#### **Linear Regression**

The classical linear model is

$$Y = X\beta + \epsilon$$

where  ${m Y}$  is  $n \times {m 1}$ ,  ${m X}$  is a fixed  $n \times p$  matrix,  ${m \beta}$  is  $p \times {m 1}$  and

$$\epsilon \sim N(0, \sigma^2 I).$$

We observe only Y and X, and the unknown parameters are the *regression coefficients*  $\beta$  and the error variance  $\sigma^2$ .

We assume that X is of full column rank p.

Let's review frequentist estimation of  $\beta$  and  $\sigma^2$ .

The maximum likelihood estimate (MLE) of  $\beta$  is the *least squares estimate*, which is

$$\widehat{\boldsymbol{\beta}} = (\boldsymbol{X}^T \boldsymbol{X})^{-1} \boldsymbol{X}^T \boldsymbol{Y}.$$

Of course,  $\widehat{\boldsymbol{\beta}}$  is the solution to the problem

$$\min_{oldsymbol{b}} (oldsymbol{Y} - oldsymbol{X} oldsymbol{b})^T (oldsymbol{Y} - oldsymbol{X} oldsymbol{b}).$$

The MLE of  $\sigma^2$  is

$$\hat{\sigma}^{2} = n^{-1} (\mathbf{Y} - \mathbf{X}\hat{\boldsymbol{\beta}})^{T} (\mathbf{Y} - \mathbf{X}\hat{\boldsymbol{\beta}})$$

$$= n^{-1} \mathbf{Y}^{T} \left[ \mathbf{I} - \mathbf{X} (\mathbf{X}^{T} \mathbf{X})^{-1} \mathbf{X}^{T} \right] \mathbf{Y}.$$

$$= n^{-1} \mathbf{Y}^{T} (\mathbf{I} - \mathbf{M}) \mathbf{Y}.$$

The Jeffreys noninformative prior for  $\beta$  is constant over all p-vectors, and hence improper.

A common noninformative prior for  $\sigma$  is  $p(\sigma) = \sigma^{-1}$ , which is also improper.

Assuming that  $\beta$  and  $\sigma$  are a priori independent, and using these noninformative priors, let's derive the posterior.

We have

$$p(\boldsymbol{\beta}, \sigma | \boldsymbol{y}) \propto \frac{1}{\sigma^{n+1}} \exp \left[ -\frac{(\boldsymbol{y} - \boldsymbol{X}\boldsymbol{\beta})^T (\boldsymbol{y} - \boldsymbol{X}\boldsymbol{\beta})}{2\sigma^2} \right].$$

Simple algebra shows that

$$(y - X\beta)^T (y - X\beta) =$$

$$y^T(I-M)y + (\beta - \hat{\beta})^T X^T X (\beta - \hat{\beta}).$$

The posterior is thus proportional to

$$\sigma^{-(n+1)} \exp \left(-\frac{\boldsymbol{y}^T (\boldsymbol{I} - \boldsymbol{M}) \boldsymbol{y}}{2\sigma^2}\right)$$

$$imes \exp\left[-rac{(oldsymbol{eta}-\widehat{oldsymbol{eta}})^Toldsymbol{X}^Toldsymbol{X}(oldsymbol{eta}-\widehat{oldsymbol{eta}})}{2\sigma^2}
ight].$$

It now follows that the conditional of  $\beta$  given  $\sigma$  and y is  $N(\hat{\beta}, \sigma^2(X^TX)^{-1})$ .

