



Machine Learning & Data Mining Coursework

Module code: 5DATA002W

Module leader: Dr. V.S. Kontogiannis

Assignment: Coursework

Handed Out: 20/02/2023.

Due Date: 04/05/2022, Submission by 13:00

Name: B.K. Praveen Navanjana

IIT ID: 20200877

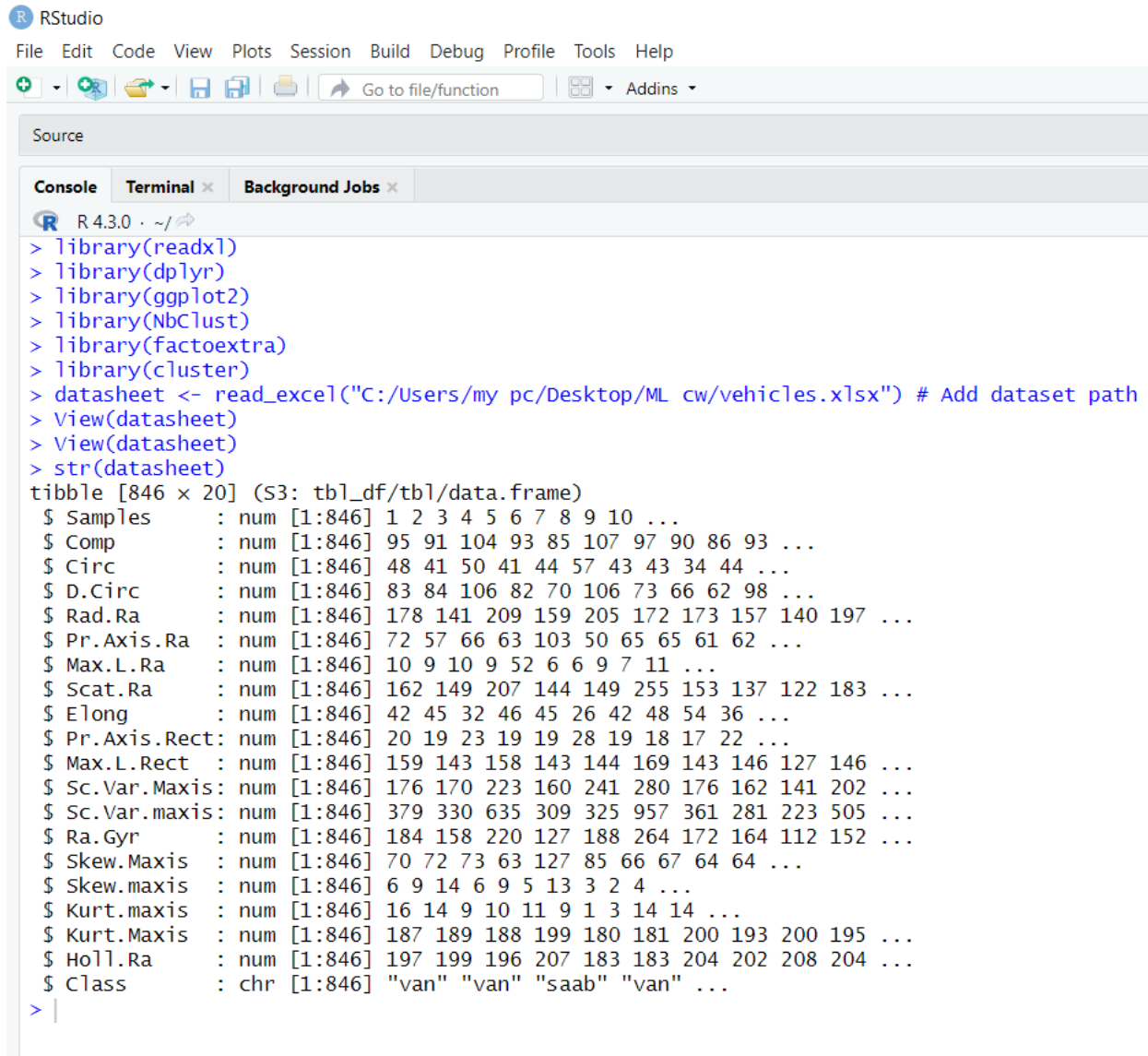
UOW ID: w1870590

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Partitioning Clustering Part

Dataset loading and checking the dataset



The screenshot shows the RStudio interface with the following components:

- Menu Bar:** File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, Help.
- Toolbar:** Includes icons for opening files, saving, and a search bar labeled "Go to file/function".
- Source Panel:** Currently empty.
- Console Panel:**
 - Header: R 4.3.0 · ~/
 - Commands entered:


```
> library(readxl)
> library(dplyr)
> library(ggplot2)
> library(NbClust)
> library(factoextra)
> library(cluster)
> datasheet <- read_excel("C:/Users/my pc/Desktop/ML cw/vehicles.xlsx") # Add dataset path
> View(datasheet)
> View(datasheet)
> str(datasheet)
```
 - Output:


```
tibble [846 × 20] (S3: tbl_df/tbl/data.frame)
 $ Samples      : num [1:846] 1 2 3 4 5 6 7 8 9 10 ...
 $ Comp         : num [1:846] 95 91 104 93 85 107 97 90 86 93 ...
 $ Circ         : num [1:846] 48 41 50 41 44 57 43 43 34 44 ...
 $ D.Circ       : num [1:846] 83 84 106 82 70 106 73 66 62 98 ...
 $ Rad.Ra       : num [1:846] 178 141 209 159 205 172 173 157 140 197 ...
 $ Pr.Axis.Ra   : num [1:846] 72 57 66 63 103 50 65 65 61 62 ...
 $ Max.L.Ra     : num [1:846] 10 9 10 9 52 6 6 9 7 11 ...
 $ Scat.Ra      : num [1:846] 162 149 207 144 149 255 153 137 122 183 ...
 $ Elong        : num [1:846] 42 45 32 46 45 26 42 48 54 36 ...
 $ Pr.Axis.Rect : num [1:846] 20 19 23 19 19 28 19 18 17 22 ...
 $ Max.L.Rect   : num [1:846] 159 143 158 143 144 169 143 146 127 146 ...
 $ Sc.Var.Maxis : num [1:846] 176 170 223 160 241 280 176 162 141 202 ...
 $ Sc.Var.maxis : num [1:846] 379 330 635 309 325 957 361 281 223 505 ...
 $ Ra.Gyr       : num [1:846] 184 158 220 127 188 264 172 164 112 152 ...
 $ Skew.Maxis   : num [1:846] 70 72 73 63 127 85 66 67 64 64 ...
 $ Skew.maxis   : num [1:846] 6 9 14 6 9 5 13 3 2 4 ...
 $ Kurt.maxis   : num [1:846] 16 14 9 10 11 9 1 3 14 14 ...
 $ Kurt.Maxis   : num [1:846] 187 189 188 199 180 181 200 193 200 195 ...
 $ Holl.Ra      : num [1:846] 197 199 196 207 183 183 204 202 208 204 ...
 $ Class        : chr [1:846] "van" "van" "saab" "van" ...
```

First 18 columns

```
> data_c
# A tibble: 846 × 18
  Comp Circ D.Circ Rad.Ra Pr.Axis.Ra Max.L.Ra Scat.Ra Elong Pr.Axis.Rect Max.L.Rect
  <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1    95    48    83    178    72    10    162    42    20    159
2    91    41    84    141    57     9    149    45    19    143
3   104    50   106    209    66    10    207    32    23    158
4    93    41    82    159    63     9    144    46    19    143
5    85    44    70    205   103    52    149    45    19    144
6   107    57   106    172    50     6    255    26    28    169
7    97    43    73    173    65     6    153    42    19    143
8    90    43    66    157    65     9    137    48    18    146
9    86    34    62    140    61     7    122    54    17    127
10   93    44    98    197    62    11    183    36    22    146
# i 836 more rows
# i 8 more variables: Sc.Var.Maxis <dbl>, Sc.Var.maxis <dbl>, Ra.Gyr <dbl>, Skew.Maxis <dbl>,
#   Skew.maxis <dbl>, Kurt.maxis <dbl>, Kurt.Maxis <dbl>, Holl.Ra <dbl>
# i Use `print(n = ...)` to see more rows
> |
```

Preprocessing tasks

Scaling

Scaling is a crucial component of machine learning because it ensures that all variables are handled similarly during the learning process, regardless of their original values or measurement units. We can improve the performance and speed of some machine learning algorithms and produce models that are more accurate and effective by scaling the features.

```

> Data_N_outliers <- data_c
> md <- mahalanobis(data_c, colMeans(data_c), cov(data_c))
> outliers <- which(md > qchisq(0.95, df = ncol(data_c)))
> Data_N_outliers <- Data_N_outliers[-outliers, ]
> D_Scaled <- scale(data_c)
> D_Scaled

