- ► The **Bayes Factor** provides a way to formally compare two **competing models**, say  $M_1$  and  $M_2$ .
- It is similar to testing a "full model" vs. "reduced model" (with, e.g., a likelihood ratio test) in classical statistics.
- However, with the Bayes Factor, one model does not have to be nested within the other.
- Given a data set x, we compare models

$$M_1: f_1(\mathbf{x}|\boldsymbol{\theta}_1) \text{ and } M_2: f_2(\mathbf{x}|\boldsymbol{\theta}_2)$$

▶ We may specify prior distributions  $p_1(\theta_1)$  and  $p_2(\theta_2)$  that lead to prior probabilities for each model  $p(M_1)$  and  $p(M_2)$ .

By Bayes' Law, the **posterior odds** in favor of Model 1 versus Model 2 is:

$$\begin{split} \frac{\pi(M_1|\mathbf{x})}{\pi(M_2|\mathbf{x})} &= \frac{\int_{\Theta_1} \frac{p(M_1) f_1(\mathbf{x}|\theta_1) p_1(\theta_1) \, \mathrm{d}\theta_1}{p(\mathbf{x})}}{\int_{\Theta_2} \frac{p(M_2) f_2(\mathbf{x}|\theta_2) p_2(\theta_2) \, \mathrm{d}\theta_2}{p(\mathbf{x})}} \\ &= \frac{p(M_1)}{p(M_2)} \cdot \frac{\int_{\Theta_1} f_1(\mathbf{x}|\theta_1) p_1(\theta_1) \, \mathrm{d}\theta_1}{\int_{\Theta_2} f_2(\mathbf{x}|\theta_2) p_2(\theta_2) \, \mathrm{d}\theta_2} \\ &= [\text{prior odds}] \times [\text{Bayes Factor } B(\mathbf{x})] \end{split}$$

Rearranging, the Bayes Factor is:

$$B(\mathbf{x}) = \frac{\pi(M_1|\mathbf{x})}{\pi(M_2|\mathbf{x})} \times \frac{p(M_2)}{p(M_1)}$$
$$= \frac{\pi(M_1|\mathbf{x})/\pi(M_2|\mathbf{x})}{p(M_1)/p(M_2)}$$

(the ratio of the posterior odds for  $M_1$  to the prior odds for  $M_1$ ).

- ▶ **Note**: If the prior model probabilities are equal, i.e.,  $p(M_1) = p(M_2)$ , then the Bayes Factor equals the posterior odds for  $M_1$ .
- ▶ Note: If  $p(M_1) = p(M_2)$  and the parameter spaces  $\Theta_1$  and  $\Theta_2$  are the same, then the Bayes Factor reduces to a likelihood ratio.

Note that:

$$B(\mathbf{x}) = \frac{\pi(M_1|\mathbf{x})}{\pi(M_2|\mathbf{x})} \times \frac{p(M_2)}{p(M_1)} = \frac{\frac{\pi(M_1,\mathbf{x})}{p(\mathbf{x})p(M_1)}}{\frac{\pi(M_2,\mathbf{x})}{p(\mathbf{x})p(M_2)}}$$
$$= \frac{\frac{\pi(M_1,\mathbf{x})}{p(M_1)}}{\frac{\pi(M_2,\mathbf{x})}{p(M_2)}} = \frac{\pi(\mathbf{x}|M_1)}{\pi(\mathbf{x}|M_2)}$$

- Clearly a Bayes Factor much greater than 1 supports Model 1 over Model 2.
- ▶ Jeffreys proposed the following rules, if Model 1 represents a null model:

#### Result Conclusion

$$B(\mathbf{x}) \geq 1 o Model \ 1$$
 supported  $0.316 \leq B(\mathbf{x}) < 1 o Minimal evidence against Model 1$  (Note  $0.316 = 10^{-1/2}$ )  $0.1 \leq B(\mathbf{x}) < 0.316 o Substantial evidence against Model 1  $0.01 \leq B(\mathbf{x}) < 0.1 o Strong evidence against Model 1  $B(\mathbf{x}) < 0.01 o Decisive evidence against Model 1$$$ 

Clearly these labels are fairly arbitrary.