The posterior is proportional to

$$\sigma^{-(n+1)}\sigma^p \exp\left(-\frac{\boldsymbol{y}^T(\boldsymbol{I}-\boldsymbol{M})\boldsymbol{y}}{2\sigma^2}\right)$$

$$\times \frac{1}{\sigma^p} \exp \left[ -\frac{(\boldsymbol{\beta} - \widehat{\boldsymbol{\beta}})^T \boldsymbol{X}^T \boldsymbol{X} (\boldsymbol{\beta} - \widehat{\boldsymbol{\beta}})}{2\sigma^2} \right].$$

Now using the fact that

$$p(\boldsymbol{\beta}, \sigma | \boldsymbol{y}) = p(\sigma | \boldsymbol{y}) p(\boldsymbol{\beta} | \sigma, \boldsymbol{y}),$$

it follows that  $\sigma^2$  given y is inverse-gamma with parameters (n-p)/2 and  $y^T(I-M)y/2$ .

The marginal posterior of  $\beta$  is proportional to

$$\left[y^{T}(I-M)y + (\beta - \widehat{\beta})^{T}X^{T}X(\beta - \widehat{\beta})\right]^{-n/2} =$$

$$\left[(n-p)s^{2} + (\beta - \widehat{\beta})^{T}X^{T}X(\beta - \widehat{\beta})\right]^{-n/2} \propto$$

$$\left[1 + \frac{1}{(n-p)}(\beta - \widehat{\beta})^T \left(\frac{X^T X}{s^2}\right) (\beta - \widehat{\beta})\right]^{-n/2}.$$

The p-variate t-distribution with parameters  $\nu$ ,  $\mu$  and  $\Sigma$  has density proportional to

$$|\Sigma|^{-1/2} \left[ 1 + \frac{1}{\nu} (x - \mu)^T \Sigma^{-1} (x - \mu) \right]^{-(\nu + p)/2}.$$

It follows that  $\beta | y$  has a multivariate t-distribution with parameters n - p,  $\hat{\beta}$  and  $s^2(X^TX)^{-1}$ .

We found the full conditional for  $\beta$  on p. 214N.

From the posterior on p. 215 we see that the full conditional of  $1/\sigma^2$  is gamma with parameters n/2 and

$$\frac{1}{2}\left[(n-p)s^2+(\beta-\widehat{\beta})^TX^TX(\beta-\widehat{\beta})\right].$$

#### Example 16 Maximal oxygen uptake

The data in this example were simulated from a model reported in a 1994 article in *Research Quarterly for Exercise and Sport*.

The study in that article involved 50 male runners. A kinesiologist wanted to know if maximal oxygen uptake could be predicted from easily measured explanatory variables.

