

# Results

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## 1 Tables of Friedman, Bonferroni-Dunn, Holm, Hochberg and Hommel Tests

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Table 1: Average Rankings of the algorithms

Algorithm	Ranking
DE	3.394736842105262
CHC	1.368421052631579
G-CMA-ES	2.0526315789473686
VXQR1	3.1842105263157894

Friedman statistic considering reduction performance (distributed according to chi-square with 3 degrees of freedom: 31.342105263157823.  
Inan and Davenport statistic considering reduction performance (distributed according to F-distribution with 3 and 54 degrees of freedom: 21.987692307692196.

Bonferroni-Dunn's procedure rejects those hypotheses that have a p-value  $\leq 0.016666666666666666$ .

Holm's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ .

Hochberg's procedure rejects those hypotheses that have a p-value  $\leq 0.025$ .

Hommel's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ .

Table 2: Holm / Hochberg Table for  $\alpha = 0.05$

$i$	algorithm	$z = (R_0 - R_i) / SE$	$p$	Holm/Hochberg/Hommel
3	DE	4.837762640769081	1.3130881465534958E-6	0.016666666666666666
2	VXQR1	4.335137950819048	1.4566874163845129E-5	0.025
1	G-CMA-ES	1.6335302423376128	0.10235752557227855	0.05

Table 3: Holm / Hochberg Table for  $\alpha = 0.10$

$i$	algorithm	$z = (R_0 - R_i) / SE$	$p$	Holm/Hochberg/Hommel
3	DE	4.837762640769081	1.3130881465534958E-6	0.033333333333333333
2	VXQR1	4.335137950819048	1.4566874163845129E-5	0.05
1	G-CMA-ES	1.6335302423376128	0.10235752557227855	0.1

Bonferroni-Dunn's procedure rejects those hypotheses that have a p-value  $\leq 0.033333333333333333$ .  
 Holm's procedure rejects those hypotheses that have a p-value  $\leq 0.1$ .  
 Hochberg's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ .  
 Hommel's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ .

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Table 4: Adjusted p-values

$i$	algorithm	unadjusted $p$	$p_{Bonf}$	$p_{Holm}$	$p_{Hoch}$	$p_{Hommel}$
1	DE	1.3130881465534958E-6	3.9392644396604875E-6	3.9392644396604875E-6	3.9392644396604875E-6	3.9392644396604875E-6
2	VXQR1	1.4566874163845129E-5	4.3700622491535399E-5	2.9133748327690257E-5	2.9133748327690257E-5	2.9133748327690257E-5
3	G-CMA-ES	0.10235752557227855	0.30707237671683563	0.10235752557227855	0.10235752557227855	0.10235752557227855

Table 5: Holm / Shaffer Table for  $\alpha = 0.05$

$i$	algorithms	$z = (R_0 - R_i) / SE$	$p$	Holm	Shaffer
6	DE vs. CHC	4.837762640769081	1.3130881465534958E-6	0.008333333333333333	0.008333333333333333
5	CHC vs. VXQR1	4.335137950819048	1.4566874163845129E-5	0.01	0.016666666666666666
4	DE vs. G-CMA-ES	3.2042323984314676	0.0013542311595120392	0.0125	0.016666666666666666
3	G-CMA-ES vs. VXQR1	2.7016077084814354	0.006900512590003988	0.016666666666666666	0.016666666666666666
2	CHC vs. G-CMA-ES	1.6335302423376128	0.10235752557227855	0.025	0.025
1	DE vs. VXQR1	0.502624689950032	0.6152281670802465	0.05	0.05

Nemenyi's procedure rejects those hypotheses that have a p-value  $\leq 0.008333333333333333$ .

Holm's procedure rejects those hypotheses that have a p-value  $\leq 0.025$ .  
 Shaffer's procedure rejects those hypotheses that have a p-value  $\leq 0.008333333333333333$ .  
 Bergmann's procedure rejects these hypotheses:

- DE vs. CHC
- DE vs. G-CMA-ES
- CHC vs. VXQR1
- G-CMA-ES vs. VXQR1

Table 6: Holm / Shaffer Table for  $\alpha = 0.10$

$i$	algorithms	$z = (R_0 - R_i) / SE$	$p$	Holm	Shaffer
6	DE vs. CHC	4.837762640769081	1.3130881465534958E-6	0.016666666666666666	0.016666666666666666
5	CHC vs. VXQR1	4.335137950819048	1.4566874163845129E-5	0.02	0.033333333333333333
4	DE vs. G-CMA-ES	3.2042323984314676	0.0013542311595120392	0.025	0.033333333333333333
3	G-CMA-ES vs. VXQR1	2.7016077084814354	0.0069000512590003988	0.033333333333333333	0.033333333333333333
2	CHC vs. G-CMA-ES	1.6335302423376128	0.10235752557227855	0.05	0.05
1	DE vs. VXQR1	0.502624689950032	0.6152281670802465	0.1	0.1

Nemenyi's procedure rejects those hypotheses that have a p-value  $\leq 0.008333333333333333$ .  
 Holm's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ .  
 Shaffer's procedure rejects those hypotheses that have a p-value  $\leq 0.016666666666666666$ .  
 Bergmann's procedure rejects these hypotheses:

- DE vs. CHC
- DE vs. G-CMA-ES
- CHC vs. VXQR1
- G-CMA-ES vs. VXQR1

Table 7: Adjusted  $p$ -values

$i$	hypothesis	unadjusted $p$	$p_{Neme}$	$p_{Holm}$	$p_{Shaf}$	$p_{Berq}$
1	DE vs .CHC	1.3130881465534958E-6	7.878528879320975E-6	7.878528879320975E-6	7.878528879320975E-6	7.878528879320975E-6
2	CHC vs .VXQR1	1.4566874163845129E-5	8.740124498307078E-5	7.283437081922564E-5	4.370062249153539E-5	4.370062249153539E-5
3	DE vs .G-CMA-ES	0.0013542311595120392	0.008125386957072236	0.005416924638048157	0.004062693478536118	0.004062693478536118
4	G-CMA-ES vs .VXQR1	0.006900512590003988	0.041403075540023926	0.020701537770011963	0.020701537770011963	0.006900512590003988
5	CHC vs .G-CMA-ES	0.1023575257227855	0.6141451534336713	0.2047150511445571	0.2047150511445571	0.2047150511445571
6	DE vs .VXQR1	0.6152281670802465	3.6913690002481479	0.6152281670802465	0.6152281670802465	0.6152281670802465