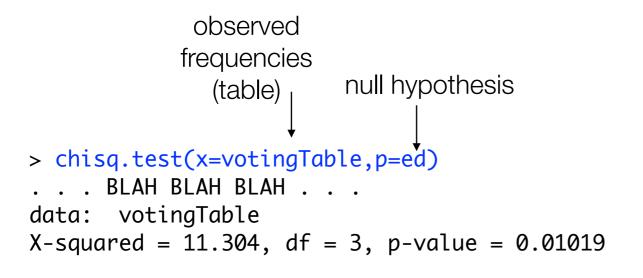
Chi-squared tests are used for categorical data: the outcome variable is nominal scale

Goodness of fit tests compare observed frequencies for *one nominal variable* to a hypothesis about the true probabilities

diagnostic test statistic	X ²	larger value = more evidence against null
distribution	chi-squared: χ^2	describes what you get when you square and sum normally distributed data, like X2
degrees of freedom	k-1 where k is the number of categories	given by # of things - constraints



Report:

- 1. Descriptives (use figure or table if you can)
- 2. Statistical test and null hypothesis
- 3. Stat reference
- 4. Interpretation for your research hypothesis

Stat reference version 1

$$\chi^2 = 11.30$$
, $df = 3$, $p = .0102$

test statistic, degrees of freedom, p-value

Stat reference version 2

$$\chi^2(3) = 11.30, p = .0102$$

sampling distribution(degrees of freedom), test statistic, p-value

Chi-squared test of independence: are two nominal variables are related to one another?

diagnostic test statistic	X ²	larger value = more evidence against null. calculated slightly differently from GOF test
distribution	chi-squared: χ^2	describes what you get when you square and sum normally distributed data, like X2
degrees of freedom	(r-1)(c-1) where r=# of categories of one variable, and c=# of categories of other	given by # of things - constraints

Two assumptions of chi-squared tests

1. "Large" expected frequencies if violated:

Fisher's exact test

> fisher.test(x=bowTable)

2. Independence of the data

if pre-and-post test: McNemar's test

> mcnemar.test(x=medTable)

contingency table

> ct <- chisq.test(x=boxesTable)</pre>

observed frequencies (contingency table) can assign to a variable that contains lots of info

> ct\$stdres

reports adjusted residuals: rule of thumb is +/- 1.96 suggests which items are "significant"

Effect size: Cramer's V for test of independence

based on X² (i.e., divergence from null) and sample size

> library(DescTools) contingency
> CramerV(x=boxesTable) ← table

Cramer's V for Goodness of Fit

based on X² (i.e., divergence from null) and sample size

- > library(rcompanion)
- > cramerVFit(x=votingTable,p=ed)