In the case when there are only **two possible models**,  $M_1$  and  $M_2$ , then given the Bayes Factor  $B(\mathbf{x})$ , we can calculate the posterior probability of Model 1 as:

$$P(M_1|\mathbf{x}) = 1 - P(M_2|\mathbf{x}) = 1 - \frac{P(\mathbf{x}|M_2)P(M_2)}{P(\mathbf{x})}$$

$$= 1 - \frac{P(\mathbf{x}|M_1)}{B(\mathbf{x})} \frac{P(M_2)}{P(\mathbf{x})}$$

$$\Rightarrow P(M_1|\mathbf{x}) = 1 - \left\{ \frac{1}{B(\mathbf{x})} \frac{P(M_2)}{P(M_1)} \right\} P(M_1|\mathbf{x})$$

$$\Rightarrow 1 = \left[ 1 + \left\{ \frac{1}{B(\mathbf{x})} \frac{P(M_2)}{P(M_1)} \right\} \right] P(M_1|\mathbf{x})$$

$$\Rightarrow P(M_1|\mathbf{x}) = \frac{1}{1 + \left\{ \frac{1}{B(\mathbf{x})} \frac{P(M_2)}{P(M_1)} \right\}}$$

**Example 2(a)**: Comparing Two Means (Bayes Factor Approach)

- ▶ Data: Blood pressure reduction was measured for 11 patients who took calcium supplements and for 10 patients who took a placebo.
- We model the data with normal distributions having common variance:

Calcium data : 
$$X_{1j} \stackrel{\text{iid}}{\sim} N(\mu_1, \sigma^2), \quad j = 1, \dots, 11$$
  
Placebo data :  $X_{2j} \stackrel{\text{iid}}{\sim} N(\mu_2, \sigma^2), \quad j = 1, \dots, 10$ 

Consider the two-sided test for whether the mean BP reduction differs for the two groups:

$$H_0: \mu_1 = \mu_2 \text{ vs. } H_a: \mu_1 \neq \mu_2$$

We will place a prior on the difference of standardized means

$$\Delta = \frac{\mu_1 - \mu_2}{\sigma}$$

with specified prior mean  $\mu_{\Delta}$  and variance  $\sigma_{\Delta}^2$ .

Consider the classical two-sample t-statistic

$$T = \frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2}{n_1 + n_2 - 2}} / \sqrt{n^*}},$$

where 
$$n^* = \left(\frac{1}{n_1} + \frac{1}{n_2}\right)^{-1}$$
.

- $\blacktriangleright$   $H_0$  and  $H_a$  define two specific models for the distribution of T.
- ▶ Under  $H_0$ ,  $T \sim$  (central) t with  $(n_1 + n_2 2)$  degrees of freedom.
- ▶ Under  $H_a$ ,  $T \sim$  noncentral t.
- ▶ With this prior, the Bayes Factor for  $H_0$  over  $H_a$  is:

$$B(\mathbf{x}) = \frac{t_{n_1 + n_2 - 2}(t^*, 0, 1)}{t_{n_1 + n_2 - 2}(t^*, \mu_{\Delta}\sqrt{n^*, 1 + n^*\sigma_{\Delta}^2})}$$

where  $t^*$  is the observed t-statistic.

▶ See R example to get  $B(\mathbf{x})$  and  $P[H_0|\mathbf{x}]$ .

**Example 2(a)**: Comparing Two Means (Gibbs Sampling Approach)

Same data set, but suppose our interest is in testing whether the calcium yields a **better** BP reduction than the placebo:

$$H_0: \mu_1 \leq \mu_2 \text{ vs. } H_a: \mu_1 > \mu_2$$

We set up the sampling model:

$$X_{1j} = \mu + \tau + \epsilon_{1j}, j = 1, \dots, 11$$
  
 $X_{2j} = \mu - \tau + \epsilon_{2j}, j = 1, \dots, 10$ 

where  $\epsilon_{ij} \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$ .

▶ Thus  $\mu_1 = \mu + \tau$  and  $\mu_2 = \mu - \tau$ .

We can assume independent priors for  $\mu$ ,  $\tau$ , and  $\sigma^2$ :

$$\mu \sim \textit{N}(\mu_{\mu}, \sigma_{\mu}^2)$$
 $au \sim \textit{N}(\mu_{ au}, \sigma_{ au}^2)$ 
 $\sigma^2 \sim \textit{IG}(\nu_1/2, \nu_1 \nu_2/2)$ 

Then it can be shown that the full conditional distributions are:

$$\begin{split} & \mu|\mathbf{x}_1,\mathbf{x}_2,\tau,\sigma^2 \sim \mathsf{Normal} \\ & \tau|\mathbf{x}_1,\mathbf{x}_2,\mu,\sigma^2 \sim \mathsf{Normal} \\ & \sigma^2|\mathbf{x}_1,\mathbf{x}_2,\mu,\tau \sim \mathit{IG} \end{split}$$

where the appropriate parameters are given in the R code.

- R example: Gibbs Sampler can obtain approximate posterior distributions for μ and (especially of interest) for τ.
- Note  $P[\mu_1 > \mu_2 | \mathbf{x}] = P[\tau > 0 | \mathbf{x}].$
- ▶ We can also find the **posterior predictive** probability  $P[X_1 > X_2]$ .