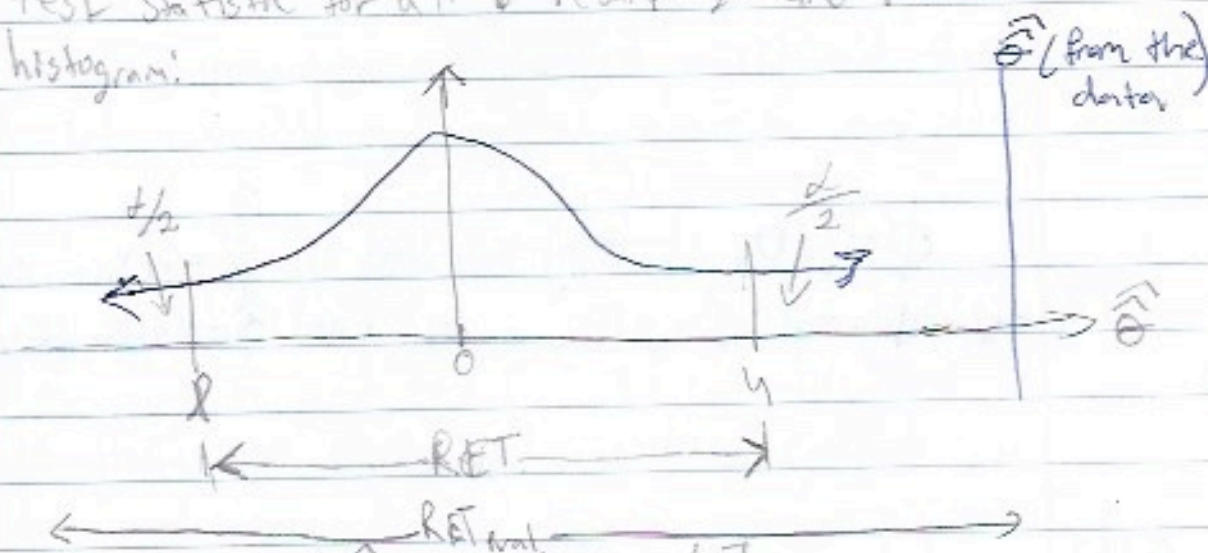


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Let's go with (a) the difference in sample averages. So, under the null, $\bar{X}_{D,1} - \bar{X}_{D,2} \approx 0$. It won't be exactly zero due to chance variations, so what is the threshold of non zero to reject H_0 ? We need the sampling distribution of this difference in sample averages under H_0 .

There are many such fake samples $\binom{200}{100} = 10^{58}$. So w/ a big computer, let's take $B = 1,000,000$. The higher the B is the more accurate, but not much more accurate. Let's calculate the test statistic for all B "resamplings" and plot a histogram.



$$l := \text{Quantile}[\{\hat{\theta}_1, \dots, \hat{\theta}_B\}, \frac{1}{2}]$$

$$u := \text{Quantile}[\{\hat{\theta}_1, \dots, \hat{\theta}_B\}, 1 - \frac{1}{2}]$$

How to get the 1-sided pval. You count the number of $\hat{\theta}_b$'s that are more extreme than the $\hat{\theta}$ from the original data.

You can also use Fisher's permutation idea to get a CI for θ . This is difficult computationally so we won't study it.

We will now see a resampling method that is extremely famous and extremely useful: Efron non-parametric boot strap (1979), ch 8 AoS and the idea is remarkably simple. Imagine you have an iid DGP $f(x; \theta_1, \dots, \theta_k)$ and you have some parameter/function of parameters you want inference for e.g.

$\phi = g(\theta_1, \dots, \theta_k)$ estimated by $\hat{\phi} = W(x_1, \dots, x_n)$
e.g. $\phi = \text{med}[x] = g(\theta_1, \dots, \theta_k)$ but $\hat{\phi} = \text{Median}[\sum x_1, \dots, x_n]$

We don't know to get the estimator's distribution. Which means we can't test hypothesis or construct CIs. But... the bootstrap gives us a way to find the asymptotic distribution of the estimator:

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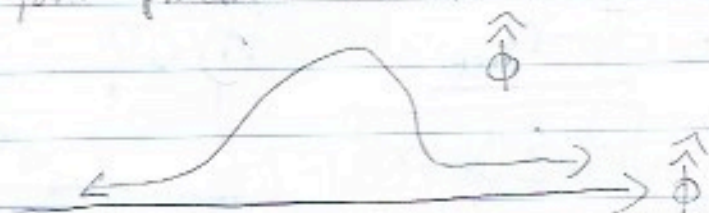
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For $b=1, \dots, B$, a large # of resamplings,
 * take a fake sample of size n by sampling
 w/ replacement of the original n data points
 (this will give you about $2/3$ of the unique data
 points and $1/3$ left out)
 * for each fake sample, I calculate $\hat{\theta}_b$
 The collection of these B statistics is called the
 "bootstrap distribution"

Thm: the bootstrap distribution approximates
 the real distribution of the estimator as $B \rightarrow \infty$
 and $n \rightarrow \infty$. So B should be as big as possible
 given your practical constraints.



One way to get a CI is to just use the
 quantiles of the bootstrap distribution:

$$CI_{\phi, 1-\alpha} = \left[\text{Quantile}[\hat{\phi}_1, \dots, \hat{\phi}_B], \frac{\alpha}{2}, \text{Quantile}[\hat{\phi}_1, \dots, \hat{\phi}_B], 1 - \frac{\alpha}{2} \right]$$

there are other ways too that we won't study. One way to do a two-sided hypothesis test (i.e. against $H_0: \phi = \phi_0$) is to just check

$$\phi_0 \in CI_{\phi, 1-\alpha} \xrightarrow{\text{Yes}} \text{Retain } H_0$$

In finance, people care about the Sharpe Ratio defined as:

$$\phi = \frac{E[X] - r_{\text{free}}}{SD[X]}$$

← a constant representing the risk free rate

$$\hat{\phi} = \frac{\overset{\uparrow \text{(estimate)}}{\bar{X}} - r_{\text{free}}}{\sqrt{\frac{1}{n} \sum (X_i - \bar{X})^2}}$$

$$\hat{\phi} \sim ?$$

Randomization and Causal Estimation

Consider two populations,

Scenario 1:

Pop. 1: Students who hand in their test at the end of the exam

Pop. 2: Students who hand in their test early.

Let $Y_{1,1}, \dots, Y_{1,n-1}$ denote test score data from Pop. 1

Let $Y_{2,1}, \dots, Y_{2,n-2}$ denote test score data from Pop. 2

We want to test if the means of these two populations are different i.e. against $H_0: \theta_1 = \theta_2$.

We know nothing about the DGPs but we can use the Wald test (2-sample z-test):

$$\text{If } \frac{\hat{\theta}_1 - \hat{\theta}_2}{\sqrt{s_1^2/n_1 + s_2^2/n_2}} \in [-1.96, 1.96] \Rightarrow \text{Retain } H_0 \text{ at } \alpha = 5\%$$

Let's say you reject and then you can conclude: "there is a difference in the mean test scores b/w students who hand