R(0,0) = SEB Var[Xi]

$$SE[O] = \sqrt{4r\left(\frac{1}{n}(\chi_1 + ... + \chi_n)\right)} = \sqrt{\frac{1}{nr}} \cdot \sum_{n} \sqrt{nk} \cdot \sum_{n} \sqrt{nk}$$

You have some well-specified mathematical theory about the DGP. For example, in the iphone survey, "I think the proportion of iPhone users in the population is NOT 52.4%. I want to prove my theory to the world (using my sample). Note: it is absolutely impossible to prove or disprove my theory because you cannot see the whole population (or go inside of the DGP). We must use inference which is always a guess.

Two ways to go about "proving" my theory:
(1) I assume I'm right and wait for other people to show me data that contradicts my theory.
(2) I assume my theory is wrong. Then I adduce (bring) evidence (i.e. data) to the contrary until people are convinced my theory

is right.

= 0.9

→20ĝ

#2 is more intellectually honest and more likely to convince.

A "hypothesis" is a mathematical statement about the DGP e.g. theta = 0.9, theta > 0.9, theta is not equal to 0.9, or theta <= or theta is in the set [0.89, 0.91], etc. The "alternative hypothesis" (H_a) is the theory you want to prove. The "null hypothesis" (H_0) is the opposite you assum in #2 for the purpose of contradicting it. Usual cases:

Ho: 0 4 Bo, Ha: 0 > Bo (right-tailed test) Ho: 0 = Bo, Ha: O < O. (left-tailed teat) Ho: 0 = 00, Ha: 0 + 00 (tho-tribl tess) How to perform this test? There are many, many options even for the same DGP. The protocol goes as follows (1) you think of a "test statistic" that could measure the departure away from H_0. (2) Derive the statistical estimator's distribution under H_0. (3) Gauge the departure.

Ha: 0 + ,524,00 Ho: 0=,524-00 (1) My test statistic is ... $\hat{B} = \bar{X}$, \hat{B} to a realization from \hat{B}

We begin with DGP: iid Bern(theta) and the "binomial exact test"

t) 0/H, ~? => 708 Ho = X, +... + Y = ~ Bihom (20, 0, =. 524)

P(B= B1H) 0.27

17.2% 1724. [14.2%. 015 right tail of the sampling distr. left tail

205 0,5 Rejection Rejection Region

Region

Retainment Region

Retain H_0 (fail to reject H_0). Not enough evidence to reject H_0 . Some authors say "accept H_0 ".

Reject H_0 / Accept H_a. My estimate is "statistically significant".

Let's say we rejected H_0 but it was really was true. This is called a Type I error. Where is the P(Type I error) on our plot? $\propto := P(Type I error) = P(\hat{\theta} \notin RET \mid H_0)$

Then in a 2-tailed test, I apportion about alpha/2 to the left tail and about alpha/2 to the right tail.

$$7\sqrt{n_y}$$
 $\sqrt{n_{z}}$

My RET, α = P(β=0|Ho)+...+P(β=0,5|Ho)+ P(β=0,75|Ho)+...+P(β=1|Ho)=7,06%. The choice of alpha is up to you. The scientific community's standard is 5% and sometimes 1%.

If I fail to reject H_0 when H_a is true that's a different error, a "type II error". Failure to prove your theory. The smaller the alpha, the larger the P(Type II error).

If you would like to prove your theory, you have to accept a positive probability of a Type I error.

As of now, we cannot calculate the P(Type II error).