```

	Comp	Circ	D.Circ	Rad.Ra	Pr.Axis.Ra	Max.L.Ra	Scat.Ra
[1,]	0.16048541	0.50864931	0.057784334	0.270645678	1.30651856	0.31135767	-0.205722605
[2,]	-0.32527723	-0.62589728	0.121189713	-0.834749808	-0.59504361	0.09402385	-0.596759110
[3,]	1.25345137	0.83280548	1.516108039	1.196787842	0.54589369	0.31135767	1.147865294
[4,]	-0.08239591	-0.62589728	-0.005621044	-0.296989842	0.16558126	0.09402385	-0.747157765
[5,]	-1.05392120	-0.13966303	-0.766485586	1.077285627	5.23641371	9.43937817	-0.596759110
[6,]	1.61777335	1.96735207	1.516108039	0.091392356	-1.48243929	-0.55797761	2.591692388
[7,]	0.40336674	-0.30174112	-0.576269450	0.121267909	0.41912288	-0.55797761	-0.476440185
[8,]	-0.44671789	-0.30174112	-1.020107099	-0.356740949	0.41912288	0.09402385	-0.957715883
[9,]	-0.93248054	-1.76044388	-1.273728613	-0.864625362	-0.08796036	-0.34064379	-1.408911850
[10,]	-0.08239591	-0.13966303	1.008865011	0.838281197	0.03881045	0.52869149	0.425951748
[11,]	-0.93248054	-1.43628771	-0.766485586	-0.774998701	-0.08796036	0.09402385	-1.078034808
[12,]	-0.44671789	-1.76044388	-1.020107099	-0.984127577	-0.84858523	-0.55797761	-1.378832119
[13,]	-0.68959922	0.18449314	-0.512864072	0.061516802	0.79943532	-0.55797761	-0.506519916
[14,]	-0.56815856	-0.46381920	0.184595091	-0.745123147	-0.46827280	0.31135767	-0.506519916
[15,]	0.03904475	0.67072739	-0.195837180	1.017534519	1.17974775	-0.77531143	0.155234168
[16,]	0.28192607	1.64319590	1.325891903	0.957783412	0.41912288	0.09402385	1.057626101
[17,]	-0.56815856	-1.43628771	-1.971187776	-1.790767526	-1.22889766	-0.55797761	-1.529230774
[18,]	0.64624806	-0.62589728	-0.322647936	0.838281197	0.92620613	-0.55797761	0.245473361
[19,]	1.25345137	1.48111782	1.135675768	0.509650107	-0.08796036	0.31135767	1.418582874
[20,]	0.88912938	1.80527399	1.135675768	1.376041164	0.92620613	0.31135767	1.177945026
[21,]	-1.17536186	0.34657122	-0.449458693	-0.476243164	0.29235207	-0.55797761	-0.446360454
[22,]	-1.17536186	-1.27420962	-1.844377019	-1.432260882	-0.34150198	-0.77531143	-1.378832119
[23,]	0.03904475	-0.30174112	-1.146917856	0.121267909	0.92620613	-0.34064379	-0.566679379
[24,]	-0.81103988	-0.95005345	-0.766485586	-0.625620932	-0.08796036	-0.34064379	-0.777237496
[25,]	0.64624806	1.31903973	1.452702660	1.495543378	0.54589369	0.52869149	1.057626101
[26,]	-1.05392120	0.02241505	-0.132431801	-0.446367610	0.29235207	0.09402385	-0.656918572
[27,]	-1.29680252	-1.43628771	-1.780971641	-1.492011989	-0.59504361	-0.55797761	-1.228433463
[28,]	1.61777335	1.48111782	1.008865011	1.017534519	0.41912288	0.52869149	1.478742337
[29,]	1.01057004	0.02241505	0.184595091	0.718778983	0.29235207	-0.55797761	0.696669328
[30,]	-1.66112451	-1.11213154	-1.210323235	-1.193256452	-0.84858523	-0.34064379	-0.686998303
[31,]	-0.56815856	-0.30174112	0.184595091	-0.267114288	0.29235207	0.52869149	-0.416280723
[32,]	-0.68959922	-0.46381920	-0.322647936	-0.535994271	-0.46827280	-0.12330997	-0.867476690
[33,]	-0.08239591	-1.59836579	-1.020107099	-0.446367610	-0.34150198	-0.55797761	-0.807317228
[34,]	0.88912938	0.50864931	1.579513417	1.585170039	0.79943532	0.31135767	1.177945026

```

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[37,] -1.41824319 -0.13966303 -0.639674829 -1.521887543 -1.22889766 -0.34064379 -0.506519916
[38,] -0.44671789 0.50864931 0.248000470 4.094716548 8.15214237 8.78737671 -0.476440185
[39,] 1.49633269 1.31903973 1.008865011 0.210894570 -0.97535604 0.31135767 1.418582874
[40,] -1.53968385 0.02241505 -0.893296343 0.001765695 1.43328937 -0.55797761 -0.536599648
[41,] 0.16048541 0.50864931 1.389297282 1.346165610 0.67266451 0.09402385 1.087705832
[42,] -0.68959922 -1.27420962 -1.971187776 -1.910269740 -1.22889766 -0.77531143 -1.499151043
[43,] 0.03904475 0.67072739 0.311405848 -0.954252023 -0.97535604 0.52869149 -0.326041530
[44,] -0.08239591 -1.27420962 -0.386053315 0.420023446 0.16558126 -0.12330997 -0.145563143
[45,] 3.07506129 1.48111782 1.516108039 1.525418932 0.41912288 0.74602532 1.328343681
[46,] -0.08239591 0.18449314 -0.005621044 -0.715247593 -0.46827280 0.52869149 -0.295961799
[47,] -0.32527723 -0.30174112 -0.766485586 -1.073754238 -0.84858523 -0.12330997 -1.168274001
[48,] -1.05392120 -0.46381920 -1.020107099 -1.402385328 -0.97535604 -0.55797761 -0.626838841
[49,] -0.56815856 0.34657122 -0.069026423 -0.655496486 0.29235207 0.52869149 -0.386200992
[50,] -0.32527723 0.02241505 -0.195837180 0.210894570 -0.34150198 0.09402385 -0.175642874
[51,] -1.90400583 -1.11213154 -1.210323235 -1.611514204 -1.35566847 -0.55797761 -0.807317228
[52,] -0.20383657 -1.11213154 -0.703080207 0.151143463 0.54589369 -0.34064379 -0.446360454
[53,] 0.52480740 1.64319590 1.199081146 1.764423361 1.05297694 0.09402385 1.238104488
[54,] 0.88912938 -0.46381920 -1.273728613 0.181019017 0.67266451 -0.55797761 -0.596759110
[55,] 0.88912938 1.80527399 1.389297282 0.479774553 -1.10212685 -0.55797761 2.651851850
      Elong Pr.Axis.Rect Max.L.Rect Sc.Var.Maxis Sc.Var.maxis Ra.Gyr
[1,] 0.136489241 -0.2248114 7.578841e-01 -0.40214560 -0.3447305820 0.285643409
[2,] 0.520535463 -0.6105933 -3.443743e-01 -0.59325983 -0.6220483430 -0.513213860
[3,] -1.143664834 0.9325342 6.889930e-01 1.09491586 1.1041132308 1.391753474
[4,] 0.648550870 -0.6105933 -3.443743e-01 -0.91178355 -0.7408988120 -1.465697528
[5,] 0.520535463 -0.6105933 -2.754832e-01 1.66825855 -0.6503460737 0.408544527
[6,] -1.911757278 2.8614436 1.446796e+00 2.91050104 2.9264870891 2.743665776
[7,] 0.136489241 -0.6105933 -3.443743e-01 -0.40214560 -0.4466024125 -0.083059946
[8,] 0.904581685 -0.9963752 -1.377009e-01 -0.84807881 -0.8993661040 -0.328862183
[9,] 1.672674129 -1.3821571 -1.446633e+00 -1.51697861 -1.2276197804 -1.926576722
[10,] -0.631603204 0.5467523 -1.377009e-01 0.42601606 0.3683722322 -0.697565538
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[14,] 0.392520055 -0.6105933 -2.754832e-01 -0.49770272 -0.5371551508 -0.421038022
[15,] -0.503587796 0.1609705 4.134284e-01 0.23490183 0.1419903864 0.961599560
[16,] -1.143664834 0.9325342 1.240122e+00 1.22232535 1.0418582233 2.190610744
[17,] 2.056720352 -1.3821571 -1.308851e+00 -1.64438810 -1.3238320648 -1.527148087
[18,] -0.631603204 0.1609705 -6.199390e-01 0.42601606 0.2551813093 -0.728290818
[19,] -1.271680241 1.3183161 1.722360e+00 1.15862060 1.3927500842 1.391753474
[20,] -1.143664834 1.3183161 1.446796e+00 1.22232535 1.1946659691 1.483929313
[21,] 0.264504648 -0.6105933 -2.065920e-01 -0.43399798 -0.4862192355 0.285643409
[22,] 1.800680537 -1.3821571 -1.584415e+00 -1.51697861 -1.2380388727 -1.281245850

