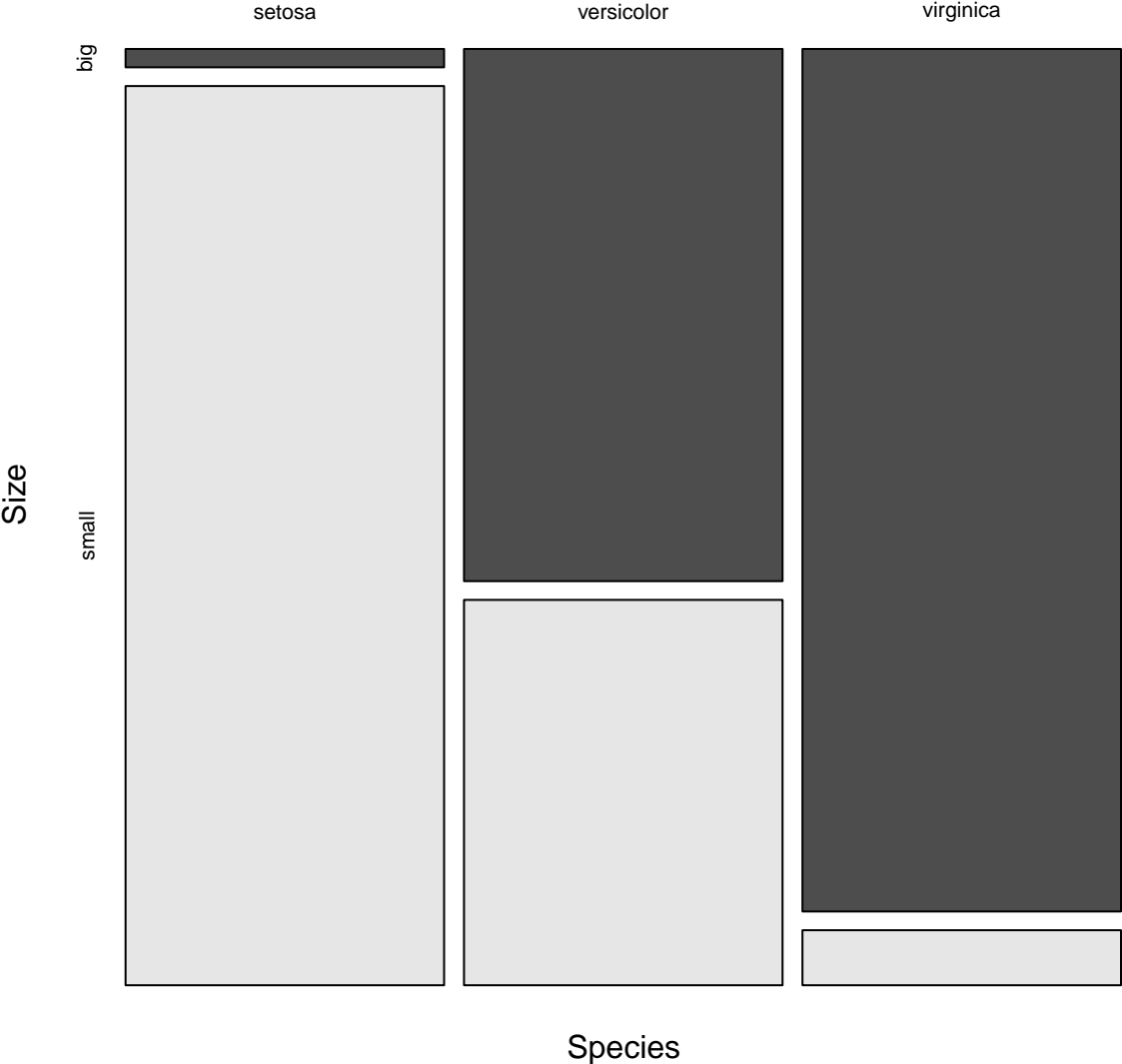
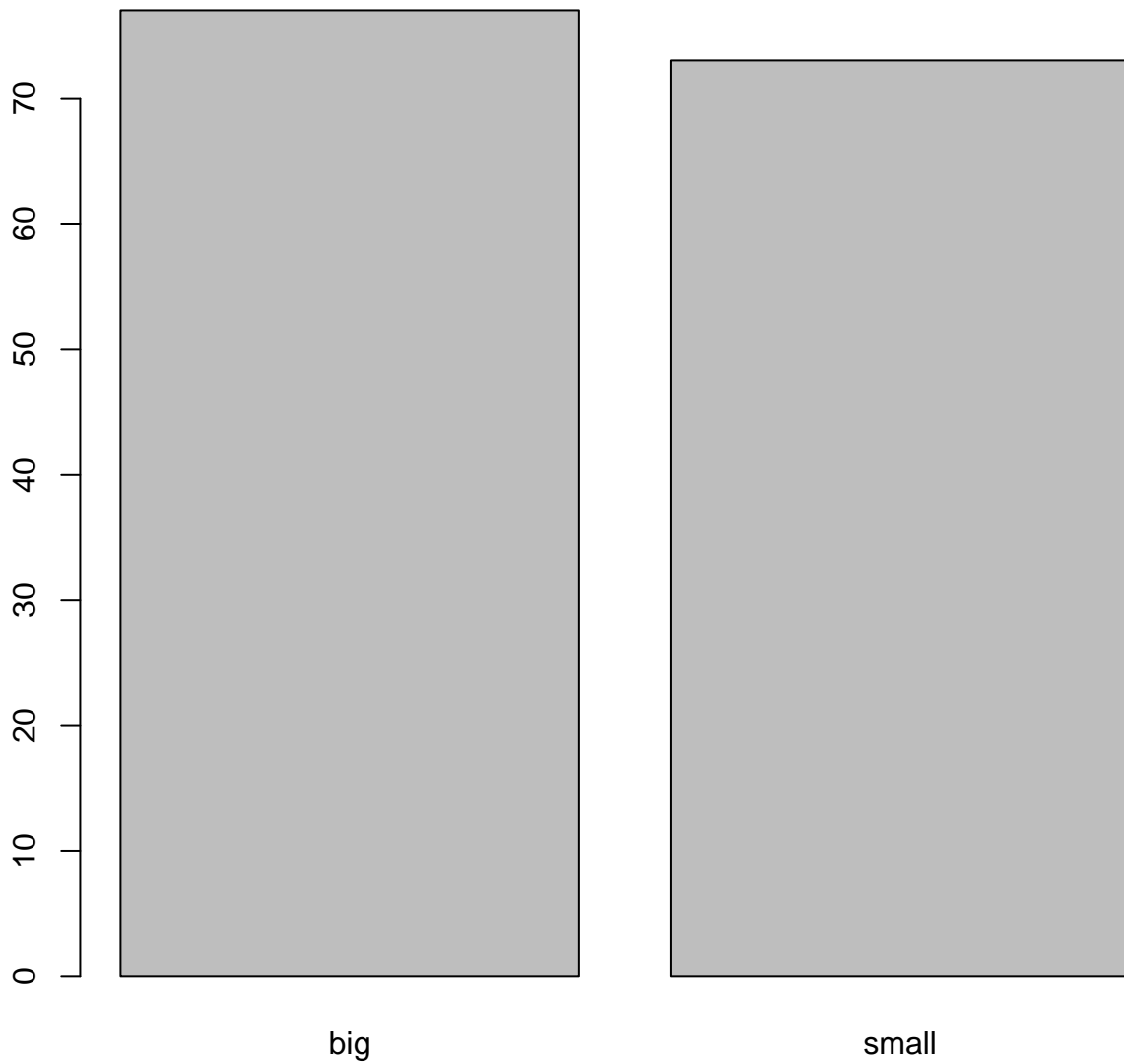
