Results

April 6, 2021

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

Table 1: Average Rankings of the algorithms

)	Ranking	2.59090909090909	2.136363636363636	1.27272727272727
)	${ m Algorithm}$	ppas	sop pos	svdd desthr

Friedman statistic considering reduction performance (distributed according to chi-square with 2 degrees of freedom: 19.7272727272542. P-value computed by Friedman Test: 5.203282570365175E-5.

Iman and Davenport statistic considering reduction performance (distributed according to F-distribution with 2 and 42 degrees of freedom: 17.067415730336787.

P-value computed by Iman and Daveport Test: 3.758234665599347E-6.

Bonferroni-Dunn's procedure rejects those hypotheses that have a p-value ≤ 0.025 .

Table 2: Holm / Hochberg Table for $\alpha = 0.05$ 1gorithm $z = (R_0 - R_i)/SE$ pHolm/Hochberg/H

Holm/Hochberg/Hommel	0.025	0.05	
d	1.2316176136565982E-5	0.0041785575681665	
$z = (R_0 - R_i)/SE$	4.371914496377571	2.864357773488755	
algorithm	ppas	sydd des	
i	2	П	

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

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

Holm/Hochberg/Hommel	0.05	0.1
d	1.2316176136565982E-5	0.0041785575681665
$z = (R_0 - R_i)/SE$	4.371914496377571	2.864357773488755
algorithm	svdd	sydd des
. 2	7	_

Bonferroni-Dunn's procedure rejects those hypotheses that have a p-value ≤ 0.05 . Hochberg's procedure rejects those hypotheses that have a p-value ≤ 0.1 . Hommel's procedure rejects all hypotheses.

Table 4: Adjusted p-values

p_{Homm}	2.4632352273131964E-5	0.0041785575681665	
p_{Hoch}	2.4632352273131964E-5	0.0041785575681665	
$_{mloHd}$	2.4632352273131964E-5	0.0041785575681665	
p_{Bonf}	2.4632352273131964E-5	0.008357115136333	
unadjusted p	1.2316176136565982E-5	0.0041785575681665	
algorithm	ppvs	sydd des	
·i	1	7	

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

. 2	algorithms	$z = (R_0 - R_i)/SE$	d	Holm	Shaffer
₂	svdd vs. svdd desthr	4.371914496377571	1.2316176136565982E-5	0.01666666666666666	0.016666666666666666
7	svdd des vs. svdd desthr	2.864357773488755	0.0041785575681665	0.025	0.05
_	sydd vs. sydd des	1.5075567228888167	0.13166801602281455	0.05	0.05

Nemenyi's procedure rejects those hypotheses that have a p-value \leq 0.0166666666666666666. 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 0.0166666666666666666. Bergmann's procedure rejects these hypotheses:

- svdd vs. svdd desthr
- sydd des vs. sydd desthr

- svdd vs. svdd desthr
- svdd des vs. svdd desthr

Table 7: Adjusted p-values

			Table 7: Adjuste	a p -values		
	hypothesis	unadjusted p	p_{Neme}	p_{Holm}	pShaf	p_{Berg}
1	svdd vs .svdd desthr	1.2316176136565982E-5	3.694852840969795E-5	3.694852840969795E-5	3.694852840969795E-5	3.694852840969795E-5
01	svdd des vs .svdd desthr	0.0041785575681665	0.012535672704499502	0.008357115136333	0.0041785575681665	0.0041785575681665
က	svdd vs. svdd des	0.13166801602281455	0.3950040480684437	0.13166801602281455	0.13166801602281455	0.13166801602281455