The variables in the study were as follows:

```
y = \text{maximal oxygen uptake (in liters per minute)}
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```
x_1 = \text{weight (in kilograms)}
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$$x_2 = age (in years)$$

 $x_3 = \text{time to walk 1 mile (in minutes)}$ 

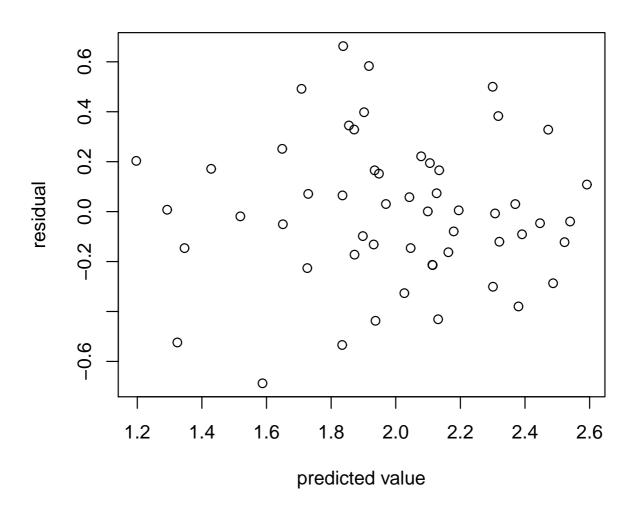
 $x_4$  = heart rate at end of walk (in beats per minute)

Least squares estimates of  $\beta_j$ s:

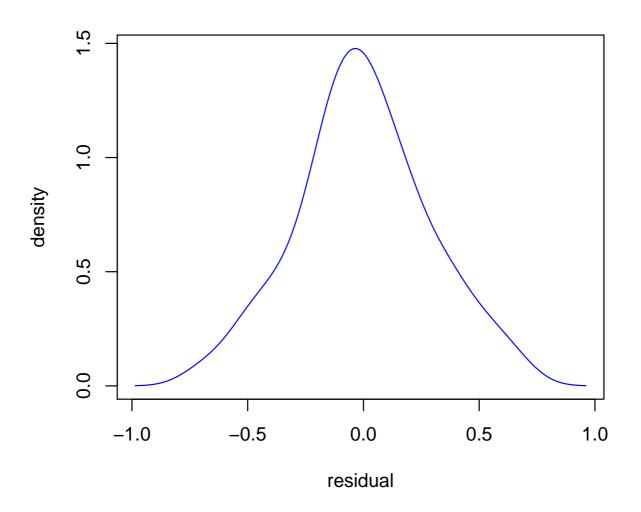
$$\hat{\beta}_0 = 5.588$$
  $\hat{\beta}_1 = 0.0129$   $\hat{\beta}_2 = -0.0830$   $\hat{\beta}_3 = -0.158$   $\hat{\beta}_4 = -0.00911$ .

The MLE of  $\sigma^2$  is  $\hat{\sigma}^2 = 0.0814$ .

## Residual plot



## Kernel estimate of density of $\epsilon_i$

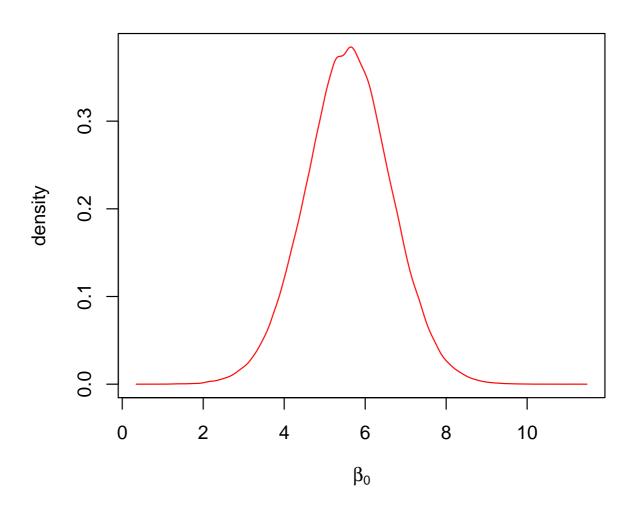


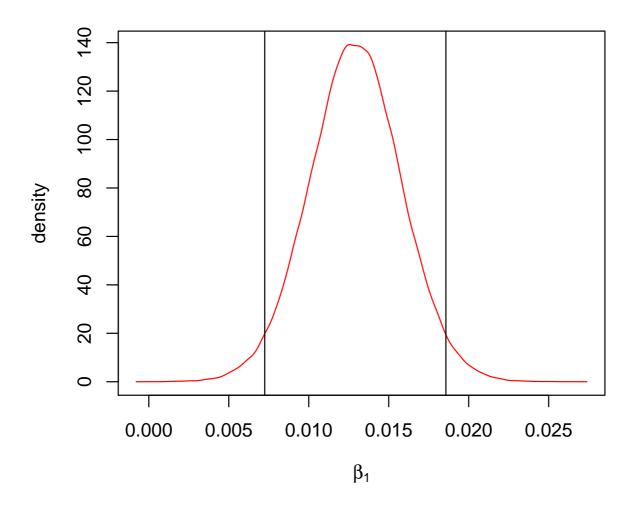
The estimate is computed from the residuals

$$\widehat{\epsilon}_i = y_i - \left[\widehat{\beta}_0 + \sum_{j=1}^4 \widehat{\beta}_j x_{ij}\right], \quad i = 1, \dots, n.$$

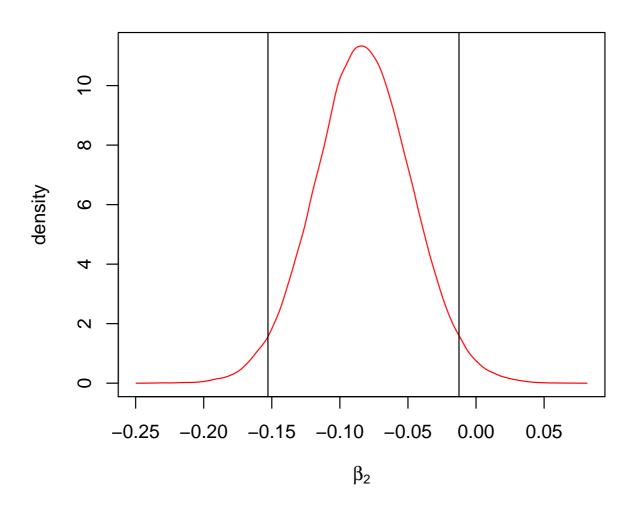
Based on the residual plot and the kernel estimate, the assumption of i.i.d. normal error terms seems reasonable.

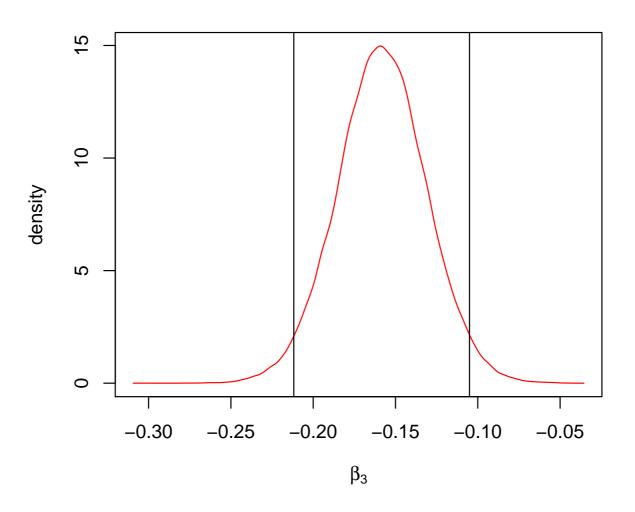
- I used Gibbs sampling to generate 100,000 observations from the posterior based on the noninformative prior on p. 213N.
- Sample autocorrelation functions for the output showed essentially no serial correlation among the generated  $\beta$ s.
- Minimal correlation was observed among the generated values of  $\sigma$ .

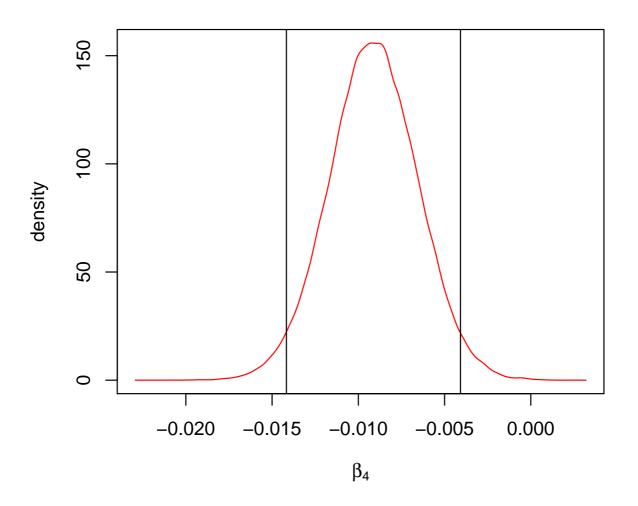




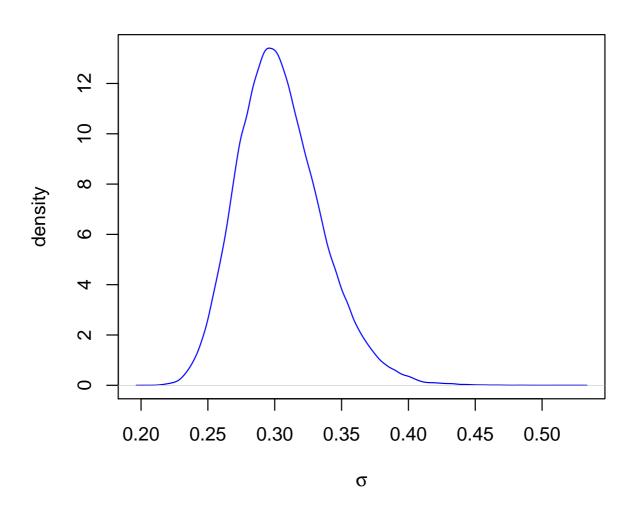
HPD region indicated by black lines.







## Kernel estimate of posterior of $\sigma$



#### Correlation matrix of $oldsymbol{eta}$ from the Gibbs output

Note that there is little correlation among  $\beta_1$ ,  $\beta_2$ ,  $\beta_3$  and  $\beta_4$ .

This is good because inference about any one regression coefficient is not affected by that for other coefficients.

The lack of correlation also shows that we don't need to worry about collinearity of predictors.

A large positive correlation between two predictors translates into a large negative correlation between the corresponding  $\beta$ s.