```


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[25,] -1.143664834 0.9325342 1.171231e+00 1.03121112 1.0361986771 1.514654593
[26,] 0.520535463 -0.6105933 8.143163e-05 -0.62511221 -0.6560056199 -0.021609387
[27,] 1.544658722 -0.9963752 -1.584415e+00 -1.45327387 -1.1427265882 -1.096994173
[28,] -1.271680241 1.7040980 1.309013e+00 1.28603009 1.4493455456 1.268852356
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[30,] 0.648550870 -0.6105933 -1.239959e+00 -0.65696458 -0.7126010813 -0.513213860
[31,] 0.264504648 -0.6105933 2.067549e-01 -0.49770272 -0.4749001433 -0.021609387
[32,] 0.776566278 -0.9963752 -4.132655e-01 -0.75252169 -0.8314515503 -0.513213860
[33,] 0.648550870 -0.9963752 -1.377742e+00 -0.84807881 -0.7691965427 -1.680774485
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[35,] -0.119541574 -0.2248114 -1.308851e+00 -0.08362189 -0.2145610206 -1.373521689
[36,] 0.264504648 -0.2248114 1.378637e-01 -0.59325983 -0.4352833203 0.285643409
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[39,] -1.271680241 1.3183161 1.584578e+00 1.47714432 1.4210478149 1.330302915
[40,] 0.392520055 -0.6105933 -1.377009e-01 -0.49770272 -0.5880910661 0.347093968
[41,] -1.143664834 0.9325342 2.067549e-01 1.22232535 1.0644964078 0.838698442
[42,] 2.056720352 -1.3821571 -1.377742e+00 -1.70809284 -1.3181725187 -1.527148087
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[49,] 0.392520055 -0.2248114 1.033449e+00 -0.59325983 -0.4975383278 0.408544527
[50,] -0.119541574 -0.2248114 8.143163e-05 -0.14732663 -0.2032419284 0.132017011
[51,] 0.776566278 -0.6105933 -1.239959e+00 -0.84807881 -0.7974942735 -0.881917215
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[53,] -1.271680241 1.3183161 1.377905e+00 1.50899669 1.2512614306 2.159885464
[54,] 0.264504648 -0.6105933 -6.199390e-01 -0.62511221 -0.5597933354 -0.298136903
[55,] -1.911757278 2.8614436 1.377905e+00 2.75123918 2.9208275430 1.699006270
Skew.Maxis Skew.maxis Kurt.maxis Kurt.Maxis Ho||.Ra
[1,] -0.32886116 -0.07666562 0.38076562 -0.31353666 0.18384857
[2,] -0.06173054 0.53329468 0.15683255 0.01093064 0.45270923
[3,] 0.07183477 1.54989517 -0.40300011 -0.15130301 0.04941824
[4,] -1.26381831 -0.07666562 -0.29103357 1.63326713 1.52815188
[5,] 7.28436143 0.53329468 -0.17906704 -1.44917221 -1.69817606
[6,] 1.67461847 -0.27998571 -0.40300011 -1.28693856 -1.69817606
[7,] -0.86312239 1.34657507 -1.29873236 1.79550078 1.12486089
[8,] -0.72955708 -0.68662591 -1.07479930 0.65986523 0.85600023
[9,] -1.13025301 -0.88994601 0.15683255 1.79550078 1.66258221

```

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```

[13,] -0.19529585 -0.27998571 -0.17906704 0.01093064 -0.08501209
[14,] -0.06173054 0.32997458 0.04486602 -0.31353666 0.18384857
[15,] -0.19529585 -0.07666562 -1.18676583 1.30879983 0.45270923
[16,] 0.20540008 -0.07666562 -1.18676583 -0.47577031 -0.21944242
[17,] 1.00679193 -0.88994601 0.15683255 -1.28693856 -1.42931540
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[24,] -0.46242646 -1.09326611 -1.18676583 0.49763158 0.45270923
[25,] -0.59599177 -1.29658621 -0.73889970 0.33539794 0.72156990
[26,] -0.19529585 -1.09326611 -0.96283277 -0.15130301 0.45270923
[27,] 1.27392255 -0.07666562 -1.07479930 -1.61140586 -1.69817606
[28,] -0.06173054 -1.09326611 1.72436400 -0.31353666 0.45270923
[29,] 0.47253070 -0.07666562 -0.62693317 0.98433253 -0.35387275
[30,] 1.40748785 0.53329468 0.82863175 -1.44917221 -1.42931540
[31,] -0.06173054 -0.27998571 -0.40300011 -0.63800396 0.04941824
[32,] -1.13025301 0.73661478 -0.17906704 1.47103348 1.25929122
[33,] -1.13025301 -0.27998571 0.04486602 1.30879983 0.85600023
[34,] -0.32886116 -0.27998571 2.84402932 0.17316429 0.85600023
[35,] -1.26381831 -1.09326611 1.38846441 1.47103348 1.25929122
[36,] -0.72955708 2.15985547 -0.62693317 0.49763158 0.58713957
[37,] 1.27392255 -0.88994601 -1.18676583 -1.44917221 -1.42931540
[38,] 6.08227365 -1.29658621 0.26879909 -0.63800396 -0.21944242
[39,] 0.20540008 -1.09326611 -0.40300011 -0.31353666 0.18384857
[40,] 0.33896539 0.12665448 -1.41069890 -0.96247126 -0.89159407
[41,] 0.20540008 -0.27998571 -0.40300011 -0.47577031 -0.35387275
[42,] 1.80818378 0.32997458 0.38076562 -1.61140586 -1.69817606
[43,] 0.33896539 -0.27998571 -0.85086623 -0.96247126 -0.21944242
[44,] -0.72955708 -0.48330581 -0.62693317 0.49763158 0.18384857
[45,] -0.86312239 2.76981576 -1.29873236 0.49763158 0.85600023
[46,] 0.60609600 -0.88994601 -0.96283277 -0.96247126 -0.21944242
[47,] -0.32886116 -1.09326611 -0.51496664 0.17316429 -0.21944242
[48,] 2.07531440 -0.07666562 0.15683255 -1.44917221 -1.83260639
[49,] 0.47253070 -0.07666562 0.04486602 -0.80023761 -0.35387275
[50,] -1.39738362 -1.29658621 -0.29103357 1.63326713 1.66258221
[51,] 0.60609600 -0.88994601 -0.96283277 -1.28693856 -1.42931540
[52,] -0.32886116 -0.48330581 1.27649787 0.01093064 -0.08501209
[53,] -0.06173054 -1.09326611 -0.73889970 -0.15130301 0.18384857

```

```

[ reached getOption("max.print") -- omitted 791 rows ]

```

```

attr(,"scaled:center")

```

Comp	Circ	D.Circ	Rad.Ra	Pr.Axis.Ra	Max.L.Ra	Scat.Ra
93.678487	44.861702	82.088652	168.940898	61.693853	8.567376	168.839243
Elong	Pr.Axis.Rect	Max.L.Rect	Sc.Var.Maxis	Sc.Var.maxis	Ra.Gyr	Skew.Maxis
40.933806	20.582742	147.998818	188.625296	439.911348	174.703310	72.462175
Skew.maxis	Kurt.maxis	Kurt.Maxis	Holl.Ra			
6.377069	12.599291	188.932624	195.632388			

```

attr(,"scaled:scale")

```

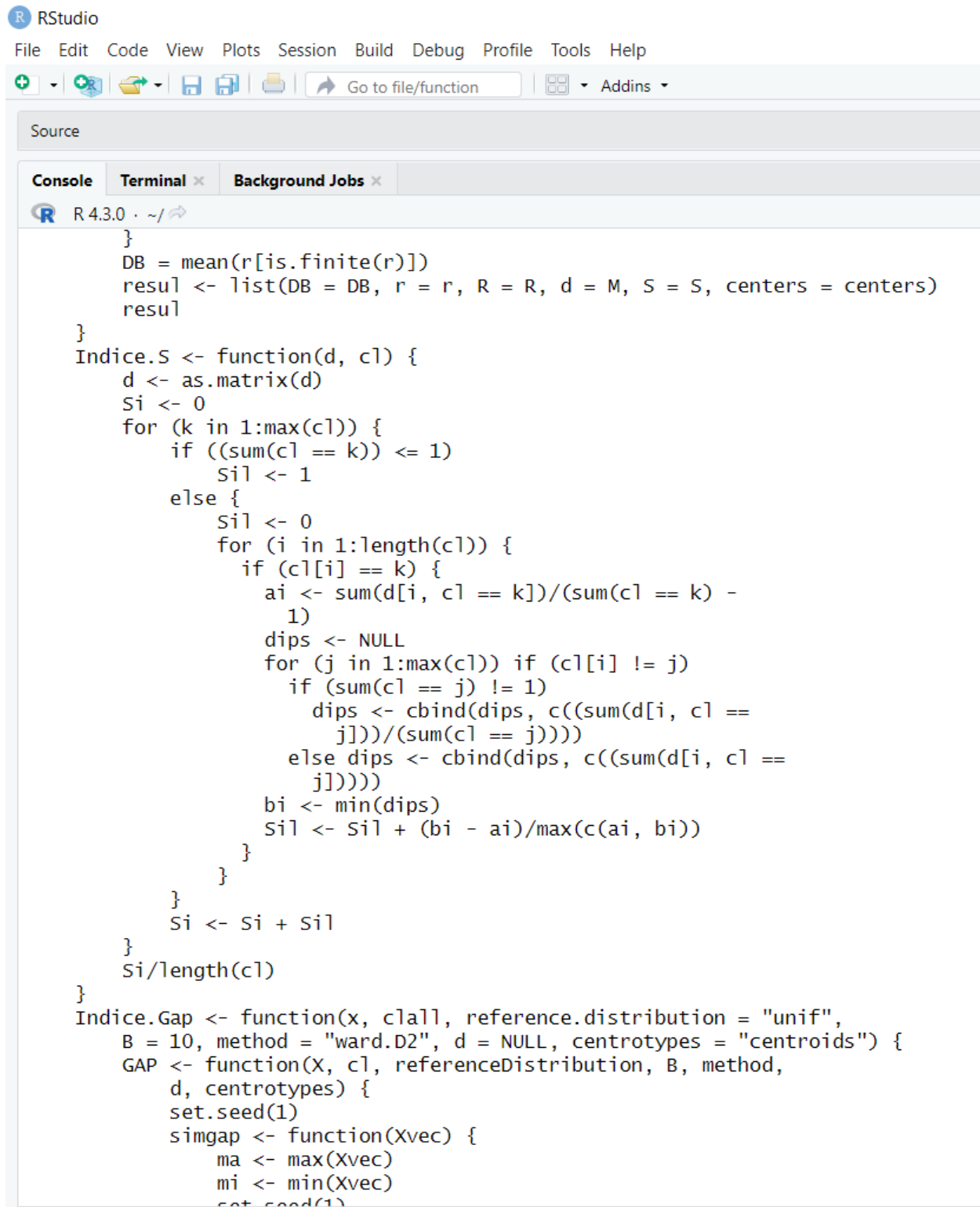
Comp	Circ	D.Circ	Rad.Ra	Pr.Axis.Ra	Max.L.Ra	Scat.Ra
8.234474	6.169866	15.771533	33.472183	7.888251	4.601217	33.244978
Elong	Pr.Axis.Rect	Max.L.Rect	Sc.Var.Maxis	Sc.Var.maxis	Ra.Gyr	Skew.Maxis
7.811560	2.592138	14.515652	31.394837	176.692614	32.546490	7.486974
Skew.maxis	Kurt.maxis	Kurt.Maxis	Holl.Ra			
4.918353	8.931240	6.163949	7.438797			

```

> |

```


Determining the number of clusters using four automated tools



The screenshot shows the RStudio interface with the following components:

- Source Editor:** Contains R code for determining the number of clusters using four automated tools.
- Console:** Shows the R version (R 4.3.0) and the current directory (~/).
- Terminal:** Closed.
- Background Jobs:** Closed.

The R code in the Source Editor is as follows:

```

}
DB = mean(r[is.finite(r)])
resul <- list(DB = DB, r = r, R = R, d = M, S = S, centers = centers)
resul
}
Indice.S <- function(d, cl) {
  d <- as.matrix(d)
  Si <- 0
  for (k in 1:max(cl)) {
    if ((sum(cl == k)) <= 1)
      Si1 <- 1
    else {
      Si1 <- 0
      for (i in 1:length(cl)) {
        if (cl[i] == k) {
          ai <- sum(d[i, cl == k])/(sum(cl == k) -
            1)
          dips <- NULL
          for (j in 1:max(cl)) if (cl[i] != j)
            if (sum(cl == j) != 1)
              dips <- cbind(dips, c((sum(d[i, cl ==
                j]))/(sum(cl == j))))
            else dips <- cbind(dips, c((sum(d[i, cl ==
              j]))))
          bi <- min(dips)
          Si1 <- Si1 + (bi - ai)/max(c(ai, bi))
        }
      }
    }
    Si <- Si + Si1
  }
  Si/length(cl)
}
Indice.Gap <- function(x, clall, reference.distribution = "unif",
  B = 10, method = "ward.D2", d = NULL, centrotypes = "centroids") {
  GAP <- function(X, cl, referenceDistribution, B, method,
    d, centrotypes) {
    set.seed(1)
    simgap <- function(Xvec) {
      ma <- max(Xvec)
      mi <- min(Xvec)
      set.seed(1)

