There is a non-parametric test for a single factor with more than two levels. This is known as the 'Kruskal-Wallace' test.

- For the model $Y_{ij} = \mu + \tau_i + \epsilon_{ij}$, tests whether any two τ_i differ.
- makes same assumptions as Wilcoxon test
- still based upon ranks.

Kruskal-Wallis

This test is based upon the Kruskal-Wallis test statistic

$$H = \frac{12}{N(N+1)} \sum_{i=1}^{k} n_i (R_{\cdot i} - R_{\cdot \cdot})^2$$

Where $R_{\cdot \cdot} = \frac{N+1}{2}$ and $R_{\cdot i}$ is the mean rank in group i.

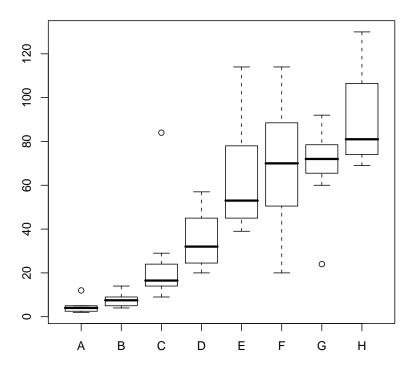
Kruskal-Wallis, Significance of H

Under H_0 , H should have a sampling distribution defined by χ^2 with df = k-1. If you calculate an H, you can use the R function pchisq(H,df) to calculate the probability of observing that value under the null

Kruskal-Wallis

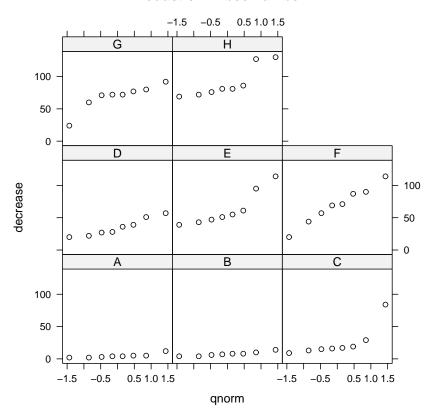
- > data(OrchardSprays)
- > boxplot(decrease ~ treatment, data = OrchardSprays, main = "Reduction in bee number

Reduction in bee number for different sprays

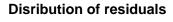


Kruskal-Wallis, cont

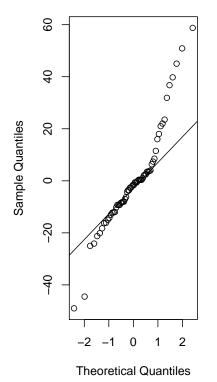
Reduction in bee number

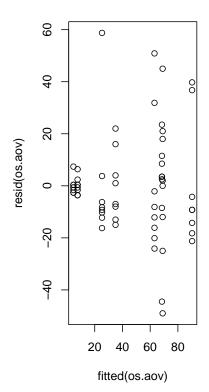


Kruskal-Wallis, cont



Fitted versus residual





Kruskal-Wallis, R implementation, cont

Kruskal-Wallis rank sum test

data: decrease by treatment

Kruskal-Wallis chi-squared = 48.8735, df = 7, p-value = 2.402e-08

> library(ctest)

> kruskal.test(decrease ~ treatment, data = OrchardSprays)