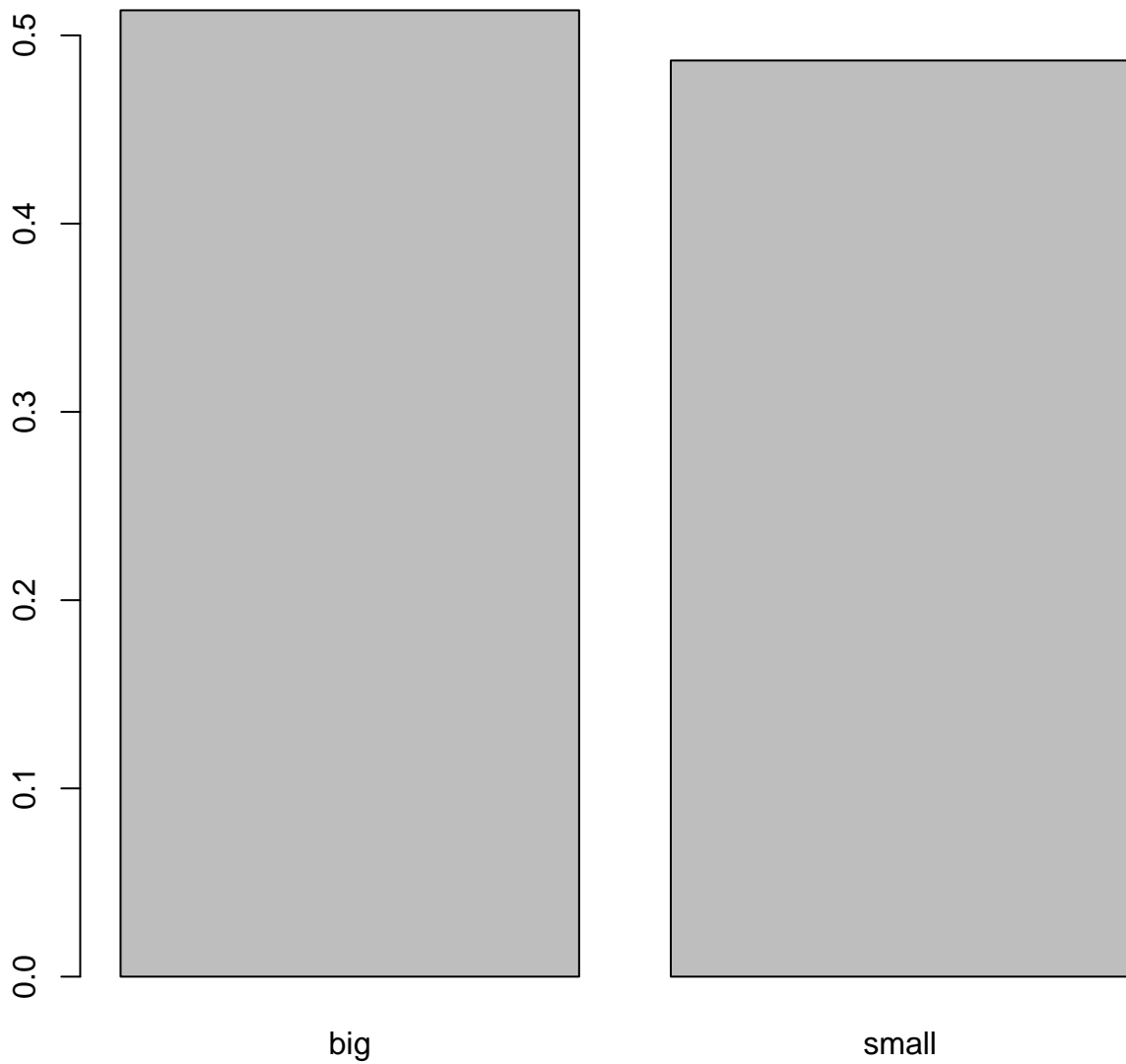


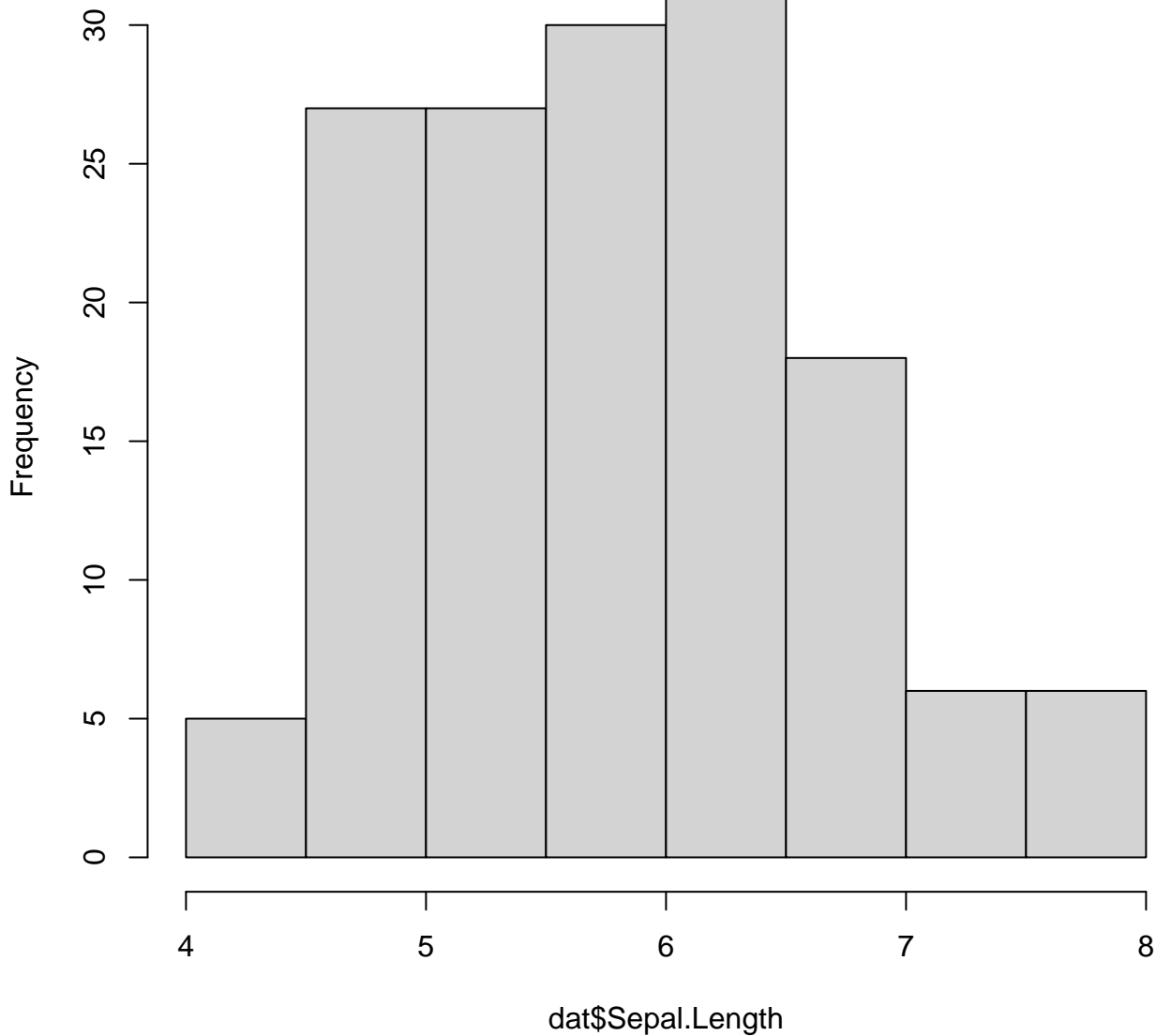