```

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

Console Terminal Background Jobs

```

R 4.3.0 ~/  

set.seed(1)  

simgap <- function(Xvec) {  
  ma <- max(Xvec)  
  mi <- min(Xvec)  
  set.seed(1)  
  Xout <- runif(length(Xvec), min = mi, max = ma)  
  return(Xout)  
}  

pcsim <- function(X, d, centrotypes) {  
  if (centrotypes == "centroids") {  
    Xmm <- apply(X, 2, mean)  
  }  
  for (k in (1:dim(X)[2])) {  
    X[, k] <- X[, k] - Xmm[k]  
  }  
  ss <- svd(X)  
  Xs <- X %*% ss$v  
  Xnew <- apply(Xs, 2, simgap)  
  Xt <- Xnew %*% t(ss$v)  
  for (k in (1:dim(X)[2])) {  
    Xt[, k] <- Xt[, k] + Xmm[k]  
  }  
  return(Xt)  
}  

if (is.null(dim(x))) {  
  dim(x) <- c(length(x), 1)  
}  

ClassNr <- max(c1)  

wk0 <- 0  

wkB <- matrix(0, 1, B)  

for (bb in (1:B)) {  
  if (reference.distribution == "unif")  
    Xnew <- apply(X, 2, simgap)  
  else if (reference.distribution == "pc")  
    Xnew <- pcsim(X, d, centrotypes)  
  else stop("Wrong reference distribution type")  
  if (bb == 1) {  
    pp <- c1  
    if (ClassNr == length(c1))  
      pp2 <- 1:ClassNr  
    else if (method == "k-means") {  
      set.seed(1)

```

RStudio

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Go to file/function Addins

Source

Console Terminal x Background Jobs x

R 4.3.0 · ~/

```

if (ClassNr == length(c1))
  pp2 <- 1:ClassNr
else if (method == "k-means") {
  set.seed(1)
  pp2 <- kmeans(Xnew, ClassNr, 100)$cluster
}
else if (method == "single" || method == "complete" ||
  method == "average" || method == "ward.D2" ||
  method == "mcquitty" || method == "median" ||
  method == "centroid" || method == "ward.D")
  pp2 <- cutree(hclust(dist(Xnew), method = method),
    ClassNr)
else stop("Wrong clustering method")
if (ClassNr > 1) {
  for (zz in (1:ClassNr)) {
    Xuse <- X[pp == zz, ]
    wk0 <- wk0 + sum(diag(var(Xuse))) * (length(pp[pp ==
      zz]) - 1)/(dim(X)[1] - ClassNr)
    Xuse2 <- Xnew[pp2 == zz, ]
    wkB[1, bb] <- wkB[1, bb] + sum(diag(var(Xuse2))) *
      (length(pp2[pp2 == zz]) - 1)/(dim(X)[1] -
        ClassNr)
  }
}
if (ClassNr == 1) {
  wk0 <- sum(diag(var(X)))
  wkB[1, bb] <- sum(diag(var(Xnew)))
}
}
if (bb > 1) {
  if (ClassNr == length(c1))
    pp2 <- 1:ClassNr
  else if (method == "k-means") {
    set.seed(1)
    pp2 <- kmeans(Xnew, ClassNr, 100)$cluster
  }
  else if (method == "single" || method == "complete" ||
    method == "average" || method == "ward.D2" ||
    method == "mcquitty" || method == "median" ||
    method == "centroid" || method == "ward.D")
    pp2 <- cutree(hclust(dist(Xnew), method = method),
      ClassNr)
}

```

RStudio

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Go to file/function Addins

Source

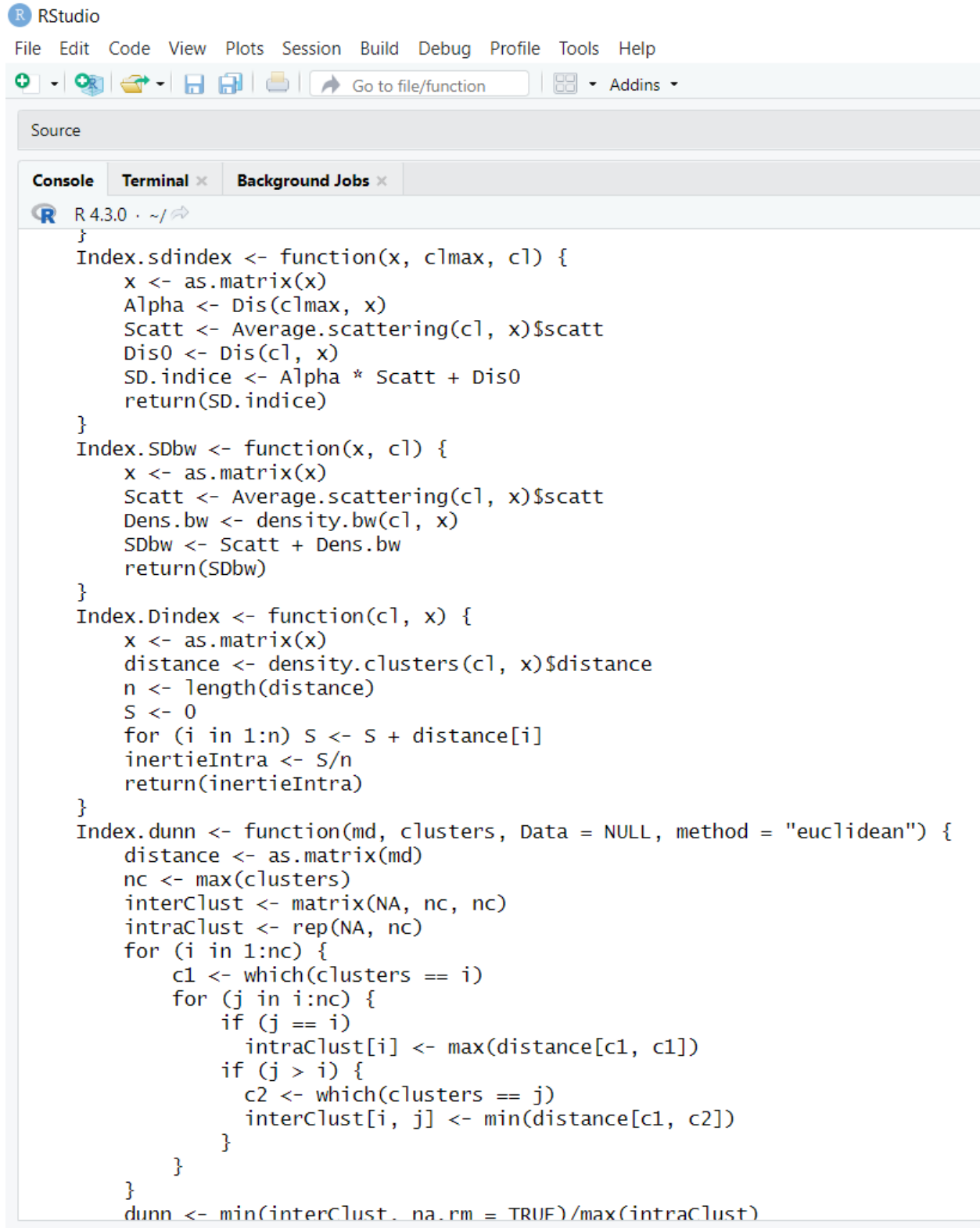
Console Terminal Background Jobs

R 4.3.0 · ~/

```

      ClassNr)
    else stop("Wrong clustering method")
    if (ClassNr > 1) {
      for (zz in (1:ClassNr)) {
        Xuse2 <- Xnew[pp2 == zz, ]
        wkB[1, bb] <- wkB[1, bb] + sum(diag(var(Xuse2))) *
          length(pp2[pp2 == zz])/(dim(X)[1] - ClassNr)
      }
    }
    if (ClassNr == 1) {
      wkB[1, bb] <- sum(diag(var(Xnew)))
    }
  }
  Sgap <- mean(log(wkB[1, ])) - log(wk0)
  Sdgap <- sqrt(1 + 1/B) * sqrt(var(log(wkB[1, ]))) *
    sqrt((B - 1)/B)
  resul <- list(Sgap = Sgap, Sdgap = Sdgap)
  resul
}
if (sum(c("centroids", "medoids") == centrotypes) ==
  0)
  stop("Wrong centrotypes argument")
if ("medoids" == centrotypes && is.null(d))
  stop("For argument centrotypes = 'medoids' d can not be null")
if (!is.null(d)) {
  if (!is.matrix(d)) {
    d <- as.matrix(d)
  }
  row.names(d) <- row.names(x)
}
X <- as.matrix(x)
gap1 <- GAP(X, clall[, 1], reference.distribution, B,
  method, d, centrotypes)
gap <- gap1$Sgap
gap2 <- GAP(X, clall[, 2], reference.distribution, B,
  method, d, centrotypes)
diffu <- gap - (gap2$Sgap - gap2$Sdgap)
resul <- list(gap = gap, diffu = diffu)
resul
}
Index sdindex <- function(x clmax cl) {

```



RStudio

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Go to file/function Addins

Source

Console Terminal Background Jobs

R 4.3.0 · ~/

```

    }
    for (nc in min_nc:max_nc) {
      if (any(method == 1) || (method == 2) || (method == 3) ||
          (method == 4) || (method == 5) || (method == 6) ||
          (method == 7) || (method == 9)) {
        cl1 <- cutree(hc, k = nc)
        cl2 <- cutree(hc, k = nc + 1)
        clall <- cbind(cl1, cl2)
        clmax <- cutree(hc, k = max_nc)
        if (nc >= 2) {
          cl0 <- cutree(hc, k = nc - 1)
          clall1 <- cbind(cl0, cl1, cl2)
        }
        if (nc == 1) {
          cl0 <- rep(NA, nn)
          clall1 <- cbind(cl0, cl1, cl2)
        }
      }
      if (method == 8) {
        set.seed(1)
        cl2 <- kmeans(jeu, nc + 1)$cluster
        set.seed(1)
        clmax <- kmeans(jeu, max_nc)$cluster
        if (nc > 2) {
          set.seed(1)
          cl1 <- kmeans(jeu, nc)$cluster
          clall <- cbind(cl1, cl2)
          set.seed(1)
          cl0 <- kmeans(jeu, nc - 1)$cluster
          clall1 <- cbind(cl0, cl1, cl2)
        }
        if (nc == 2) {
          set.seed(1)
          cl1 <- kmeans(jeu, nc)$cluster
          clall <- cbind(cl1, cl2)
          cl0 <- rep(1, nn)
          clall1 <- cbind(cl0, cl1, cl2)
        }
        if (nc == 1) {
          stop("Number of clusters must be higher than 2")
        }
      }
    }
  }

