Example of Data-Determined Statistical Distance: Biting-Fly Data

In this example the sample mean vector and sample variance-covariance matrix are used in place of the population mean vector and population variance-covariance matrix to calculate statistical distances. Table 6.15 on page 352 of the textbook contains data from a study that examines the differences between two morphologically-similar species of biting flies. Measurements were taken on n = 70 specimens, 35 from each of the two species. For the moment we will ignore the fact that there are two species and simply consider p = 2 variables measured on each of the 70 specimens using wing length (x_1) and wing width (x_2) .

On the next page is the series of \mathbf{R} code commands and output supporting the calculations given below. From the output we note that the sample mean vector is $\mathbf{\bar{x}} = \begin{pmatrix} 97.90 \\ 43.33 \end{pmatrix}$ and the sample $\begin{pmatrix} 37.60 & 14.99 \end{pmatrix}$

variance-covariance matrix is $\mathbf{S} = \begin{pmatrix} 37.60 & 14.99 \\ 14.99 & 16.57 \end{pmatrix}$. The eigenvalues and eigenvectors of \mathbf{S} are

calculated using the \mathbf{R} command eigen (S). The eigenvalues of \mathbf{S} are $\lambda_1 = 45.39$ and $\lambda_2 = 8.78$ with corresponding eigenvectors $\mathbf{e}_1 = \begin{pmatrix} 0.89 \\ 0.46 \end{pmatrix}$ and $\mathbf{e}_1 = \begin{pmatrix} -0.46 \\ 0.89 \end{pmatrix}$. Since $\mathbf{A} = \mathbf{S}^{-1} = \begin{pmatrix} 0.042 & -0.038 \\ -0.038 & 0.094 \end{pmatrix}$

(found using the \mathbf{R} command ginv(S) – after first loading the MASS library), this means that the eigenvalues of \mathbf{A} are $\lambda_1^{-1} = (45.39)^{-1} = 0.02$ and $\lambda_2^{-1} = (8.78)^{-1} = 0.11$, with the same eigenvectors as those of \mathbf{S} . The axes of the ellipse centered at $\overline{\mathbf{x}}$ (that is the set of points all of which have statistical distance c from $\overline{\mathbf{x}}$) are

$$\overline{\mathbf{x}} \pm c\sqrt{\lambda_1} \mathbf{e}_1 = \begin{pmatrix} 97.90 \\ 43.33 \end{pmatrix} \pm c\sqrt{45.39} \begin{pmatrix} 0.89 \\ 0.46 \end{pmatrix} = \begin{pmatrix} 97.90 \pm 5.98c \\ 43.33 \pm 3.11c \end{pmatrix},$$

and

$$\overline{\mathbf{x}} \pm c\sqrt{\lambda_2} \mathbf{e}_2 = \begin{pmatrix} 97.90 \\ 43.33 \end{pmatrix} \pm c\sqrt{8.78} \begin{pmatrix} -0.46 \\ 0.88 \end{pmatrix} = \begin{pmatrix} 97.90 \pm 1.36c \\ 43.33 \pm 2.61c \end{pmatrix}.$$

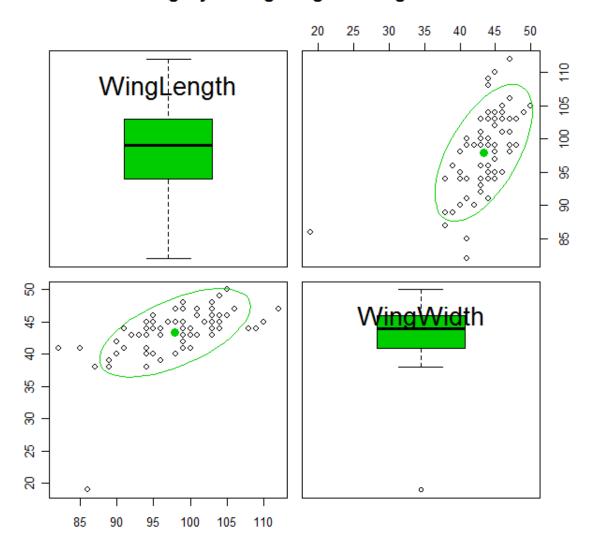
And finally, the formula for the squared statistical distance of a point $\mathbf{x} = \begin{pmatrix} x_1 \\ x_2 \end{pmatrix}$ from $\overline{\mathbf{x}}$ is:

$$d^{2}(\mathbf{x}, \overline{\mathbf{x}}) = (\mathbf{x} - \overline{\mathbf{x}})' \mathbf{A} (\mathbf{x} - \overline{\mathbf{x}}) = (\mathbf{x} - \overline{\mathbf{x}})' \mathbf{S}^{-1} (\mathbf{x} - \overline{\mathbf{x}})$$
$$= 0.042(x_{1} - 97.90)^{2} - 0.076(x_{1} - 97.90)(x_{2} - 43.33) + 0.094(x_{2} - 43.33)^{2}$$

R code and Output for Biting Flies Example

```
> Flys <-
data.frame(WingLength=X[,1],WingWidth=X[,2],ThirdPalpLength=X[,3],ThirdPalpWidth=X[,4],FourthPalpLe
ngth=X[,5],LengthSeg12=X[,6],LengthSeg13=X[,7],Species=X[,8])
> mean(Flys)
   WingLength
                    WingWidth ThirdPalpLength ThirdPalpWidth
    97.900000
                  43.328571
                                 37.342857
                                                14.585714
                    LengthSeg12
                                    LengthSeg13
FourthPalpLength
                                                      Species
    27.814286
                   9.614286
                                 9.542857
                                               0.500000
> # construct the covariance matrix for the first two variables
> # (wing length and wing width)
>#
> S < -cov(Flys[,1:2])
>#
> # Find the e-values / e-vectors of S
>#
> eigen(S)
$values
[1] 45.394341 8.775846
$vectors
      [,1]
             [,2]
[1,] -0.8871910 0.4614024
[2,] -0.4614024 -0.8871910
> # need the MASS library to use the ginv command
>#
> library(MASS)
>#
> # calculate the inverse of S
> A < -ginv(S)
> A
       [,1]
              [,2]
[1,] 0.04159821 -0.03762762
[2,] -0.03762762  0.09438010
>#
> # Find the e-values / e-vectors of A (compare to those for S)
> eigen(A)
$values
[1] 0.11394913 0.02202918
$vectors
      [,1]
             [,2]
[1,] -0.4614024 -0.8871910
[2,] 0.8871910 -0.4614024
> spm(Flys[,1:2],diagonal=list(method="boxplot"),smooth=FALSE,regLine=FALSE,ellipse=
list(levels=c(0.75), robust=FALSE, fill=FALSE),main=c("Biting Flys: Wing Length & Wing Width"))
```

Biting Flys: Wing Length & Wing Width



(Courtesy of Dr. Roy St. Laurent)