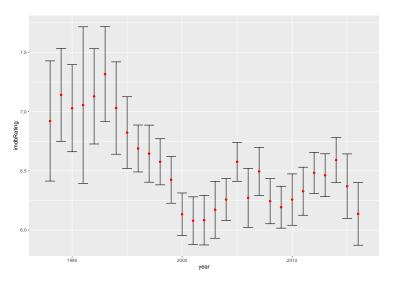
• mean_cl_normal will create 95% CI by default

```
ggplot(movies_small, aes(x = year, y = gross_adjusted)) +
  stat_summary(fun.data = mean_cl_normal, geom = "errorbar") +
  stat_summary(fun = mean, geom='point', color = 'red')
```



Subset the data with the years starting from 1987

```
movies_small %>% filter(year > 1987) %>%
ggplot(aes(x = year, y = imdbRating)) +
   stat_summary(fun.data = mean_cl_normal, geom = "errorbar") +
   stat_summary(fun = mean, geom='point', color = 'red')
```



Error bars with confidence intervals can help us to do initial hypothesis testing

$$H_0: \mu_0 = \mu_1$$

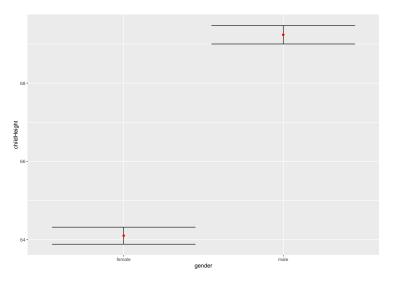
$$H_1: \mu_0 \neq \mu_1$$

We can look at the error bars

- No intersection of the bars is an indicator of rejecting the null hypothesis
- ullet Larger is the gap between error bars, more likely you will reject H_0

Confidence intervals for Height by gender

```
ggplot(GaltonFamilies, aes(x = gender, y = childHeight)) +
   stat_summary(fun.data = mean_cl_normal, geom = "errorbar") +
   stat_summary(fun = mean, geom='point', color = 'red')
```



Testing the hypothesis

```
H_1: \mu_{male} \neq \mu_{female}
t.test(GaltonFamilies$childHeight~GaltonFamilies$gender)
##
##
    Welch Two Sample t-test
##
  data: GaltonFamilies$childHeight by GaltonFamilies$gender
   t = -31.476, df = 929.89, p-value < 0.0000000000000022
  alternative hypothesis: true difference in means is not equal to 0
   95 percent confidence interval:
## -5.449979 -4.810266
  sample estimates:
## mean in group female mean in group male
               64 10397
                                     69.23410
##
```

 $H_0: \mu_{male} = \mu_{female}$

Look at the mtcars data

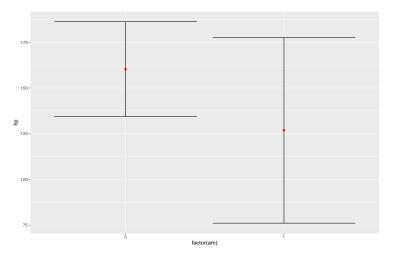
- hp Horsepower
- am transmission automatics vs manual

First - t.test

```
t.test(mtcars$hp~mtcars$am)
##
## Welch Two Sample t-test
##
## data: mtcars$hp by mtcars$am
## t = 1.2662, df = 18.715, p-value = 0.221
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -21.87858 88.71259
## sample estimates:
## mean in group 0 mean in group 1
## 160.2632 126.8462
```

Confidence interval

```
ggplot(mtcars, aes(x = factor(am), y = hp)) +
  stat_summary(fun.data = mean_cl_normal, geom = "errorbar") +
  stat_summary(fun = mean, geom='point', color = 'red')
```



Section 3

Uncertainty in curve fit

When we fit a model into the data, we sometimes look for answers to the following questions:

- What is the mean response for a particular value of x?
- In which interval will these value lie

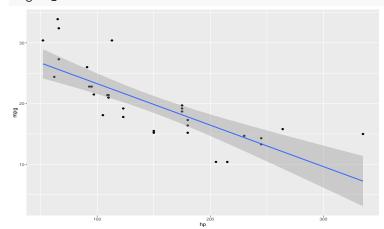
To answer these questions, we can construct confidence interval around the regression line, using the following formula:

$$\hat{y}_h \pm t_{lpha/2,n-2} \sqrt{ extit{MSE}\left(rac{1}{n} + rac{(x_k - ar{x})^2}{\sum (x_i - ar{x}^2)}
ight)}$$

Wider is the confidence interval, more uncertainty we would have for that specific x value.

To get the confidence interval on the graph, specify se=TRUE (the default option) either in stat_smooth or geom_smooth

```
ggplot(mtcars, aes(x = hp, y = mpg)) + geom_point() +
geom_smooth(method = 'lm')
```



As you can see from the formula, further away you go from the mean, larger becomes the confidence interval

```
ggplot(mtcars, aes(x = hp, y = mpg)) + geom_point() +
  geom_smooth(method = 'lm') +
  geom_vline(xintercept = mean(mtcars$hp), color = 'red')
```