```


RStudio

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Source

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```

R 4.3.0 · ~/
}
j <- table(c11)
s <- sum(j == 1)
j2 <- table(c12)
s2 <- sum(j2 == 1)
if (any(indice == 3) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 3] <- Indices.Traces(jeu, md,
clall1, index = "hart")
}
if (any(indice == 4) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 4] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$ccc
}
if (any(indice == 5) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 5] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$scott
}
if (any(indice == 6) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 6] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$marriot
}
if (any(indice == 7) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 7] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$trcovw
}
if (any(indice == 8) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 8] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$tracew
}
if (any(indice == 9) || (indice == 31) || (indice ==
32)) {
  res[nc - min_nc + 1, 9] <- Indices.WBT(x = jeu, cl = c11,
P = TT, s = ss, vv = vv)$friedman
}
if (any(indice == 10) || (indice == 31) || (indice ==
32)) {

```

RStudio

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Go to file/function

Addins

Source

Console Terminal Background Jobs

```

R 4.3.0 · ~/
res[nc - min_nc + 1, 9] <- Indices.WBT(x = jeu, c1 = c11,
    P = TT, s = ss, vv = vv)$friedman
}
if (any(indice == 10) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 10] <- Indices.WBT(x = jeu,
        c1 = c11, P = TT, s = ss, vv = vv)$rubin
}
if (any(indice == 14) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 14] <- Indices.WKWL(x = jeu,
        c11 = c11, c12 = c12)$duda
}
if (any(indice == 15) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 15] <- Indices.WKWL(x = jeu,
        c11 = c11, c12 = c12)$pseudot2
}
if (any(indice == 16) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 16] <- beale <- Indices.WKWL(x = jeu,
        c11 = c11, c12 = c12)$beale
}
if (any(indice == 14) || (indice == 15) || (indice ==
    16) || (indice == 31) || (indice == 32)) {
    NM <- Indices.WKWL(x = jeu, c11 = c11, c12 = c12)$NM
    NK <- Indices.WKWL(x = jeu, c11 = c11, c12 = c12)$NK
    NL <- Indices.WKWL(x = jeu, c11 = c11, c12 = c12)$NL
    zz <- 3.2
    zzz <- zz * sqrt(2 * (1 - 8/((pi^2) * pp)))/(NM *
        pp)
    if (any(indice == 14) || (indice == 31) || (indice ==
        32)) {
        resCritical[nc - min_nc + 1, 1] <- critValue <- 1 -
            (2/(pi * pp)) - zzz
    }
    if ((indice == 15) || (indice == 31) || (indice ==
        32)) {
        critValue <- 1 - (2/(pi * pp)) - zzz
        resCritical[nc - min_nc + 1, 2] <- ((1 - critValue)/critValue) *
            (NK + NL - 2)
    }
}

```

RStudio

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Source

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R 4.3.0 · ~/

```

    }
    if (any(indice == 16) || (indice == 31) || (indice ==
      32)) {
      df2 <- (NM - 2) * pp
      resCritical[nc - min_nc + 1, 3] <- 1 - pf(beale,
        pp, df2)
    }
  }
  if (any(indice == 18) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 18] <- Indices.Traces(jeu, md,
      clall1, index = "ball")
  }
  if (any(indice == 19) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 19] <- Indice.ptbiserial(x = jeu,
      md = md, cl1 = cl1)
  }
  if (any(indice == 20) || (indice == 32)) {
    if (method == 1) {
      resultSGAP <- Indice.Gap(x = jeu, clall = clall,
        reference.distribution = "unif", B = 10, method = "ward.D2",
        d = NULL, centrotypes = "centroids")
    }
    if (method == 2) {
      resultSGAP <- Indice.Gap(x = jeu, clall = clall,
        reference.distribution = "unif", B = 10, method = "single",
        d = NULL, centrotypes = "centroids")
    }
    if (method == 3) {
      resultSGAP <- Indice.Gap(x = jeu, clall = clall,
        reference.distribution = "unif", B = 10, method = "complete",
        d = NULL, centrotypes = "centroids")
    }
    if (method == 4) {
      resultSGAP <- Indice.Gap(x = jeu, clall = clall,
        reference.distribution = "unif", B = 10, method = "average",
        d = NULL, centrotypes = "centroids")
    }
    if (method == 5) {
      resultSGAP <- Indice.Gap(x = jeu, clall = clall,

```

RStudio

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R 4.3.0 · ~/

```

    resultSGAP <- Indice.Gap(x = jeu, clall = clall,
      reference.distribution = "unif", B = 10, method = "mcquitty",
      d = NULL, centrotypes = "centroids")
  }
  if (method == 6) {
    resultSGAP <- Indice.Gap(x = jeu, clall = clall,
      reference.distribution = "unif", B = 10, method = "median",
      d = NULL, centrotypes = "centroids")
  }
  if (method == 7) {
    resultSGAP <- Indice.Gap(x = jeu, clall = clall,
      reference.distribution = "unif", B = 10, method = "centroid",
      d = NULL, centrotypes = "centroids")
  }
  if (method == 9) {
    resultSGAP <- Indice.Gap(x = jeu, clall = clall,
      reference.distribution = "unif", B = 10, method = "ward.D",
      d = NULL, centrotypes = "centroids")
  }
  if (method == 8) {
    resultSGAP <- Indice.Gap(x = jeu, clall = clall,
      reference.distribution = "unif", B = 10, method = "k-means",
      d = NULL, centrotypes = "centroids")
  }
  res[nc - min_nc + 1, 20] <- resultSGAP$gap
  resCritical[nc - min_nc + 1, 4] <- resultSGAP$diffu
}
if (nc >= 2) {
  if (any(indice == 1) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 1] <- Indices.Traces(jeu,
      md, clall1, index = "kl")
  }
  if (any(indice == 2) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 2] <- Indices.Traces(jeu,
      md, clall1, index = "ch")
  }
  if (any(indice == 11) || (indice == 31) || (indice ==
    32)) {
    res[nc - min_nc + 1, 11] <- Indice.cindex(d = md,

```

RStudio

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Go to file/function Addins

Source

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R 4.3.0 · ~/

```

        cl = cl1)
    }
    if (any(indice == 12) || (indice == 31) || (indice ==
        32)) {
        res[nc - min_nc + 1, 12] <- Indice.DB(x = jeu,
            cl = cl1, d = NULL, centrotypes = "centroids",
            p = 2, q = 2)$DB
    }
    if (any(indice == 13) || (indice == 31) || (indice ==
        32)) {
        res[nc - min_nc + 1, 13] <- Indice.S(d = md,
            cl = cl1)
    }
    if (any(indice == 17) || (indice == 31) || (indice ==
        32)) {
        res[nc - min_nc + 1, 17] <- Indices.Traces(jeu,
            md, clall1, index = "ratkowsky")
    }
    if (any(indice == 21) || (indice == 31) || (indice ==
        32)) {
        res[nc - min_nc + 1, 21] <- Index.15and28(cl1 = cl1,
            cl2 = cl2, md = md)$frey
    }
    if (any(indice == 22) || (indice == 31) || (indice ==
        32)) {
        res[nc - min_nc + 1, 22] <- Index.15and28(cl1 = cl1,
            cl2 = cl2, md = md)$mcclain
    }
    if (any(indice == 23) || (indice == 32)) {
        res[nc - min_nc + 1, 23] <- Index.sPlussMoins(cl1 = cl1,
            md = md)$gamma
    }
    if (any(indice == 24) || (indice == 32)) {
        res[nc - min_nc + 1, 24] <- Index.sPlussMoins(cl1 = cl1,
            md = md)$gplus
    }
    if (any(indice == 25) || (indice == 32)) {
        res[nc - min_nc + 1, 25] <- Index.sPlussMoins(cl1 = cl1,
            md = md)$tau
    }
    if (any(indice == 26) || (indice == 31) || (indice ==
        32)) {

```

RStudio

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Go to file/function Addins

Source

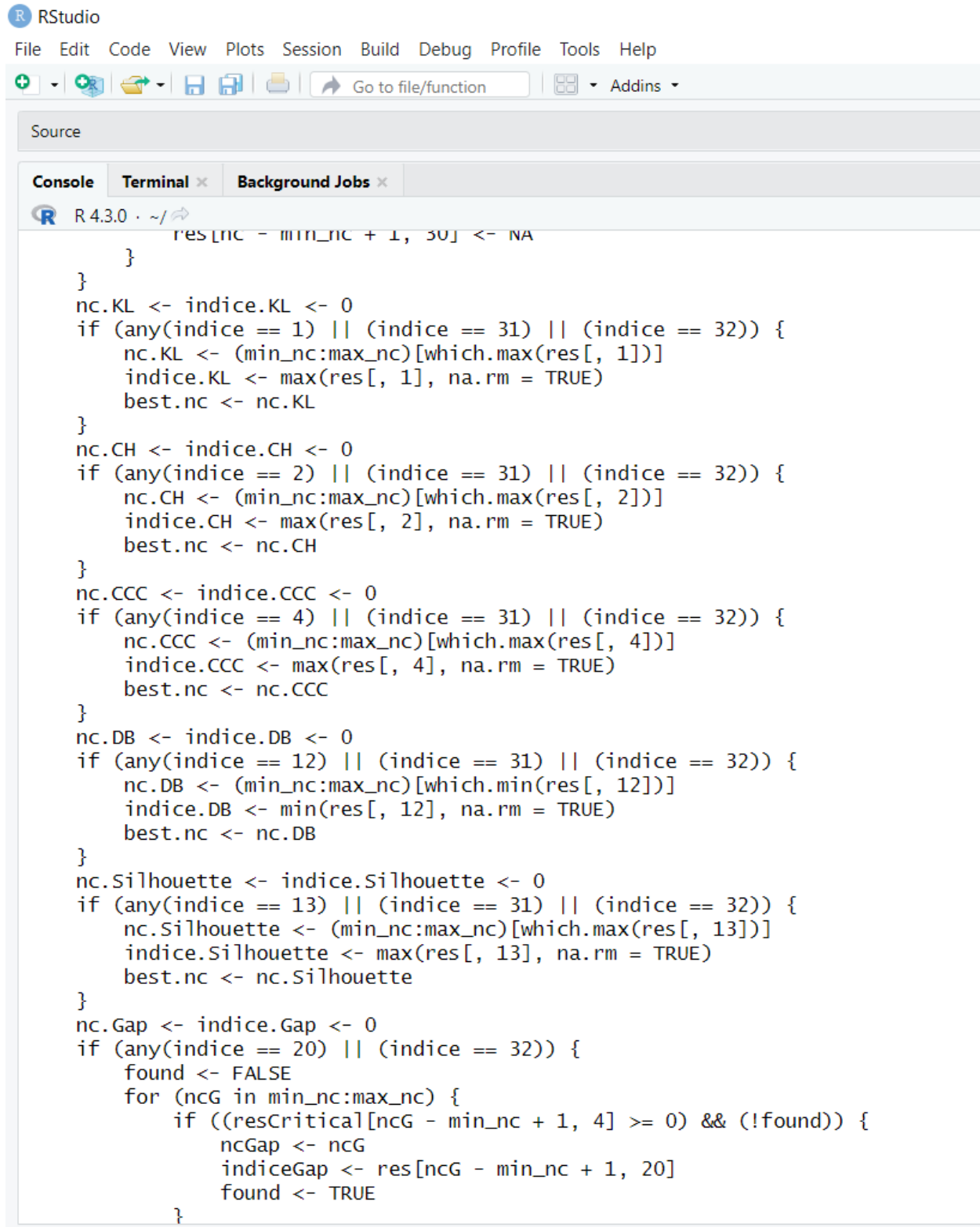
Console Terminal Background Jobs

R 4.3.0 · ~/

