Results

April 5, 2021

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

Table 1: Average Rankings of the algorithms

| Ranking | 2.749999999999999 | 1.99999999999993 | 1.2499999999999996 |
|-----------|-------------------|------------------|--------------------|
| Algorithm | parzen | parzen des | parzen desthr |

Friedman statistic considering reduction performance (distributed according to chi-square with 2 degrees of freedom: 24.749999999905. P-value computed by Friedman Test: 4.2228836869551145E-6.

Iman and Davenport statistic considering reduction performance (distributed according to F-distribution with 2 and 42 degrees of freedom: 26.9999999999513.

P-value computed by Iman and Daveport Test: 2.8876144374466767E-8.

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

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

| ·z | algorithm | $z = (R_0 - R_i)/SE$ | d | Holm/Hochberg/Hommel |
|----|------------|----------------------|----------------------|----------------------|
| 2 | parzen | 4.974937185533098 | 6.526880098667021E-7 | 0.025 |
| | parzen des | 2.487468592766549 | 0.01286558129455608 | 0.05 |

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 | 6.526880098667021E-7 | 0.01286558129455608 | |
| $z = (R_0 - R_i)/SE$ | 4.974937185533098 | 2.487468592766549 | |
| algorithm | parzen | parzen des | |
| . 2 | 21 | _ | |

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

| | | | I months is organi | and I have | | |
|---|------------|----------------------|-----------------------|-----------------------|-----------------------|-----------------------|
| | algorithm | unadjusted p | p_{Bonf} | pHolm | p_{Hoch} | p_{Homm} |
| 1 | parzen | 6.526880098667021E-7 | 1.3053760197334043E-6 | 1.3053760197334043E-6 | 1.3053760197334043E-6 | 1.3053760197334043E-6 |
| 7 | parzen des | 0.01286558129455608 | 0.02573116258911216 | 0.01286558129455608 | 0.01286558129455608 | 0.01286558129455608 |
| | | | | | | |

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

| | | TOTAL OF TROUTE | DIRECT TRDIC TOT C | 00:0 2 | |
|---|------------------------------|----------------------|----------------------|---------------------|----------------------|
| i | algorithms | $z = (R_0 - R_i)/SE$ | d | Holm | Shaffer |
| m | parzen vs. parzen desthr | 4.974937185533098 | 6.526880098667021E-7 | 0.01666666666666666 | 0.016666666666666666 |
| 2 | parzen vs. parzen des | 2.487468592766549 | 0.01286558129455608 | 0.025 | 0.05 |
| - | parzen des vs. parzen desthr | 2.487468592766549 | 0.01286558129455608 | 0.05 | 0.05 |

- parzen vs. parzen des
- parzen vs. parzen desthr
- parzen des vs. parzen desthr

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

| Shaffer | 33333 0.0333333333333333 | 0.1 | 0.1 |
|----------------------|--------------------------|-----------------------|------------------------------|
| Holm | 0.033333333333 | 0.02 | 0.1 |
| d | 6.526880098667021E-7 | 0.01286558129455608 | 0.01286558129455608 |
| $z = (R_0 - R_i)/SE$ | 4.974937185533098 | 2.487468592766549 | 2.487468592766549 |
| algorithms | parzen vs. parzen desthr | parzen vs. parzen des | parzen des vs. parzen desthr |
| ·z | က | 7 | |

Bergmann's procedure rejects these hypotheses:

- parzen vs. parzen des
- $\bullet\,$ parzen vs. parzen desthr
- $\bullet\,$ parzen des vs. parzen desthr

| | | | Table 7: Adjusted p -v | p-values | | |
|---|------------------------------|----------------------|--------------------------|-----------------------|-----------------------|-----------------------|
| | hypothesis | unadjusted p | pNeme | p_{Holm} | pShaf | pBerg |
| - | parzen vs .parzen desthr | 6.526880098667021E-7 | 1.9580640296001065E-6 | 1.9580640296001065E-6 | 1.9580640296001065E-6 | 1.9580640296001065E-6 |
| 7 | parzen vs .parzen des | 0.01286558129455608 | 0.03859674388366824 | 0.02573116258911216 | 0.01286558129455608 | 0.01286558129455608 |
| 8 | parzen des vs .parzen desthr | 0.01286558129455608 | 0.03859674388366824 | 0.02573116258911216 | 0.01286558129455608 | 0.01286558129455608 |