table(dat\$Species, dat\$size)

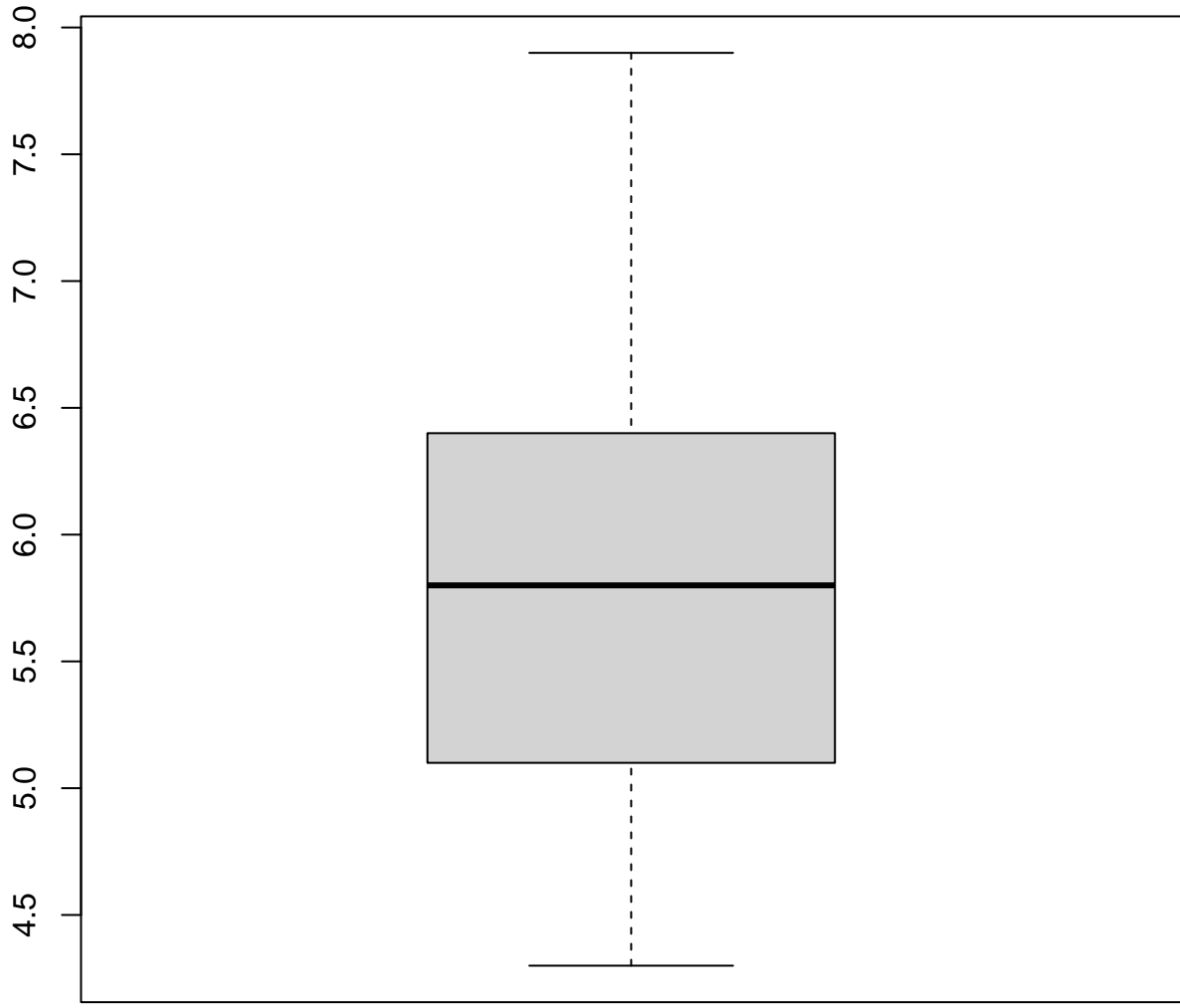


