

**Example:** For Example 3.1.1 on page 81, we can apply the Kruskal-Wallis test to the ranks. The data is as follows:

Samples from normal populations with  $\mu_1 = 15$ ,  $\mu_2 = 25$ ,  $\mu_3 = 30$ ,  $\sigma = 9$ .

Treatment 1	6.08	22.29	7.51	34.36	23.68
Rank 1	1	3	2	13	5
Treatment 2	30.45	22.71	44.52	31.47	36.81
Rank 2	9	4	15	10	14
Treatment 3	32.04	28.03	32.74	23.84	29.64
Rank 3	11	7	12	6	8

Since there are 756,756 possible permutations, the exact Kruskal-Wallis test will take a significant amount of time to run. Instead, we use the MC option which randomly samples 10,000 permutations.

```
data oct18;
input treat resp @@;
cards;
1 1 1 3 1 2 1 13 1 5
2 9 2 4 2 15 2 10 2 14
3 11 3 7 3 12 3 6 3 8
;
proc npar1way data=oct18 anova scores=data;
class treat;
exact scores=data /MC; /* The MC option randomly samples the permutations */
var resp;
run;
```

The first and third screens displayed by SAS are as follows.

The NPARIWAY Procedure

Analysis of Variance for Variable resp  
Classified by Variable treat

treat	N	Mean
1	5	4.80
2	5	10.40
3	5	8.80

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Among	2	83.20	41.60	2.5366	0.1206
Within	12	196.80	16.40		

The NPAR1WAY Procedure

Monte Carlo Estimate for the Exact Test

Pr >= Chi-Square	
Estimate	0.1281
99% Lower Conf Limit	0.1195
99% Upper Conf Limit	0.1367
Number of Samples	10000
Initial Seed	725846367

In this case, we find a  $p$ -value of 0.1281. This is not very significant evidence to reject  $H_0$ .