# Permutation Tests

#### Context

- So far, all of our tests have been in a setting where:
  - we have identified a statistical model we are confident is correct
  - we had a test statistic whose sampling distribution under  $H_0$  we could obtain either
    - \* analytically
    - \* via a large-sample approximation
- What if we are not confident we have a good statistical model, or we can't derive the sampling disribution of our statistic, and our sample size is small?
  - Use computational/sampling approaches to approximate the sampling distribution.
  - Many variations on this idea; here we will discuss permutation tests for paired data.

## Example

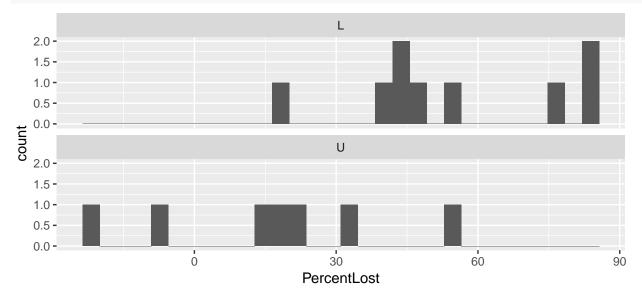
It is sometimes claimed that logging can help forests recover more quickly after forest fires. Is there evidence for this claim?

Here's a quote from the Statistical Sleuth (Ramsey and Schafer, 2013) describing our data:

The 2002 Biscuit Fire in southwest Oregon provided a test case. Researchers selected 16 fire-affected plots in 2014 – before any logging was done – and counted tree seedlings along a randomly located transect pattern in each plot. They returned in 2005, after nine of the plots had been logged, and counted the tree seedlings along the same transects. ( Data from D.C. Donato et al., 2006. "Post-Wildfire Logging Hinders Regeneration and Increases Fire Risk," *Science*, 311: 352.) The numbers of seedlings in the logged (L) and unlogged (U) plots are [loaded in the R code below].

```
logging <- read_csv("http://www.evanlray.com/data/sleuth3/ex0429_logging.csv")

ggplot(data = logging, mapping = aes(x = PercentLost)) +
   geom_histogram() +
   facet_wrap(vars(Action), ncol = 1)</pre>
```



Let  $\mu_1$  = mean percent lost in "population" of plots that are logged.

Let  $\mu_2$  = mean percent lost in "population" of plots that are unlogged.

Test

 $H_0: \mu_1 = \mu_2 \text{ vs } H_A: \mu_1 \neq \mu_2$ 

## Option 1: Likelihood Ratio Test based on parametric model

- If we assume that  $X_{1i} \stackrel{\text{i.i.d.}}{\sim} \text{Normal}(\mu_1, \sigma^2)$  and  $X_{2i} \stackrel{\text{i.i.d.}}{\sim} \text{Normal}(\mu_2, \sigma^2)$ , we can derive a likelihood ratio test based on t distributions.
- If we don't trust that the data are normally distributed, this is risky (we have seen that t-based methods can fail with moderate sample sizes if conditions are not met).

## Option 2: Large-sample $\chi^2$ approximation to sampling distribution of likelihood ratio

• 15 is not large, this is a worse idea than #1

#### Option 3: Permutation test

• Easier to motivate with a slight modification to the hypotheses:

 $H_0$ : The distributions of percent lost are the same whether or not logging is done. (In particular,  $\mu_1 = \mu_2$ )

 $H_A$ : The distributions of percent lost are different depending on whether or not logging is done. (In particular,  $\mu_1 \neq \mu_2$ )

- Our test statistic will be the difference in means:  $W = \bar{X}_1 \bar{X}_2$ .
- To calculate a p-value, we need an estimate of the sampling distribution of W under the condition that  $H_0$  is true.
- Key ideas:
  - If  $H_0$  is true, the observed percent lost for our plots would have been equally likely to be observed in either the logged or unlogged plots.
  - We will simulate many data sets that might have been observed if  $H_0$  was true by permuting assignments of percent lost to different plots.
- 1. Allocate storage space for difference in group means from nsims different samples
- 2. For  $i = 1, \ldots, nsims$ :
  - a. Permute the assignments of observed values to groups
  - b. Calculate the difference in group means, save in allocated space
- 3. Calculate approximate p-value as proportion of simulated samples with a difference in group means at least as large as our observed difference in group means