# Bayesian hypothesis testing (cont.)

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#### Outline

- Review of formal Bayesian hypothesis testing
- Likelihood ratio tests
- Jeffrey-Lindley paradox

#### Bayes tests = evaluate predictive models

Consider a standard hypothesis test scenario:

$$H_0: \theta = \theta_0, \qquad H_1: \theta \neq \theta_0$$

A Bayesian measure of the support for the null hypothesis is the Bayes Factor:

$$BF(H_0: H_1) = \frac{p(y|H_0)}{p(y|H_1)} = \frac{p(y|\theta_0)}{\int p(y|\theta)p(\theta|H_1)d\theta}$$

where  $p(\theta|H_1)$  is the prior distribution for  $\theta$  under the alternative hypothesis. Thus the Bayes Factor measures the predictive ability of the two Bayesian models. Both models say  $p(y|\theta)$  are the data model if we know  $\theta$ , but

- 1. Model 0 says  $\theta = \theta_0$  and thus  $p(y|\theta_0)$  is our predictive distribution for y while
- 2. Model 1 says  $p(\theta|H_1)$  is our uncertainty about  $\theta$  and thus

$$p(y|H_1) = \int p(y|\theta)p(\theta|H_1)d\theta$$

is our predictive distribution for y.

#### Normal example

Consider  $y \sim N(\theta, 1)$  and

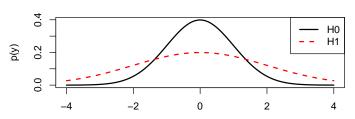
$$H_0: \theta = 0, \qquad H_1: \theta \neq 0$$

and we assume  $\theta|H_1 \sim N(0, C)$ . Thus,

$$BF(H_0: H_1) = \frac{p(y|H_0)}{p(y|H_1)} = \frac{p(y|\theta_0)}{\int p(y|\theta)p(\theta|H_1)d\theta} = \frac{N(y; 0, 1)}{N(y; 0, 1 + C)}.$$

Now, as  $C \to \infty$ , our predictions about y become less sharp.

#### **Predictive distributions**



#### Likelihood Ratio Tests

Consider a likelihood  $L(\theta) = p(y|\theta)$ , then the liklihood ratio test statistic for testing  $H_0: \theta \in \Theta_0$  and  $H_1: \theta \in \Theta_0^c$  with  $\Theta = \Theta_0 \cup \Theta_0^c$  is

$$\lambda(y) = \frac{\sup_{\Theta_0} L(\theta)}{\sup_{\Theta} L(\theta)} = \frac{L(\hat{\theta}_{0,MLE})}{L(\hat{\theta}_{MLE})}$$

where  $\hat{\theta}_{MLE}$  and  $\hat{\theta}_{0,MLE}$  are the (restricted) MLEs. The likelihood ratio test (LRT) is any test that has a rejection region of the form  $\{y:\lambda(y)\leq c\}$ . (Casella & Berger Def 8.2.1)

Under certain conditions (see Casella & Berger 10.3.3), as  $n \to \infty$ 

$$-2\log\lambda(y)\to\chi^2_{\nu}$$

where  $\nu$  us the difference between the number of free parameters specified by  $\theta \in \theta_0$  and the number of free parameters specified by  $\theta \in \Theta$ .

# Binomial example

Consider a coin flipping experiment so that  $Y_i \stackrel{iid}{\sim} Ber(\theta)$  and the null hypothesis  $H_0: \theta = 0.5$  versus the alternative  $H_1: \theta \neq 0.5$ . Then

$$\lambda(y) = \frac{\sup_{\Theta_0} L(\theta)}{\sup_{\Theta} L(\theta)} = \frac{0.5^n}{\hat{\theta}_{MLE}^{n\overline{y}} (1 - \hat{\theta}_{MLE})^{n - n\overline{y}}} = \frac{0.5^n}{\overline{y}^{n\overline{y}} (1 - \overline{y})^{n - n\overline{y}}}$$

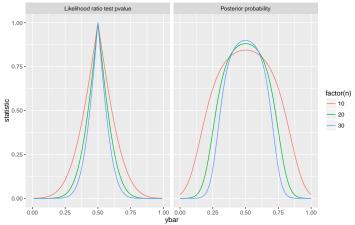
and  $-2\log\lambda(y) \to \chi_1^2$  as  $n\to\infty$  so

pvalue 
$$\approx P(\chi_1^2 > -2 \log \lambda(y)).$$

If pvalue  $< \alpha$ , then we reject  $H_0$  at level  $\alpha$ . Typically  $\alpha = 0.05$ .