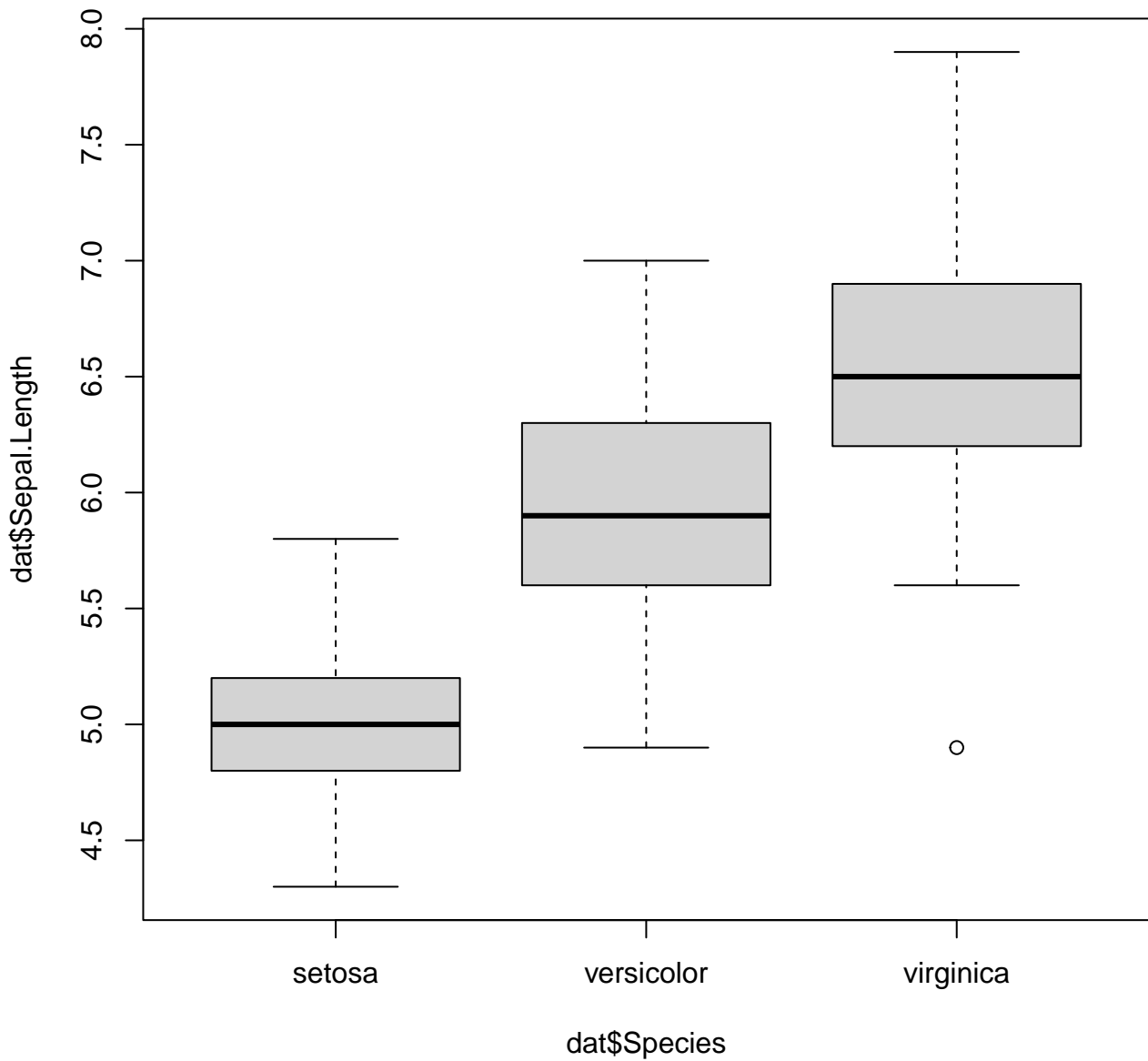


