**> drinks<-read.table(file.choose(),header=T)**

**> head(drinks)**

Carbohydrate carbohydrateprotein

1 10.58 10.53

2 10.89 11.16

3 10.17 10.31

4 11.61 11.79

5 11.79 11.88

6 9.72 9.81

**> attach(drinks)**

**> shapiro.test(Carbohydrate-carbohydrateprotein)**

Shapiro-Wilk normality test

data: Carbohydrate - carbohydrateprotein

W = 0.96995, p-value = 0.7538

1. .
2. .