

R version 2.9.0 (2009-04-17)
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Tippen Sie 'demo()' für einige Demos, 'help()' für on-line
Hilfe, oder
'help.start()' für eine HTML Browserschnittstelle zur Hilfe.
Tippen Sie 'q()', um R zu verlassen.

[R.app GUI 1.28 (5395) i386-apple-darwin8.11.1]

[Workspace restored from /Users/oppl/.RData]

```
> daten = read.csv("/Users/oppl/Desktop/Auswertung/
daten.csv")
> attach(daten)
> summary(PTA)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
0.4062 0.4286 0.5000 0.5228 0.5625 0.8571
> summary(PTB)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
0.1429 0.2647 0.3333 0.3242 0.3750 0.4412
> summary(PTC)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's
0.03125 0.14880 0.21240 0.18580 0.23330 0.33330 3.00000
> summary(S1TA)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's
0.5652 0.5789 0.5862 0.6852 0.7826 0.8667 8.0000
> summary(S1TB)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's
0.1333 0.2174 0.4138 0.3148 0.4211 0.4348 8.0000
```

```

> summary(S1TC)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
0.2069 0.2069 0.2069 0.2069 0.2069 0.2069 16.0000
> summary(S2TA)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
0.5294 0.5556 0.6250 0.6529 0.7143 0.8462  8.0000
> summary(S2TB)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
0.1538 0.2857 0.3750 0.3471 0.4444 0.4706  8.0000
> summary(S2TC)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
0.1579 0.1579 0.1579 0.1579 0.1579 0.1579 16.0000
> shapiro.test(PTA)

```

Shapiro-Wilk normality test

```

data:  PTA
W = 0.8179, p-value = 0.003611

```

```

> shapiro.test(PTB)

```

Shapiro-Wilk normality test

```

data:  PTB
W = 0.9485, p-value = 0.4329

```

```

> shapiro.test(PTC)

```

Shapiro-Wilk normality test

```

data:  PTC
W = 0.9447, p-value = 0.4818

```

```

> shapiro.test(S1TA)

```

Shapiro-Wilk normality test

```

data:  S1TA
W = 0.7717, p-value = 0.009657

```

```

> shapiro.test(S1TB)

```

Shapiro-Wilk normality test

```
data: S1TB
W = 0.7717, p-value = 0.009657
```

```
> shapiro.test(S1TC)
Fehler in shapiro.test(S1TC) :
  Stichprobengröße muss zwischen 3 und 5000 liegen
> shapiro.test(S2TA)
```

Shapiro-Wilk normality test

```
data: S2TA
W = 0.8783, p-value = 0.1508
```

```
> shapiro.test(S2TB)
```

Shapiro-Wilk normality test

```
data: S2TB
W = 0.8783, p-value = 0.1508
```

```
> shapiro.test(S2TC)
Fehler in shapiro.test(S2TC) :
  Stichprobengröße muss zwischen 3 und 5000 liegen
> kruskal.test(list(PTA, PTB, PTC))
```

Kruskal-Wallis rank sum test

```
data: list(PTA, PTB, PTC)
Kruskal-Wallis chi-squared = 36.5432, df = 2,
p-value = 1.161e-08
```

```
> wilcox.test(S1TA, S1TB, exact=F, a="less", paired=T)
```

Wilcoxon signed rank test with continuity correction

```
data: S1TA and S1TB
V = 45, p-value = 0.9968
alternative hypothesis: true location shift is less than 0
```

```
> wilcox.test(S2TA, S2TB, exact=F, a="less", paired=T)
```

Wilcoxon signed rank test with continuity correction

```
data: S2TA and S2TB  
V = 45, p-value = 0.9968  
alternative hypothesis: true location shift is less than 0
```

```
>
```