#### Selecting a linear regression model

To this point we have not addressed the question of model selection. We have always assumed that we know the correct parametric model.

Linear regression is a natural place to address model selection.

Often investigators throw in "all variables but the kitchen sink," and then go through a process of finding a best subset of variables.

Bayesian methods provide a means of estimating a best subset, i.e., a best model.

Let's first consider the general problem of model selection, and then apply what we learn to linear regression.

Let  $M_1, M_2, \ldots, M_N$  be the models under consideration. These might be of different dimensions, i.e., they might have different numbers of parameters.

- ullet  $heta_k$ : vector of parameters for model  $M_k$
- $p(\theta_k|k)$ : prior for  $\theta_k$  assuming that  $M_k$  is the correct model
- ullet  $p_k$ : prior probability of  $M_k$
- $p(y|\theta_k,k)$ : likelihood of data assuming true model is  $M_k$  and parameters are  $\theta_k$

We have

$$p(\boldsymbol{\theta}_k, k|\boldsymbol{y}) = \frac{p(\boldsymbol{y}|\boldsymbol{\theta}_k, k)p(\boldsymbol{\theta}_k|k)p_k}{m(\boldsymbol{y})},$$

where

$$m(\mathbf{y}) = \sum_{k=1}^{N} p_k m_k(\mathbf{y}),$$

and

$$m_k(y) = \int_{\Theta_k} p(y|\theta_k, k) p(\theta_k|k) d\theta_k.$$

Why? Because of the law of total probabilities.

$$m(\mathbf{y}) = \sum_{k=1}^{N} m(\mathbf{y}|M_k) p_k$$

$$= \sum_{k=1}^{N} \int_{\Theta_k} p(\mathbf{y}, \boldsymbol{\theta}_k | k) d\boldsymbol{\theta}_k \cdot p_k$$

$$= \sum_{k=1}^{N} \int_{\Theta_k} p(\mathbf{y} | \boldsymbol{\theta}_k, k) p(\boldsymbol{\theta}_k | k) d\boldsymbol{\theta}_k \cdot p_k$$

$$= \sum_{k=1}^{N} p_k m_k(\mathbf{y}).$$

An apparently reasonable way to select a model is to choose one that maximizes posterior probability:

$$P(M_k|\mathbf{y}) = \int_{\Theta_k} p(\boldsymbol{\theta}_k, k|\mathbf{y}) d\boldsymbol{\theta}_k$$
$$= \frac{m_k(\mathbf{y})p_k}{m(\mathbf{y})}.$$

A problem that arises is that when using  $P(M_k|y)$  as a criterion for model selection, one should **not** use improper priors for any of  $p(\theta_1|1), \ldots, p(\theta_N|N)$ .

#### Why?

When considering a single model, the arbitrary constant in front of the improper prior cancels out when computing the posterior.

