

Monte Carlo Tests

Suppose we have a null hypothesis H_0 , represented by a completely specified model, and a test statistic T , such as a goodness-of-fit statistic, for which small values indicate departure from H_0 .

Denote the observed data by y and let the observed value of T be $t = T(y)$. small $T \implies$ rejection arbitrarily chosen.

Monte Carlo Tests

To perform a pure significance test we need to know the distribution of T under H_0 . This may be difficult or impossible to obtain analytically, but it may be possible to simulate from the model to produce m samples y_1^*, \dots, y_m^* and the corresponding values t_1^*, \dots, t_m^* of T . We could then estimate the distribution of T under H_0 by the empirical distribution of (t_1^*, \dots, t_m^*) , or a smoothed version thereof. Informally, we estimate the critical point of a level α test by the 100α th percentile of (t_1^*, \dots, t_m^*) .

Monte Carlo Tests

Monte Carlo tests are a related idea. If H_0 is true, we have $m + 1$ values from the distribution of T , m by simulation and one by observation. Thus the probability that t is the k th smallest or smaller of $\{t, t_1^*, \dots, t_m^*\}$ is $k/(m + 1)$, ignoring ties. [Assume a continuous distribution for T .] If we choose k and m so that $\alpha = k/(m + 1)$ is a conventional significance level (e.g. 0.01 or 0.05), we have a ‘Monte Carlo test’, which rejects H_0 if t is the k th smallest or smaller of the $m + 1$ values. [Two sided versions of the Monte Carlo test follow an obvious construction].

Example

Consider n random points in $[0, 1] \times [0, 1]$. The null hypothesis is one of uniformity, to be tested against one of inhibition between points. Let $T = \#$ 'close' pairs of points. Small values of T reject H_0 . A Monte Carlo test of the significance of t is performed by generating m samples of n points uniformly on $[0, 1] \times [0, 1]$. Notice that in performing the test we can stop when k samples t_i^* are smaller than t , for then we know we will not reject. Also, if $m - k + 1$ values exceed t , we will certainly reject and can stop.

Monte Carlo Tests

Let F be the distribution function of T under H_0 .

A 'conventional test of size α ', possible if F were known, rejects H_0 if

$$u = F(t) = P(T \leq t \mid H_0) < \alpha.$$

The Monte Carlo test rejects H_0 if

$$t < t_{(k)},$$

where $t_{(k)}$ is the k th order statistic of (t_1^*, \dots, t_m^*) .

Monte Carlo Tests

So, the Monte Carlo test rejects H_0 with probability p , where

$$\begin{aligned} p &= P(r \text{ simulated values of } T \text{ are } \leq t, 0 \leq r \leq k-1) \\ &= \sum_{r=0}^{k-1} P(r \text{ simulated values of } T \text{ are } \leq t) \\ &= \sum_{r=0}^{k-1} \binom{m}{r} u^r (1-u)^{m-r}. \quad (u = F(t)) \end{aligned}$$

This should be interpreted as the proportion of times the Monte Carlo test will reject H_0 with $t = F^{-1}(u)$ as observation. Note: we are not assuming $t \sim F$ here, just using F^{-1} as a transformation, as we have $t_i^* \sim F_T$

Monte Carlo Tests

Compare the power function $\beta^{(m)}(\alpha)$ of the Monte Carlo test with the power $\beta(\alpha)$ of the conventional test. We expect, and it is true, that $\beta^{(m)} \leq \beta$, but how large can the power loss be? Power = 1 - P(type II error) = P(reject | H_0 false)

The conventional test rejects H_0 if $T < F^{-1}(\alpha)$ and has power against an alternative denoted by F_θ of

$$\begin{aligned}\beta(\alpha) &= P(T < F^{-1}(\alpha) \mid T \sim F_\theta) \\ &= F_\theta(F^{-1}(\alpha)).\end{aligned}$$