# Binomial example

 $Y \sim Bin(n, \theta)$  and, for the Bayesian analysis,  $\theta|H_1 \sim Be(1, 1)$  and  $p(H_0) = p(H_1) = 0.5$ :



### Do pvalues and posterior probabilities agree?

Suppose n = 10,000 and y = 4,900, then the pvalue is

$$pvalue \approx P(\chi_1^2 > -2\log(0.135)) = 0.045$$

so we would reject  $H_0$  at the 0.05 level.

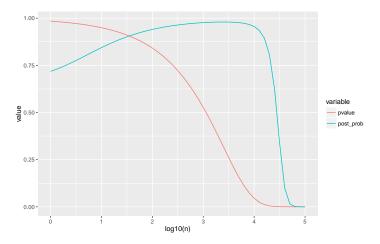
The posterior probability of  $H_0$  is

$$p(H_0|y) \approx \frac{1}{1+1/10.8} = 0.96,$$

so the probability of  $H_0$  being true is 96%.

It appears the Bayesian and LRT pvalue completely disagree!

### Binomial $\overline{y} = 0.49$ with $n \to \infty$



### Jeffrey-Lindley Paradox

#### Definition

The Jeffrey-Lindley Paradox concerns a situation when comparing two hypotheses  $H_0$  and  $H_1$  given data y and find

- ullet a frequentist test result is significant leading to rejection of  $H_0$ , but
- the posterior probability of  $H_0$  is high.

#### This can happen when

- the effect size is small,
- n is large,
- H<sub>0</sub> is relatively precise,
- H<sub>1</sub> is relative diffuse, and
- the prior model odds is  $\approx 1$ .

### Comparison

The test statistic with point null hypotheses:

$$\lambda(y) = \frac{p(y|\theta_0)}{p(y|\hat{\theta}_{MLE})}$$

$$BF(H_0: H_1) = \frac{p(y|\theta_0)}{\int p(y|\theta)p(\theta|H_1)d\theta} = \frac{p(y|H_0)}{p(y|H_1)}$$

#### A few comments:

- The LRT chooses the best possible alternative value.
- The Bayesian test penalizes for vagueness in the prior.
- The LRT can be interpreted as a Bayesian point mass prior exactly at the MLE.
- Generally, pvalues provide a measure of lack-of-fit of the null model.
- Bayesian tests compare predictive performance of two Bayesian models (model+prior).

### Normal mean testing

Let  $y \sim N(\theta, 1)$  and we are testing

$$H_0: \theta = 0$$
 vs  $H_1: \theta \neq 0$ 

We can compute a two-sided pvalue via

$$\mathsf{pvalue} = 2\Phi(-|y|)$$

where  $\Phi(\cdot)$  is the cumulative distribution function for a standard normal.

Typically, we set our Type I error rate at level  $\alpha$ , i.e.

$$P(\text{reject } H_0|H_0 \text{ true}) = \alpha.$$

But, if the pvalue is less than  $\alpha$ , we should be interested in

$$P(H_0 \text{ true}|\text{reject } H_0).$$

### Pvalue interpretation

Let  $y \sim N(\theta, 1)$  and we are testing

$$H_0: \theta = 0$$
 vs  $H_1: \theta \neq 0$ 

For the following activity, you need to tell me

- 1. the observed pvalue,
- 2. the relative frequencies of null and alternative hypotheses, and
- 3. the distribution for  $\theta$  under the alternative.

Then this pvalue app below will calculate (via simulation) the probability the null hypothesis is true.

shiny::runGitHub('jarad/pvalue')

# Pvalue app approach

The idea is that a scientist performs a series of experiments. For each experiment,

- whether  $H_0$  or  $H_1$  is true is randomly determined,
- $oldsymbol{ heta}$  is sampled according to which hypothesis is true, and
- the pvalue is calculated.

This process is repeated until a pvalue of the desired value is achieved, e.g. pvalue=0.05, and the true hypothesis is recorded. Thus,

$$P(H_0 \text{ true} \mid \text{pvalue} = 0.05) \approx \frac{1}{K} \sum_{k=1}^{K} I(H_0 \text{ true} \mid \text{pvalue} \approx 0.05).$$

Thus, there is nothing Bayesian happening here except that the probability being calculated has the unknown quantity on the left and the known quantity on the right.

### Prosecutor's Fallacy

It is common for those using statistics to equate the following

pvalue 
$$\stackrel{?}{=} P(\text{data}|H_0 \text{ true}) \neq P(H_0 \text{ true}|\text{data}).$$

but we can use Bayes rule to show us that these probabilities cannot be equated

$$p(H_0|y) = \frac{p(y|H_0)p(H_0)}{p(y)} = \frac{p(y|H_0)p(H_0)}{p(y|H_0)p(H_0) + p(y|H_0)p(H_0)}$$

This situation is common enough that it is called The Prosecutor's Fallacy.