However, these constants do not cancel out when computing  $P(M_k|\mathbf{y})$ .

Each improper prior  $\operatorname{could}$  be  $C_k p(\boldsymbol{\theta}_k|k)$ , where  $C_k$  is arbitrary. That means we can write

$$P(M_k|\mathbf{y}) = \frac{C_k m_k(\mathbf{y}) p_k}{\sum_{j=1}^N p_j C_j m_j(\mathbf{y})}$$
$$= \frac{m_k(\mathbf{y}) p_k}{\sum_{j=1}^N p_j (C_j/C_k) m_j(\mathbf{y})},$$

and hence  $P(M_k|y)$  is not well-defined.

So, when doing model selection, we need to use proper priors.

Unfortunately, even weakly informative priors, such as unit information priors, don't always work very well for model selection. This will be seen in Example 17.

In the linear regression setting, consider the problem of selecting a best subset of predictors.

Suppose we have a total of p predictors, and let  $z = (z_1, \ldots, z_p)$  be a p-vector of 0s and 1s.

We have  $z_j = 1$  if predictor  $x_j$  is in the model and 0 otherwise.

For a given model z, let  $X_z$  and  $\beta_z$  be the corresponding design matrix and vector of regression coefficients.

For example, suppose z = (1, 0, 1, 0). Then

$$\boldsymbol{\beta}_{\boldsymbol{z}}^T = (\beta_1, \beta_3)$$

and  $X_z$  is an  $n \times 2$  matrix whose first and third columns contain  $x_{11}, \ldots, x_{n1}$  and  $x_{13}, \ldots, x_{n3}$ , respectively.

For the prior  $p(\boldsymbol{\beta_z}, \sigma^2 | \boldsymbol{z})$  suppose that we use

$$\boldsymbol{\beta_z}|(\sigma^2, \boldsymbol{z}) \sim N(\boldsymbol{0}, n\sigma^2(\boldsymbol{X_z^T X_z})^{-1})$$

and

$$\sigma^2 | z \sim \text{inverse gamma} \left( \frac{1}{2}, \frac{s_z^2}{2} \right),$$

where  $s_z^2$  is the estimated error variance under model z.

#### Notes:

- The mean vector 0 in the prior for the regression coefficients results from an invariance argument (pp. 156-57).
- Using  $s_z^2$  in the prior for  $\sigma^2$  is "cheating," but is consistent with the principle of "unbiased but weak prior information." The weak part comes from using 1/2 as the first parameter.

For a model z define

$$RSS_z = y^T \left[ I - \left( \frac{n}{n+1} \right) X_z (X_z^T X_z)^{-1} X_z^T \right] y.$$

Note that the usual RSS, i.e., residual sum of squares, does not have the factor of n/(n+1).

Hoff shows that using the prior on the previous page leads to