```

    if (any(indice == 26) || (indice == 31) || (indice ==
      32)) {
      res[nc - min_nc + 1, 26] <- Index.dunn(md, cl1,
        Data = jeu, method = NULL)
    }
    if (any(indice == 27) || (indice == 31) || (indice ==
      32)) {
      res[nc - min_nc + 1, 27] <- Index.Hubert(jeu,
        cl1)
    }
    if (any(indice == 28) || (indice == 31) || (indice ==
      32)) {
      res[nc - min_nc + 1, 28] <- Index.sdindex(jeu,
        clmax, cl1)
    }
    if (any(indice == 29) || (indice == 31) || (indice ==
      32)) {
      res[nc - min_nc + 1, 29] <- Index.Dindex(cl1,
        jeu)
    }
    if (any(indice == 30) || (indice == 31) || (indice ==
      32)) {
      res[nc - min_nc + 1, 30] <- Index.SDbw(jeu, cl1)
    }
  }
  else {
    res[nc - min_nc + 1, 1] <- NA
    res[nc - min_nc + 1, 2] <- NA
    res[nc - min_nc + 1, 11] <- NA
    res[nc - min_nc + 1, 12] <- NA
    res[nc - min_nc + 1, 13] <- NA
    res[nc - min_nc + 1, 17] <- NA
    res[nc - min_nc + 1, 21] <- NA
    res[nc - min_nc + 1, 22] <- NA
    res[nc - min_nc + 1, 23] <- NA
    res[nc - min_nc + 1, 24] <- NA
    res[nc - min_nc + 1, 25] <- NA
    res[nc - min_nc + 1, 26] <- NA
    res[nc - min_nc + 1, 27] <- NA
    res[nc - min_nc + 1, 28] <- NA
    res[nc - min_nc + 1, 29] <- NA
    res[nc - min_nc + 1, 30] <- NA
  }

```

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

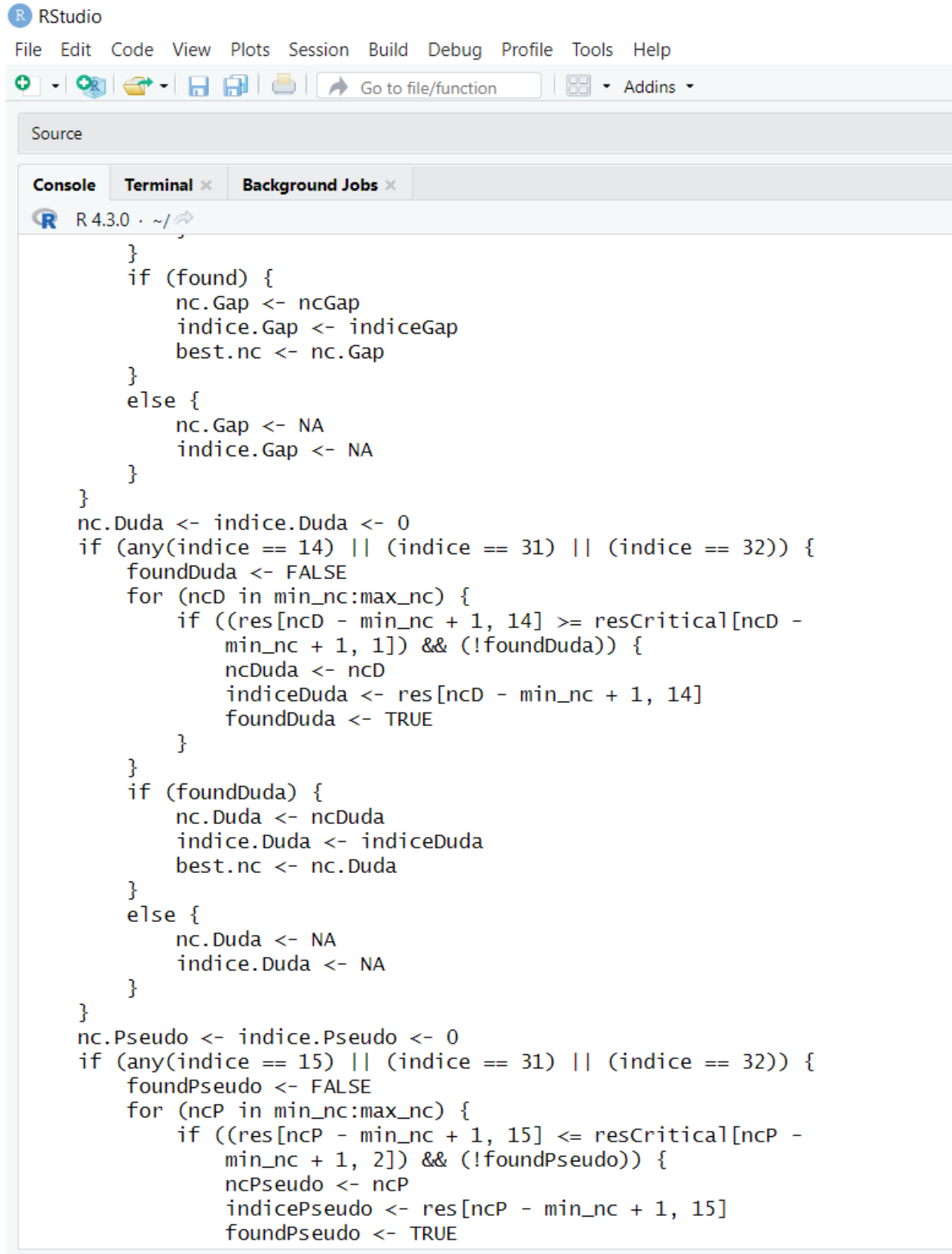
Console Terminal x Background Jobs x

R 4.3.0 · ~/

```

    res[nc - min_nc + 1, 30] <- NA
  }
}
nc.KL <- indice.KL <- 0
if (any(indice == 1) || (indice == 31) || (indice == 32)) {
  nc.KL <- (min_nc:max_nc)[which.max(res[, 1])]
  indice.KL <- max(res[, 1], na.rm = TRUE)
  best.nc <- nc.KL
}
nc.CH <- indice.CH <- 0
if (any(indice == 2) || (indice == 31) || (indice == 32)) {
  nc.CH <- (min_nc:max_nc)[which.max(res[, 2])]
  indice.CH <- max(res[, 2], na.rm = TRUE)
  best.nc <- nc.CH
}
nc.CCC <- indice.CCC <- 0
if (any(indice == 4) || (indice == 31) || (indice == 32)) {
  nc.CCC <- (min_nc:max_nc)[which.max(res[, 4])]
  indice.CCC <- max(res[, 4], na.rm = TRUE)
  best.nc <- nc.CCC
}
nc.DB <- indice.DB <- 0
if (any(indice == 12) || (indice == 31) || (indice == 32)) {
  nc.DB <- (min_nc:max_nc)[which.min(res[, 12])]
  indice.DB <- min(res[, 12], na.rm = TRUE)
  best.nc <- nc.DB
}
nc.Silhouette <- indice.Silhouette <- 0
if (any(indice == 13) || (indice == 31) || (indice == 32)) {
  nc.Silhouette <- (min_nc:max_nc)[which.max(res[, 13])]
  indice.Silhouette <- max(res[, 13], na.rm = TRUE)
  best.nc <- nc.Silhouette
}
nc.Gap <- indice.Gap <- 0
if (any(indice == 20) || (indice == 32)) {
  found <- FALSE
  for (ncG in min_nc:max_nc) {
    if ((resCritical[ncG - min_nc + 1, 4] >= 0) && (!found)) {
      ncGap <- ncG
      indiceGap <- res[ncG - min_nc + 1, 20]
      found <- TRUE
    }
  }
}

```



The image shows the RStudio interface. At the top, there's a menu bar with 'File', 'Edit', 'Code', 'View', 'Plots', 'Session', 'Build', 'Debug', 'Profile', 'Tools', and 'Help'. Below the menu bar is a toolbar with icons for file operations and a search bar labeled 'Go to file/function'. The main area is divided into three tabs: 'Source', 'Console', and 'Background Jobs'. The 'Console' tab is active, showing R code being executed. The code defines variables for 'nc.Gap', 'indice.Gap', 'best.nc', 'nc.Duda', 'indice.Duda', 'best.nc', 'nc.Pseudo', 'indice.Pseudo', and 'best.nc'. It uses conditional statements and loops to find the best values based on 'resCritical' and 'res' arrays.