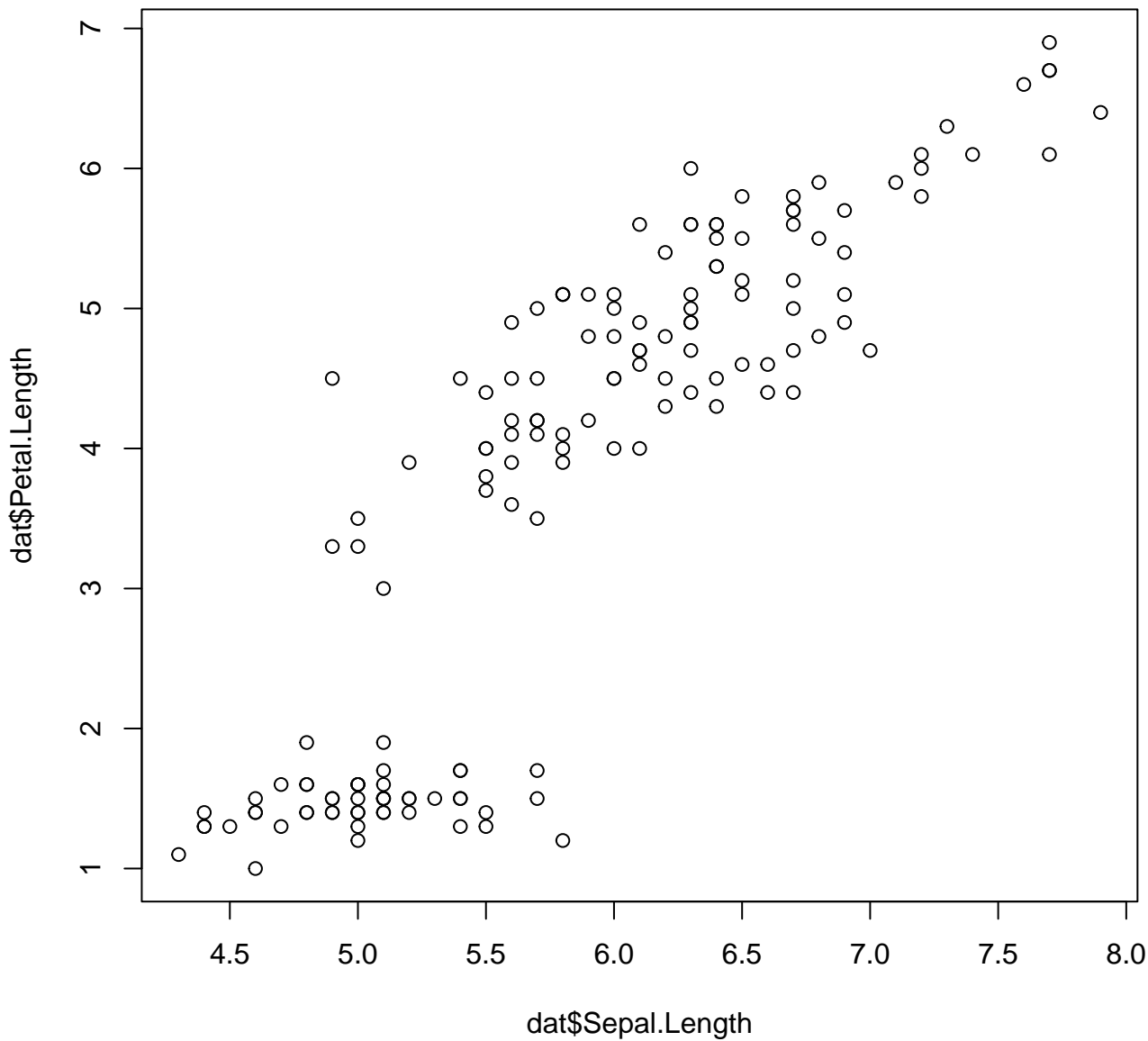


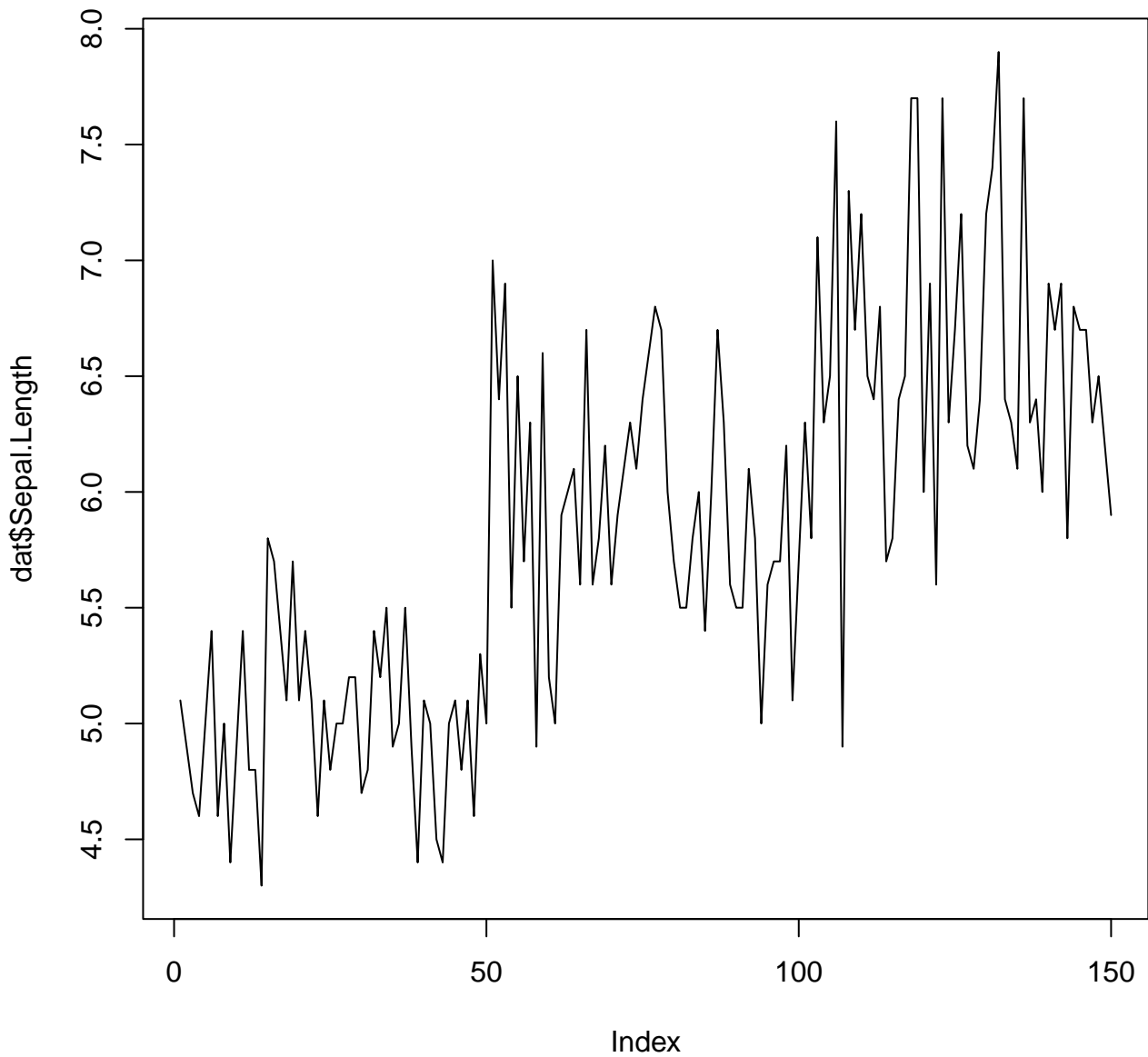