$$\frac{p(\boldsymbol{z}_a|\boldsymbol{y})}{p(\boldsymbol{z}_b|\boldsymbol{y})} = \frac{p(\boldsymbol{z}_a)}{p(\boldsymbol{z}_b)} \cdot \text{Bayes factor},$$

where p(z) is the prior probability of model z and

Bayes factor = 
$$(1+n)^{(p_b-p_a)/2} \left(\frac{s_{\boldsymbol{z}_a}^2}{s_{\boldsymbol{z}_b}^2}\right)^{1/2}$$

$$\times \left(\frac{s_{\boldsymbol{z}_b}^2 + \text{RSS}_{\boldsymbol{z}_b}}{s_{\boldsymbol{z}_a}^2 + \text{RSS}_{\boldsymbol{z}_a}}\right)^{(n+1)/2}$$

where  $p_a$  and  $p_b$  are the numbers of 1s in the vectors  $z_a$  and  $z_b$ , respectively.

# Example 17 Model selection in the oxygen uptake example

We'll use the priors of p. 234N and evaluate each of the 15 subsets of predictors. We'll use the full model (with all four predictors) as a benchmark.

In other words,  $z_b$  in the odds ratio on p. 235N will always be the full model.

I will assign an equal prior probability of 1/15 to each of the 15 models, meaning that the Bayes factor on p. 235 is the ratio of the posterior probabilities of two models.

#### Results for oxygen uptake data

	Probability relative	
Model	to full model	BIC
$\overline{x_1}$	0.007	66.61
$x_2$	0.001	72.12
$x_3$	0.049	60.84
$x_{4}$	0.000	75.51
$x_{1}, x_{2}$	0.004	66.09
$x_1, x_3$	0.411	51.52
$x_1, x_4$	0.002	67.88
$x_{2}, x_{3}$	0.031	59.96
$x_{2}, x_{4}$	0.000	74.66
$x_3, x_4$	0.053	58.32
$x_1, x_2, x_3$	0.230	50.62
$x_{1}, x_{2}, x_{4}$	0.002	67.08
$x_{1}, x_{3}, x_{4}$	1.578	43.63
$x_{2}, x_{3}, x_{4}$	0.037	56.85
$x_1, x_2, x_3, x_4$	1.000	41.70

The procedure of p. 235 points to the model with the predictors  $x_1$ ,  $x_3$  and  $x_4$  as being best.

Our Gibbs sampling suggested that all four predictors should be in the model.

An all purpose model selector with a Bayesian motivation is BIC.

According to this criterion we choose the model with the smallest value of

$$BIC(k) = -2 \log p(y|\widehat{\theta}_k, k) + d_k \log n,$$

where  $\hat{\theta}_k$  is the MLE for model k and  $d_k$  is the number of parameters in model k.

The BIC values on p. 237 point to the full model as being the best, which agrees with the results of our Gibbs sampling.

This example is to be continued ...

It turns out that if the prior variance for  $\beta_z$  is too large, then there is a strong tendency for too small a model to have the largest posterior probability.

