

# Bayesian inference

Statistical Inference

Brian Caffo, Roger Peng, Jeff Leek Johns Hopkins Bloomberg School of Public Health

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# **Bayesian analysis**

- · Bayesian statistics posits a *prior* on the parameter of interest
- · All inferences are then performed on the distribution of the parameter given the data, called the posterior
- In general, \mbox{Posterior} \propto \mbox{Likelihood} \times \mbox{Prior}
- Therefore (as we saw in diagnostic testing) the likelihood is the factor by which our prior beliefs are updated to produce conclusions in the light of the data

## **Prior specification**

- The beta distribution is the default prior for parameters between 0 and 1.
- The beta density depends on two parameters \alpha and \beta \frac{\Gamma(\alpha + \beta)} {\Gamma(\alpha)\Gamma(\beta)} p ^ {\alpha 1} (1 p) ^ {\beta 1} ~~~ \mbox{for} ~~ 0 \leq p \leq 1
- The mean of the beta density is \alpha / (\alpha + \beta)
- The variance of the beta density is \frac{\alpha \beta}{(\alpha + \beta)^2 (\alpha + \beta + 1)}
- The uniform density is the special case where \alpha = \beta = 1

```
## Exploring the beta density
library(manipulate)
pvals <- seq(0.01, 0.99, length = 1000)
manipulate(
    plot(pvals, dbeta(pvals, alpha, beta), type = "1", lwd = 3, frame = FALSE),
    alpha = slider(0.01, 10, initial = 1, step = .5),
    beta = slider(0.01, 10, initial = 1, step = .5)
)</pre>
```

#### **Posterior**

- Suppose that we chose values of \alpha and \beta so that the beta prior is indicative of our degree of belief regarding p in the absence of data
- Then using the rule that \mbox{Posterior} \propto \mbox{Likelihood} \times \mbox{Prior} and throwing out anything that doesn't depend on p, we have that \begin{align} \mbox{Posterior} & \propto p^x(1 p)^{n-x} \times p^{\alpha-1} (1 p)^{\theta-1} \ = p^{x + \alpha-1} (1 p)^{n x} + \beta = 1 \end{align}
- This density is just another beta density with parameters \tilde \alpha = x + \alpha and \tilde \beta = n x + \beta

#### **Posterior mean**

## **Thoughts**

- The posterior mean is a mixture of the MLE (\hat p) and the prior mean
- · \pi goes to 1 as n gets large; for large n the data swamps the prior
- · For small n, the prior mean dominates
- Generalizes how science should ideally work; as data becomes increasingly available, prior beliefs should matter less and less
- · With a prior that is degenerate at a value, no amount of data can overcome the prior

### **Example**

- · Suppose that in a random sample of an at-risk population 13 of 20 subjects had hypertension. Estimate the prevalence of hypertension in this population.
- x = 13 and n = 20
- Consider a uniform prior, \alpha = \beta = 1
- The posterior is proportional to (see formula above)  $p^{x + \alpha 1} (1 p)^{n x + \beta 1} = p^x (1 p)^{n-x}$  That is, for the uniform prior, the posterior is the likelihood
- Consider the instance where  $\alpha = 2$  (recall this prior is humped around the point .5) the posterior is  $p^x + \alpha 1$  (1 p) $\alpha x + \beta = p^x + 1$  (1 p) $\alpha x + 1$
- The "Jeffrey's prior" which has some theoretical benefits puts \alpha = \beta = .5

```
pvals < seq(0.01, 0.99, length = 1000)
x < -13: n < -20
myPlot <- function(alpha, beta){</pre>
    plot(0:1,0:1, type = "n", xlab = "p", ylab = "", frame = FALSE)
    lines(pvals, dbeta(pvals, alpha, beta) / max(dbeta(pvals, alpha, beta)),
            lwd = 3, col = "darkred")
    lines(pvals, dbinom(x,n,pvals) / dbinom(x,n,x/n), lwd = 3, col = "darkblue")
    lines(pvals, dbeta(pvals, alpha+x, beta+(n-x)) / max(dbeta(pvals, alpha+x, beta+(n-x))),
        lwd = 3, col = "darkgreen")
    title("red=prior, green=posterior, blue=likelihood")
manipulate(
   myPlot(alpha, beta),
    alpha = slider(0.01, 100, initial = 1, step = .5),
    beta = slider(0.01, 100, initial = 1, step = .5)
```

#### **Credible intervals**

- · A Bayesian credible interval is the Bayesian analog of a confidence interval
- A 95\% credible interval, [a, b] would satisfy  $P(p \in [a, b] \sim x) = .95$
- The best credible intervals chop off the posterior with a horizontal line in the same way we did for likelihoods
- These are called highest posterior density (HPD) intervals

# Getting HPD intervals for this example

Install the \texttt{binom} package, then the command

```
## Error: there is no package called 'binom'

binom.bayes(13, 20, type = "highest")

## Error: could not find function "binom.bayes"
```

gives the HPD interval.

• The default credible level is 95\% and the default prior is the Jeffrey's prior.

```
pvals < seq(0.01, 0.99, length = 1000)
x < -13; n < -20
mvPlot2 <- function(alpha, beta, cl){</pre>
    plot(pvals, dbeta(pvals, alpha+x, beta+(n-x)), type = "1", lwd = 3,
   xlab = "p", ylab = "", frame = FALSE)
    out <- binom.bayes(x, n, type = "highest",
        prior.shape1 = alpha,
        prior.shape2 = beta,
        conf.level = cl)
    p1 <- out$lower; p2 <- out$upper
    lines(c(p1, p1, p2, p2), c(0, dbeta(c(p1, p2), alpha+x, beta+(n-x)), 0),
        type = "1", lwd = 3, col = "darkred")
manipulate(
   myPlot2(alpha, beta, cl),
    alpha = slider(0.01, 10, initial = 1, step = .5),
   beta = slider(0.01, 10, initial = 1, step = .5),
    cl = slider(0.01, 0.99, initial = 0.95, step = .01)
```