Monte Carlo Tests

Similarly, the power of the Monte Carlo test against this alternative is

$$\begin{aligned}\beta^{(m)}(\alpha) &= P(T < t_{(k)} \mid T \sim F_\theta, t_1^*, \dots, t_m^* \sim F) \\&= \int_{-\infty}^{\infty} \sum_{r=0}^{k-1} \binom{m}{r} F(t)^r [1 - F(t)]^{m-r} dF_\theta(t) \\&= \int_{-\infty}^{\infty} \int_{F(t)}^1 b(\alpha, m, \xi) d\xi dF_\theta(t) \\&= \int_0^1 F_\theta(F^{-1}(\xi)) b(\alpha, m, \xi) d\xi \\&= \int_0^1 \beta(\xi) b(\alpha, m, \xi) d\xi,\end{aligned}$$

where $b(\alpha, m, \cdot)$ is the pdf of a beta distribution with parameters $\alpha(m+1)$ and $(1-\alpha)(m+1)$.

Monte Carlo Tests

If m is big, the Monte Carlo test is OK.

Avoid small values of k , which blur the critical region a lot and lose power. Since $\alpha = k/(m+1)$, this implies m is large. Usually $m = 99$ will be sufficient.

Resampling Techniques: The Jackknife

Suppose we have a statistic θ , with an estimator

$$\hat{\theta} = \hat{\theta}(X_1, \dots, X_n).$$

Construct n distinct $(n - 1)$ -length samples from our original sample X_1, \dots, X_n ; do this by leaving out each datapoint in turn.

$$\underline{X}_{-i} = \underline{X}_{(i)} = (X_1, \dots, X_{i-1}, X_{i+1}, \dots, X_n), i = 1, \dots, n$$

Use each sample $\underline{X}_{(i)}$ to obtain a different estimate $\hat{\theta}_{(i)} = \hat{\theta}(\underline{X}_{(i)})$

Resampling Techniques: The Jackknife

Consider

$$\hat{\theta}_{(\cdot)} = \frac{1}{n} \sum_{i=1}^n \hat{\theta}_{(i)}.$$

with variance estimator

$$\hat{V}_{jack} [\hat{\theta}] = \frac{n-1}{n} \sum_{i=1}^n \left(\hat{\theta}_{(i)} - \hat{\theta}_{(\cdot)} \right)^2.$$

Resampling Techniques: Bootstrap Resampling

Suppose we have an i.i.d. sample from an unknown distribution, with cdf $F(\cdot)$. Given the data $X_1, \dots, X_n \stackrel{i.i.d.}{\sim} F(x)$, construct an empirical estimate \hat{F} of the distribution function

$$\hat{F}(x) = \frac{i}{n} \quad X_{(i)} \leq x < X_{(i+1)}$$

where $X_{(i)}$ is the i^{th} order statistic. Sample B i.i.d. samples from this empirical estimate,

$$X_1^{*(b)}, \dots, X_n^{*(b)} \stackrel{i.i.d.}{\sim} \hat{F}, \quad b = 1, \dots, B.$$

Resampling Techniques: Bootstrap Resampling

We define the **Monte Carlo approximation of the bootstrap estimator** for the variance of

$$\hat{\theta}^{*(b)} = \hat{\theta} \left(X_1^{*(b)}, \dots, X_n^{*(b)} \right), \quad b = 1, \dots, B.$$

to be

$$\widehat{\text{Var}}_{boot}^* \left[\hat{\theta} \right] = \frac{1}{B-1} \sum_{b=1}^B \left(\hat{\theta}^{*(b)} - \frac{1}{B} \sum_{k=1}^B \hat{\theta}^{*(k)} \right)^2.$$

Resampling Techniques: Bootstrap Resampling

- ▶ This methodology extends flexibly to estimating any distributional characteristics of $\hat{\theta}$, not just the variance!
- ▶ number of bootstrap samples is not restricted by the size of the original data sample, unlike with the Jackknife
- ▶ Monte Carlo sampling from $\hat{F} \equiv$ resampling with replacement from the original data!
- ▶ Increasing B will reduce the Monte Carlo error, but not the error due to using \hat{F} instead of F .