## Results

## April 6, 2021

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

Table 1: Average Rankings of the algorithms
Algorithm

Ranking	2.954545454545453	1.90909090909085	1.1363636363636358
Algorithm	kmeans	kmeans des	kmeans desthr

Friedman statistic considering reduction performance (distributed according to chi-square with 2 degrees of freedom: 36.6363636363333. P-value computed by Friedman Test: 1.1112804454427305E-8. Iman and Davenport statistic considering reduction performance (distributed according to F-distribution with 2 and 42 degrees of freedom: 104.48148148147622.

P-value computed by Iman and Daveport Test: 4.9708745378137235E-17.

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

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

Holm/Hochberg/Hommel	0.025	0.05
<i>a</i>	1.637296451224841E-9	0.010381795789701729
$z = (R_0 - R_i)/SE$	6.030226891555269	2.5628464289109907
algorithm	kmeans	kmeans des
.5	2	-

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	1.637296451224841E-9	0.010381795789701729
$z = (R_0 - R_i)/SE$	6.030226891555269	2.5628464289109907
algorithm	kmeans	kmeans des
i	61	1

Bonferroni-Dunn's procedure rejects those hypotheses that have a p-value  $\leq 0.05$ . Hoch berg's procedure rejects those hypotheses that have a p-value  $\leq 0.1.$  Hommel's procedure rejects all hypotheses.

Table 4: Adjusted p-values

		L	_	l
	pHomm	3.274592902449682E-9	0.010381795789701729	
	$p_{Hoch}$	3.274592902449682E-9	0.010381795789701729	
,	$p_{Holm}$	3.274592902449682E-9	0.010381795789701729	
	pBonf	3.274592902449682E-9	0.020763591579403457	
	unadjusted $p$	1.637296451224841E-9	0.010381795789701729	
	algorithm	kmeans	kmeans des	
		1	7	ĺ

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

		Table 9. House /	DITCHT TOTAL TOT	00:01	
.2	algorithms	$z = (R_0 - R_i)/SE$	d	Holm	Shaffer
8	kmeans vs. kmeans desthr	6.030226891555269	1.637296451224841E-9	0.01666666666666666	0.01666666666666666
7	kmeans vs. kmeans des	3.467380462644278	5.25557438272822E-4	0.025	0.05
_	kmeans des vs. kmeans desthr	2.5628464289109907	0.010381795789701729	0.05	0.05

Bergmann's procedure rejects these hypotheses:

- kmeans vs. kmeans des
- kmeans vs. kmeans desthr
- kmeans des vs. kmeans desthr

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

	ı		
Shaffer	0.03333333333333333	0.1	0.1
Holm	0.033333333333333333	0.05	0.1
d	1.637296451224841E-9	5.255574382728222E-4	0.010381795789701729
$z = (R_0 - R_i)/SE$	6.030226891555269	3.467380462644278	2.5628464289109907
algorithms	kmeans vs. kmeans desthr	kmeans vs. kmeans des	kmeans des vs. kmeans desthr
.2	က	61	1

Bergmann's procedure rejects these hypotheses:

- kmeans vs. kmeans des
- kmeans vs. kmeans desthr
- kmeans des vs. kmeans desthr

Table 7: Adjusted p-values

			Table 1: Aujusteu	b-values		
	hypothesis	unadjusted $p$	$p_{Neme}$	$^{pHolm}$	pShaf	$p_{Berg}$
-	kmeans vs .kmeans desthr	1.637296451224841E-9	4.911889353674523E-9	4.911889353674523E-9	4.911889353674523E-9	4.911889353674523E-9
73	kmeans vs .kmeans des	5.255574382728222E-4	0.0015766723148184665	0.0010511148765456444	5.255574382728222E-4	5.255574382728222E-4
က	kmeans des vs .kmeans desthr	0.010381795789701729	0.031145387369105187	0.010381795789701729	0.010381795789701729	0.010381795789701729