This point has been made by a number of authors, including those below:

- Chipman, H., George, E.I. and McCulloch, R.E. (2001). The practical implementation of Bayesian model selection (with discussion). *IMS Lecture Notes - Monograph* Series 38 65-116.
- Berger, J.O. and Pericchi, L.R. (1996).
   The intrinsic Bayes factor for model selection and prediction. *Journal of the American Statistical Association* 91 109-122.

Instead of the prior on p. 235N, suppose we use

$$\boldsymbol{\beta_z}|(\sigma^2, \boldsymbol{z}) \sim N(\boldsymbol{0}, g\sigma^2(\boldsymbol{X_z^T X_z})^{-1})$$

and

$$\sigma^2|z\sim \text{inverse gamma}\left(\frac{\nu_0}{2},\frac{s^2}{2}\right),$$

where  $s^2$  is the sample variance of y.

The use of  $s^2$  was suggested by Chipman, George and McCulloch (2001) (CGM), since it represents an upper bound for  $\sigma^2$ .

CGM recommend a small value, such as 2 or 3, for  $\nu_0$ .

The prior for  $\beta_z$  is the so-called g-prior proposed by Zellner (1986).

Previously we used g = n, but as we saw in our example this can produce a model smaller than what seems reasonable.

A common way of standardizing the independent variables is as follows:

$$v_{ij} = \frac{x_{ij} - \bar{x}_i}{s_i}, \quad i = 1, \dots, p, \ j = 1, \dots, n,$$

where  $\bar{x}_i$  and  $s_i$  are the mean and standard deviation, respectively, of  $x_{i1}, \ldots, x_{in}$ .

If the predictors are so standardized, CGM recommend using  $g = (2.85)^2$ .

Using the prior on p. 241N, we have

$$\frac{p(\boldsymbol{z}_a|\boldsymbol{y})}{p(\boldsymbol{z}_b|\boldsymbol{y})} = \frac{p(\boldsymbol{z}_a)}{p(\boldsymbol{z}_b)} \cdot (1+g)^{(p_b-p_a)/2} \\
\times \left(\frac{\nu_0 \sigma_0^2 + \text{RSS}_{g,\boldsymbol{z}_b}}{\nu_0 \sigma_0^2 + \text{RSS}_{g,\boldsymbol{z}_a}}\right)^{(n+\nu_0)/2},$$

where  $RSS_{g,z}$  is defined as was  $RSS_z$  but with g in place of n.

### Example 17 continued

I recomputed the odds ratios, this time using the formula at the bottom of p. 242N. (I standardized the predictors as on p. 242N.)

I took  $\nu_0 = 2$ ,  $\sigma_0^2 = 0.198$  (the sample variance of the  $y_i$ s) and  $g = (2.85)^2$ .

As before I gave equal prior probability to each of the fifteen models.

The resulting odds ratios are given on the next page.