**Histogram of dat\$Sepal.Length**





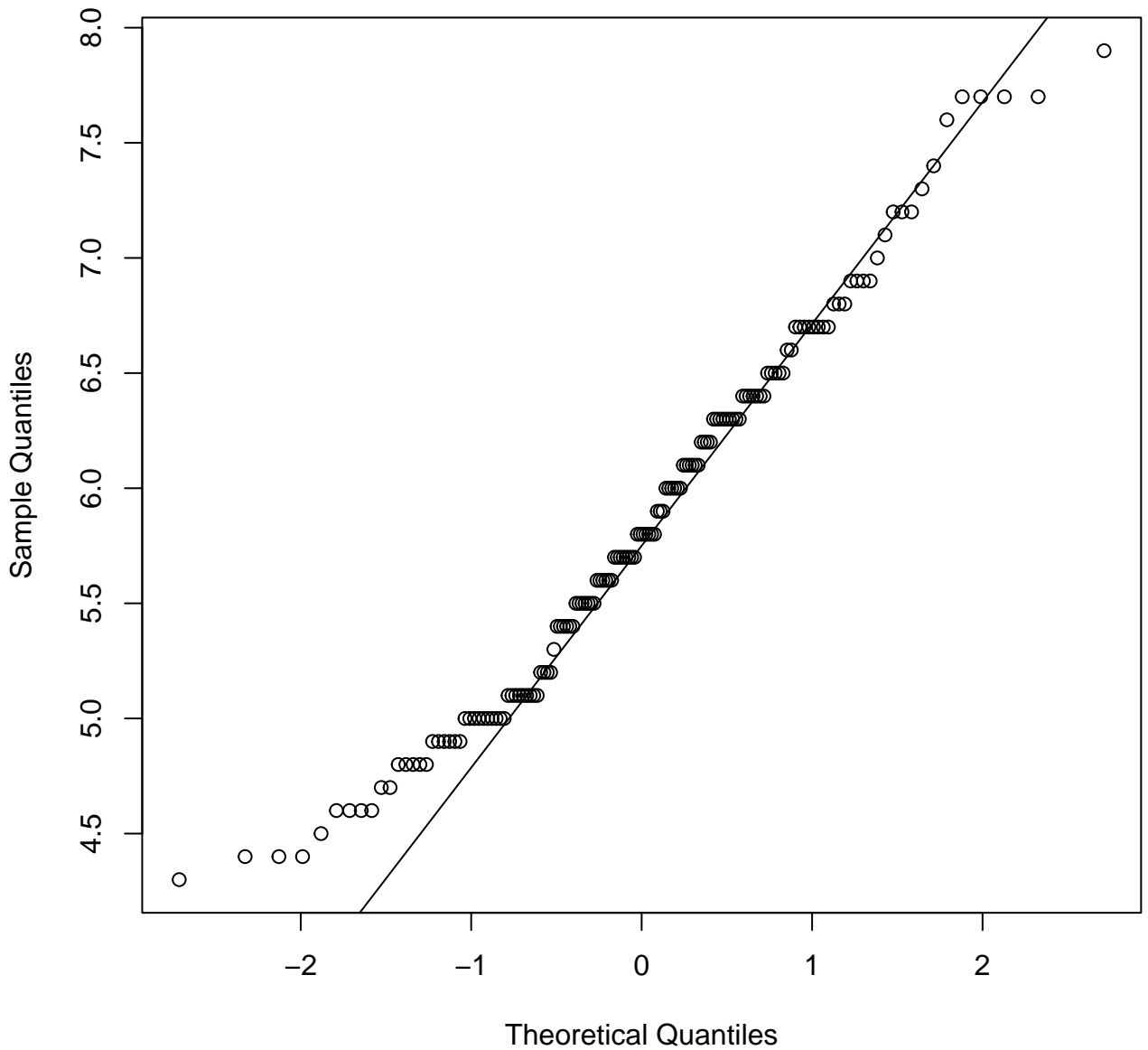




