## An Observation About Subsection 6.6: On Average Training MSE Underestimates Validation MSE

The validating mean squared error is often used to evaluate the performance of a predictor. The expected value of the validating mean squared error, taken over the distribution of training data y and validating data  $y_v$  is shown in equation (6.65) on page 285:

$$E_{y,y_v}(MSE_v) = \sigma^2 + \frac{1}{N} \sum_{i} Var(\hat{y}_{v,i}) + bias^2.$$
(6.65)

On the other hand, the expected value of the training mean squared error, taken over the distribution of training data y is shown in equation (6.66):

$$E_y(MSE_t) = \sigma^2 + \frac{1}{N} \sum_i Var(\hat{y}_i) + bias^2 - \frac{2}{N} \sum_i Cov(y_i, \hat{y}_i), \qquad (6.66)$$

indicating that the training mean squared error

$$MSE_t = \frac{1}{N} \sum_{i=1}^{N} (y_i - \hat{y}_i)^2$$

is a poor estimate of (6.65).

The predicted values in training and validating data are  $\hat{y}_i = \hat{f}(x_i)$  and  $\hat{y}_{v,i} = \hat{f}(x_{v,i})$ , a function of known covariates  $x_i$  and  $x_{v,i}$ , respectively. These covariates are not necessarily the same. However with common covariates such as genetic markers, for a given number of these covariates, as the number of records increases the second terms after the equal sign of these expressions become smaller and more alike. Regardless of the number of records, if the covariates in the training and validating data take the same values,  $\frac{1}{N} \sum_i \text{Var}(\hat{y}_{v,i}) = \frac{1}{N} \sum_i \text{Var}(\hat{y}_i)$ . This is not made explicit in the book. Then it follows that

$$E_y(MSE_t) = E_{y,y_v}(MSE_v) - \frac{2}{N} \sum_i Cov(y_i, \hat{y}_i).$$
(6.67)

## References