## Posterior probability of each model in the oxygen uptake example relative to that of the full model

	Probability relative
Model	to full model
$\overline{x_1}$	0.000
$x_2$	0.000
$x_3$	0.000
$x_{4}$	0.000
$x_{1}, x_{2}$	0.000
$x_{1}, x_{3}$	0.009
$x_{1}, x_{4}$	0.000
$x_{2}, x_{3}$	0.000
$x_{2}, x_{4}$	0.000
$x_{3}, x_{4}$	0.001
$x_1, x_2, x_3$	0.021
$x_1, x_2, x_4$	0.000
$x_{1}, x_{3}, x_{4}$	0.324
$x_2, x_3, x_4$	0.002
$x_1, x_2, x_3, x_4$	1.000

This analysis places the highest probability on the full model. The model with  $x_1$ ,  $x_3$  and  $x_4$  is reasonably probable, but none of the others seem viable.

Note that knowing all the odds ratios tells us the posterior probability of each model.

Let  $P_j$  be the posterior probability of model j. The odds ratios are  $o_j = P_j/P_N$ , and so

$$\sum_{j=1}^{N} o_j = \frac{1}{P_N}.$$

Therefore,

$$P_j = \frac{o_j}{\sum_{i=1}^N o_i}, \quad j = 1, \dots, n.$$

Using this fact, the posterior probability of the full model is 0.729 and that of the model with  $x_1$ ,  $x_3$  and  $x_4$  is 0.245.

No other model has posterior probability higher than 0.0168.

It is interesting to look at how BIC compares with the Bayesian posterior probabilities. For our linear model,

$$BIC(z) = n \left[ \log \hat{\sigma}_z^2 + 1 + \log(2\pi) \right] + (p+1) \log n,$$

where  $\hat{\sigma}_{z}^{2}$  is the MLE of  $\sigma^{2}$  for model z.

I can compare  $z_a$  and  $z_b$  by taking the difference of their BIC values. This gives

$$BIC(z_b) - BIC(z_a) =$$

$$n \log(\hat{\sigma}_{z_b}^2 / \hat{\sigma}_{z_a}^2) + (p_b - p_a) \log n.$$

Now, from page 242N, if we take  $p(z_a) = p(z_b)$ , then

$$2\log\left[\frac{p(\boldsymbol{z}_a|\boldsymbol{y})}{p(\boldsymbol{z}_b|\boldsymbol{y})}\right] =$$

$$(n+\nu_0)\log\left[\frac{\nu_0\sigma_0^2 + \mathsf{RSS}_{g,\boldsymbol{z}_b}}{\nu_0\sigma_0^2 + \mathsf{RSS}_{g,\boldsymbol{z}_a}}\right]$$

$$+(p_b-p_a)\log(g+1).$$

So, there is a strong similarity between the difference of BIC values and

$$2\log\left[rac{p(oldsymbol{z}_a|oldsymbol{y})}{p(oldsymbol{z}_b|oldsymbol{y})}
ight].$$

In fact, for large n they are essentially the same if we take g=n.

However, if we take g to be a fixed constant, as suggested by CGM, then the two criteria can be quite different.