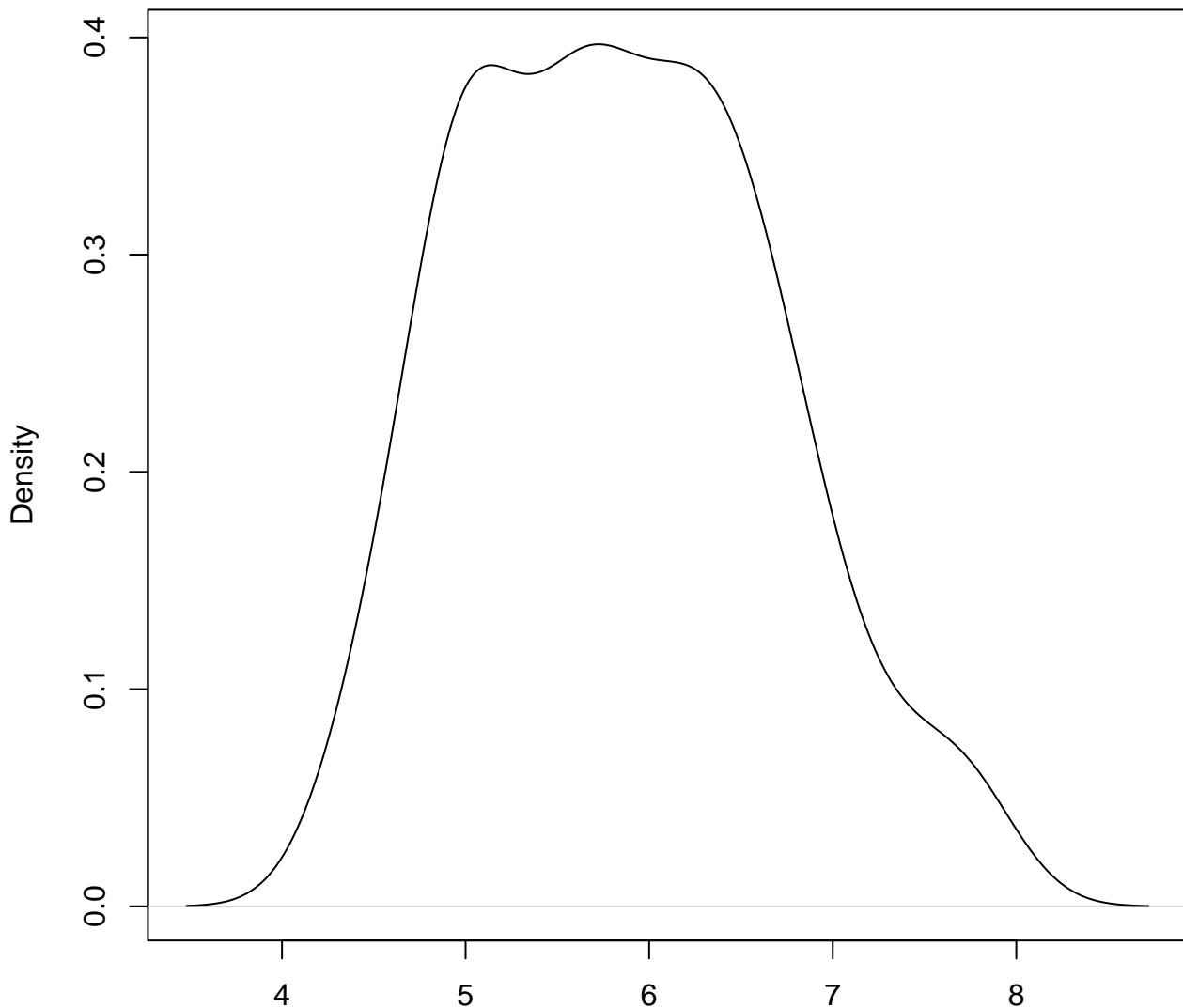




Normal Q-Q Plot



**density.default(x = dat\$Sepal.Length)**



N = 150 Bandwidth = 0.2736