```

}
if (found) {
  nc.Gap <- ncGap
  indice.Gap <- indiceGap
  best.nc <- nc.Gap
}
else {
  nc.Gap <- NA
  indice.Gap <- NA
}
}
nc.Duda <- indice.Duda <- 0
if (any(indice == 14) || (indice == 31) || (indice == 32)) {
  foundDuda <- FALSE
  for (ncD in min_nc:max_nc) {
    if ((res[ncD - min_nc + 1, 14] >= resCritical[ncD -
      min_nc + 1, 1]) && (!foundDuda)) {
      ncDuda <- ncD
      indiceDuda <- res[ncD - min_nc + 1, 14]
      foundDuda <- TRUE
    }
  }
  if (foundDuda) {
    nc.Duda <- ncDuda
    indice.Duda <- indiceDuda
    best.nc <- nc.Duda
  }
  else {
    nc.Duda <- NA
    indice.Duda <- NA
  }
}
nc.Pseudo <- indice.Pseudo <- 0
if (any(indice == 15) || (indice == 31) || (indice == 32)) {
  foundPseudo <- FALSE
  for (ncP in min_nc:max_nc) {
    if ((res[ncP - min_nc + 1, 15] <= resCritical[ncP -
      min_nc + 1, 2]) && (!foundPseudo)) {
      ncPseudo <- ncP
      indicePseudo <- res[ncP - min_nc + 1, 15]
      foundPseudo <- TRUE
    }
  }
}

```

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

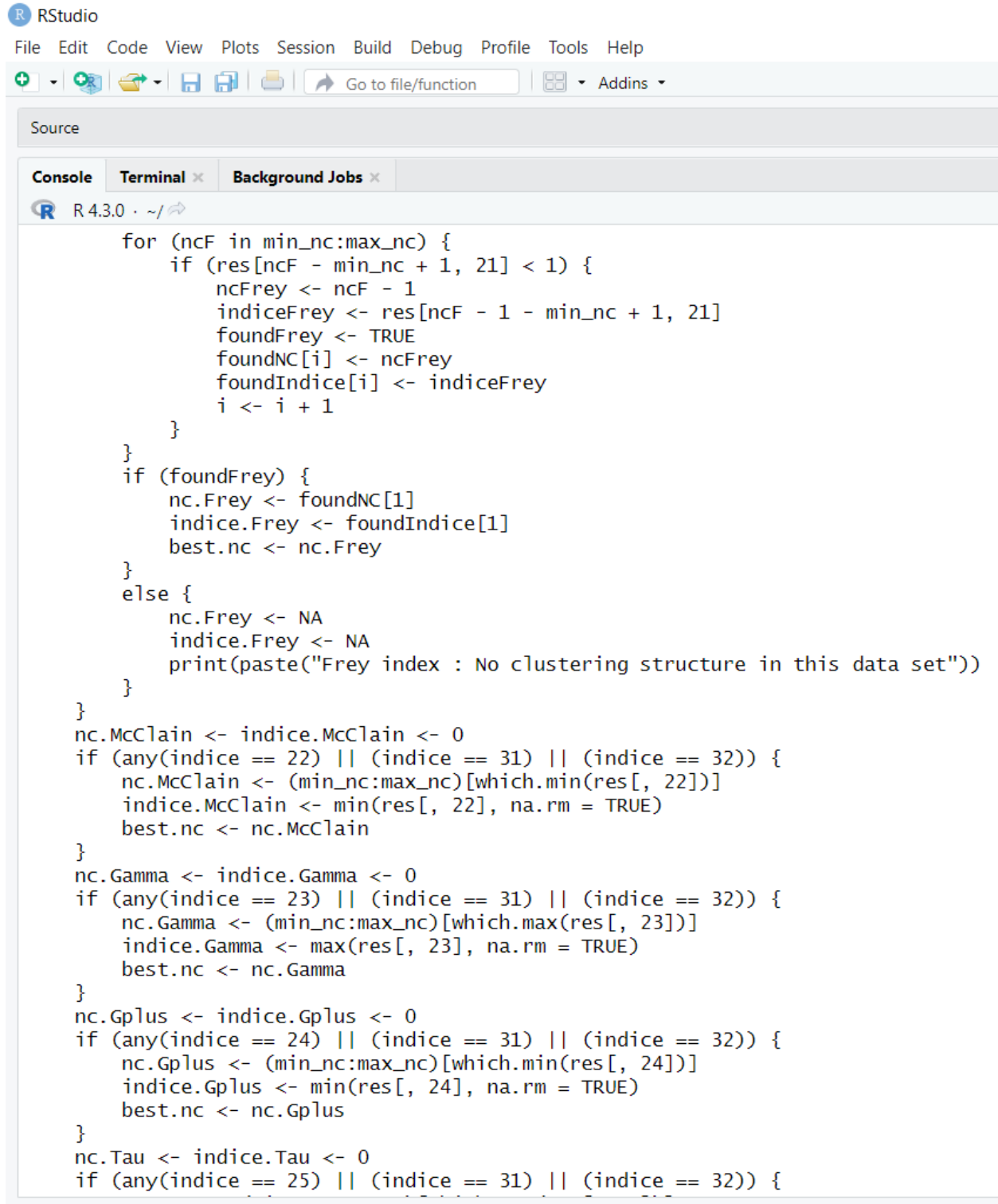
Console Terminal Background Jobs

R 4.3.0 · ~/

```

    }
  }
  if (foundPseudo) {
    nc.Pseudo <- ncPseudo
    indice.Pseudo <- indicePseudo
    best.nc <- nc.Pseudo
  }
  else {
    nc.Pseudo <- NA
    indice.Pseudo <- NA
  }
}
nc.Beale <- indice.Beale <- 0
if (any(indice == 16) || (indice == 31) || (indice == 32)) {
  foundBeale <- FALSE
  for (ncB in min_nc:max_nc) {
    if ((resCritical[ncB - min_nc + 1, 3] >= alphaBeale) &&
        (!foundBeale)) {
      ncBeale <- ncB
      indiceBeale <- res[ncB - min_nc + 1, 16]
      foundBeale <- TRUE
    }
  }
  if (foundBeale) {
    nc.Beale <- ncBeale
    indice.Beale <- indiceBeale
    best.nc <- nc.Beale
  }
  else {
    nc.Beale <- NA
    indice.Beale <- NA
  }
}
nc.ptbiseria1 <- indice.ptbiseria1 <- 0
if (any(indice == 19) || (indice == 31) || (indice == 32)) {
  nc.ptbiseria1 <- (min_nc:max_nc)[which.max(res[, 19])]
  indice.ptbiseria1 <- max(res[, 19], na.rm = TRUE)
  best.nc <- nc.ptbiseria1
}
foundNC <- foundIndice <- numeric(0)
nc.Frey <- indice.Frey <- 0
if (any(indice == 31) || (indice == 32) || (indice == 33)) {

```

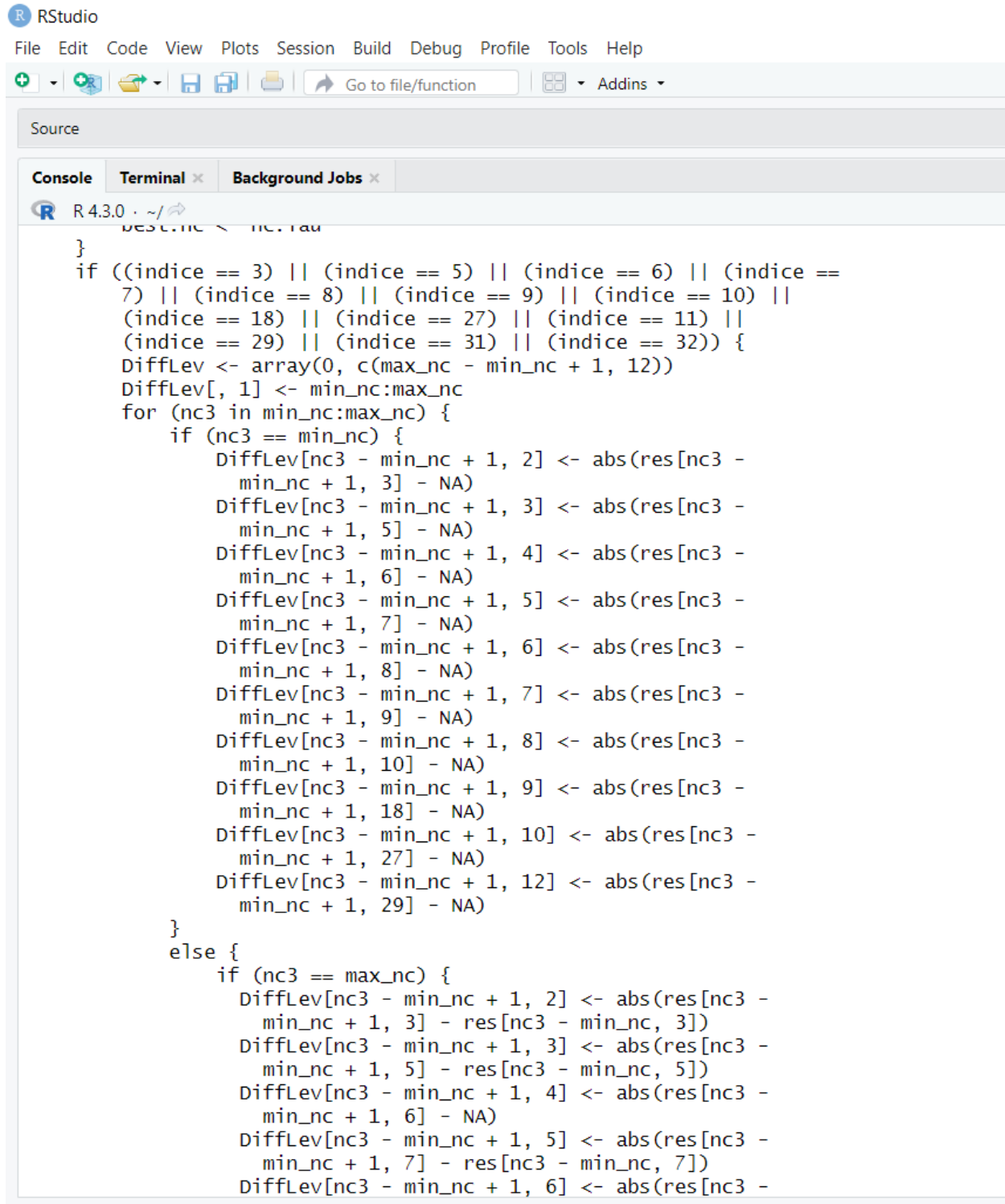


The image shows the RStudio interface. At the top, the title bar says 'RStudio'. Below it is a menu bar with 'File', 'Edit', 'Code', 'View', 'Plots', 'Session', 'Build', 'Debug', 'Profile', 'Tools', and 'Help'. Below the menu bar is a toolbar with icons for file operations and a search bar labeled 'Go to file/function'. Below the toolbar is a tab bar with 'Source', 'Console', 'Terminal', and 'Background Jobs'. The 'Console' tab is active, showing R code. The code is as follows:

