

## Agenda

1. Difference of paired samples
2. Difference of two means

**Warmup: Gifted Children's Parents** Since in this data set, the IQ of both parents is recorded for all children, the IQ data is naturally paired.

```
require(mosaic)
require(openintro)
favstats(~motheriq, data = gifted)

## min      Q1 median      Q3 max      mean      sd  n missing
## 101 113.75    118 122.25 131 118.1667 6.504943 36      0

favstats(~fatheriq, data = gifted)

## min  Q1 median      Q3 max      mean      sd  n missing
## 110 112    115 116.25 126 114.7778 3.48147 36      0
```

We can define a new variable, **diff**, to be the difference between the mother's IQ and the father's for each gifted child.

```
gifted <- gifted %>%
  mutate(diff = motheriq - fatheriq)
favstats(~diff, data = gifted)

## min  Q1 median      Q3 max      mean      sd  n missing
## -15 -2.5    4.5 8.25   18 3.388889 7.453773 36      0
```

Recall the conditions for using a  $t$ -based sampling distribution for a single mean:

1. The samples come from independent observations
2. The distribution of the original variable is approximately normal, or the sample size is large

We return to our original questions:

1. Find a 90% confidence interval for the mean IQ of the mothers. Do the same for the fathers. Do they overlap?
  - (a) State the null and alternative hypotheses
  - (b) Check that **diff** meets the conditions listed above
  - (c) Compute the standard error of the mean ( $SE_{\text{diff}}$ ) and the degrees of freedom
2. Test the hypothesis that the mothers of gifted children have higher IQs, on average, than the fathers. Write out all of the steps. What do you conclude?

- (d) Compute the test statistic ( $T$ )
- (e) Compute the p-value and draw a conclusion [Use the table at the back of the book, or the `pt()` function in R.]
- (f) Write a sentence that provides an interpretation of your result

**Difference of two means** Often the data are *not* naturally paired. In particular, we are often interested in comparing mean from two groups of unequal sizes. For example, the 11 children whose fathers had higher IQs than mothers had a lower average score on the skills test than the 25 children whose mothers had higher IQs than the fathers.

```
favstats(score ~ (diff > 0), data = gifted)
```

```
##   (diff > 0) min    Q1 median  Q3 max    mean      sd  n missing
## 1     FALSE 150 152.5    156 161 164 156.5455 4.906397 11      0
## 2      TRUE 154 159.0    160 163 169 160.2800 4.097967 25      0
```

Now the samples are *not* naturally paired. How do we know if the observed difference in means between these two groups is meaningful? Let  $X$  be the random variable that gives the analytical skills test score for a gifted child whose father has a higher IQ than her mother, and let  $Y$  be the random variable that gives the test score for a gifted child whose mother has a higher IQ. Then we need to understand the sampling distribution of the test statistic  $D = \bar{X} - \bar{Y}$ .

Just as we did with proportions, the standard error of the difference is a combination of the standard errors of the variables.

$$SE_D = \sqrt{(SE_X)^2 + (SE_Y)^2}$$

If both  $X$  and  $Y$  meet the conditions for a  $t$ -based sampling distribution, then  $D$  will meet those conditions as well. We typically use  $\min(n_1 - 1, n_2 - 1)$  for the degrees of freedom.

The hypothesis test for a difference of two means constructed in this manner is called the *two-sample t-test*, and it is a commonly applied statistical technique.

1. Use the information above to conduct a two-sample  $t$ -test for a difference in mean test score between gifted children whose fathers have higher IQs vs. those whose mothers have higher IQs.