```

for (ncF in min_nc:max_nc) {
  if (res[ncF - min_nc + 1, 21] < 1) {
    ncFrey <- ncF - 1
    indiceFrey <- res[ncF - 1 - min_nc + 1, 21]
    foundFrey <- TRUE
    foundNC[i] <- ncFrey
    foundIndice[i] <- indiceFrey
    i <- i + 1
  }
}
if (foundFrey) {
  nc.Frey <- foundNC[1]
  indice.Frey <- foundIndice[1]
  best.nc <- nc.Frey
}
else {
  nc.Frey <- NA
  indice.Frey <- NA
  print(paste("Frey index : No clustering structure in this data set"))
}
}
nc.McClain <- indice.McClain <- 0
if (any(indice == 22) || (indice == 31) || (indice == 32)) {
  nc.McClain <- (min_nc:max_nc)[which.min(res[, 22])]
  indice.McClain <- min(res[, 22], na.rm = TRUE)
  best.nc <- nc.McClain
}
nc.Gamma <- indice.Gamma <- 0
if (any(indice == 23) || (indice == 31) || (indice == 32)) {
  nc.Gamma <- (min_nc:max_nc)[which.max(res[, 23])]
  indice.Gamma <- max(res[, 23], na.rm = TRUE)
  best.nc <- nc.Gamma
}
nc.Gplus <- indice.Gplus <- 0
if (any(indice == 24) || (indice == 31) || (indice == 32)) {
  nc.Gplus <- (min_nc:max_nc)[which.min(res[, 24])]
  indice.Gplus <- min(res[, 24], na.rm = TRUE)
  best.nc <- nc.Gplus
}
nc.Tau <- indice.Tau <- 0
if (any(indice == 25) || (indice == 31) || (indice == 32)) {

```



RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

Console Terminal Background Jobs

```

R 4.3.0 . ~/
best.nc <- nc.fau
}
if ((indice == 3) || (indice == 5) || (indice == 6) || (indice ==
7) || (indice == 8) || (indice == 9) || (indice == 10) ||
(indice == 18) || (indice == 27) || (indice == 11) ||
(indice == 29) || (indice == 31) || (indice == 32)) {
  DiffLev <- array(0, c(max_nc - min_nc + 1, 12))
  DiffLev[, 1] <- min_nc:max_nc
  for (nc3 in min_nc:max_nc) {
    if (nc3 == min_nc) {
      DiffLev[nc3 - min_nc + 1, 2] <- abs(res[nc3 -
min_nc + 1, 3] - NA)
      DiffLev[nc3 - min_nc + 1, 3] <- abs(res[nc3 -
min_nc + 1, 5] - NA)
      DiffLev[nc3 - min_nc + 1, 4] <- abs(res[nc3 -
min_nc + 1, 6] - NA)
      DiffLev[nc3 - min_nc + 1, 5] <- abs(res[nc3 -
min_nc + 1, 7] - NA)
      DiffLev[nc3 - min_nc + 1, 6] <- abs(res[nc3 -
min_nc + 1, 8] - NA)
      DiffLev[nc3 - min_nc + 1, 7] <- abs(res[nc3 -
min_nc + 1, 9] - NA)
      DiffLev[nc3 - min_nc + 1, 8] <- abs(res[nc3 -
min_nc + 1, 10] - NA)
      DiffLev[nc3 - min_nc + 1, 9] <- abs(res[nc3 -
min_nc + 1, 18] - NA)
      DiffLev[nc3 - min_nc + 1, 10] <- abs(res[nc3 -
min_nc + 1, 27] - NA)
      DiffLev[nc3 - min_nc + 1, 12] <- abs(res[nc3 -
min_nc + 1, 29] - NA)
    }
    else {
      if (nc3 == max_nc) {
        DiffLev[nc3 - min_nc + 1, 2] <- abs(res[nc3 -
min_nc + 1, 3] - res[nc3 - min_nc, 3])
        DiffLev[nc3 - min_nc + 1, 3] <- abs(res[nc3 -
min_nc + 1, 5] - res[nc3 - min_nc, 5])
        DiffLev[nc3 - min_nc + 1, 4] <- abs(res[nc3 -
min_nc + 1, 6] - NA)
        DiffLev[nc3 - min_nc + 1, 5] <- abs(res[nc3 -
min_nc + 1, 7] - res[nc3 - min_nc, 7])
        DiffLev[nc3 - min_nc + 1, 6] <- abs(res[nc3 -

```

RStudio

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Go to file/function Addins

Source

Console Terminal Background Jobs

R 4.3.0 · ~/

```

min_nc + 1, 9] - res[nc3 - min_nc, 9])
DiffLev[nc3 - min_nc + 1, 8] <- abs(res[nc3 -
min_nc + 1, 10] - NA)
DiffLev[nc3 - min_nc + 1, 9] <- abs(res[nc3 -
min_nc + 1, 18] - res[nc3 - min_nc, 18])
DiffLev[nc3 - min_nc + 1, 10] <- abs(res[nc3 -
min_nc + 1, 27] - NA)
DiffLev[nc3 - min_nc + 1, 12] <- abs(res[nc3 -
min_nc + 1, 29] - NA)
}
else {
DiffLev[nc3 - min_nc + 1, 2] <- abs(res[nc3 -
min_nc + 1, 3] - res[nc3 - min_nc, 3])
DiffLev[nc3 - min_nc + 1, 3] <- abs(res[nc3 -
min_nc + 1, 5] - res[nc3 - min_nc, 5])
DiffLev[nc3 - min_nc + 1, 4] <- ((res[nc3 -
min_nc + 2, 6] - res[nc3 - min_nc + 1, 6]) -
(res[nc3 - min_nc + 1, 6] - res[nc3 - min_nc,
6]))
DiffLev[nc3 - min_nc + 1, 5] <- abs(res[nc3 -
min_nc + 1, 7] - res[nc3 - min_nc, 7])
DiffLev[nc3 - min_nc + 1, 6] <- ((res[nc3 -
min_nc + 2, 8] - res[nc3 - min_nc + 1, 8]) -
(res[nc3 - min_nc + 1, 8] - res[nc3 - min_nc,
8]))
DiffLev[nc3 - min_nc + 1, 7] <- abs(res[nc3 -
min_nc + 1, 9] - res[nc3 - min_nc, 9])
DiffLev[nc3 - min_nc + 1, 8] <- ((res[nc3 -
min_nc + 2, 10] - res[nc3 - min_nc + 1, 10]) -
(res[nc3 - min_nc + 1, 10] - res[nc3 - min_nc,
10]))
DiffLev[nc3 - min_nc + 1, 9] <- abs(res[nc3 -
min_nc + 1, 18] - res[nc3 - min_nc, 18])
DiffLev[nc3 - min_nc + 1, 10] <- abs((res[nc3 -
min_nc + 1, 27] - res[nc3 - min_nc, 27]))
DiffLev[nc3 - min_nc + 1, 12] <- ((res[nc3 -
min_nc + 2, 29] - res[nc3 - min_nc + 1, 29]) -
(res[nc3 - min_nc + 1, 29] - res[nc3 - min_nc,
29]))
}
}
}

```


RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

Console Terminal Background Jobs

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```

if (any(indice == 3) || (indice == 31) || (indice == 32)) {
  nc.Hartigan <- DiffLev[, 1][which.max(DiffLev[, 2])]
  indice.Hartigan <- max(DiffLev[, 2], na.rm = TRUE)
  best.nc <- nc.Hartigan
}
nc.Ratkowsky <- indice.Ratkowsky <- 0
if (any(indice == 17) || (indice == 31) || (indice == 32)) {
  nc.Ratkowsky <- (min_nc:max_nc)[which.max(res[, 17])]
  indice.Ratkowsky <- max(res[, 17], na.rm = TRUE)
  best.nc <- nc.Ratkowsky
}
nc.cindex <- indice.cindex <- 0
if (any(indice == 11) || (indice == 31) || (indice == 32)) {
  nc.cindex <- (min_nc:max_nc)[which.min(res[, 11])]
  indice.cindex <- min(res[, 11], na.rm = TRUE)
  best.nc <- nc.cindex
}
nc.Scott <- indice.Scott <- 0
if (any(indice == 5) || (indice == 31) || (indice == 32)) {
  nc.Scott <- DiffLev[, 1][which.max(DiffLev[, 3])]
  indice.Scott <- max(DiffLev[, 3], na.rm = TRUE)
  best.nc <- nc.Scott
}
nc.Marriot <- indice.Marriot <- 0
if (any(indice == 6) || (indice == 31) || (indice == 32)) {
  nc.Marriot <- DiffLev[, 1][which.max(DiffLev[, 4])]
  round(nc.Marriot, digits = 1)
  indice.Marriot <- max(DiffLev[, 4], na.rm = TRUE)
  best.nc <- nc.Marriot
}
nc.TrCovW <- indice.TrCovW <- 0
if (any(indice == 7) || (indice == 31) || (indice == 32)) {
  nc.TrCovW <- DiffLev[, 1][which.max(DiffLev[, 5])]
  indice.TrCovW <- max(DiffLev[, 5], na.rm = TRUE)
  best.nc <- nc.TrCovW
}
nc.TraceW <- indice.TraceW <- 0
if (any(indice == 8) || (indice == 31) || (indice == 32)) {
  nc.TraceW <- DiffLev[, 1][which.max(DiffLev[, 6])]
  indice.TraceW <- max(DiffLev[, 6], na.rm = TRUE)
  best.nc <- nc.TraceW
}

```

RStudio

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Go to file/function Addins

Source

Console Terminal Background Jobs

```

R 4.3.0 · ~/
nc.Friedman <- DiffLev[, 1][which.max(DiffLev[, 7])]
indice.Friedman <- max(DiffLev[, 7], na.rm = TRUE)
best.nc <- nc.Friedman
}
nc.Rubin <- indice.Rubin <- 0
if (any(indice == 10) || (indice == 31) || (indice == 32)) {
  nc.Rubin <- DiffLev[, 1][which.min(DiffLev[, 8])]
  indice.Rubin <- min(DiffLev[, 8], na.rm = TRUE)
  best.nc <- nc.Rubin
}
nc.Ball <- indice.Ball <- 0
if (any(indice == 18) || (indice == 31) || (indice == 32)) {
  nc.Ball <- DiffLev[, 1][which.max(DiffLev[, 9])]
  indice.Ball <- max(DiffLev[, 9], na.rm = TRUE)
  best.nc <- nc.Ball
}
nc.Dunn <- indice.Dunn <- 0
if (any(indice == 26) || (indice == 31) || (indice == 32)) {
  nc.Dunn <- (min_nc:max_nc)[which.max(res[, 26])]
  indice.Dunn <- max(res[, 26], na.rm = TRUE)
  best.nc <- nc.Dunn
}
nc.Hubert <- indice.Hubert <- 0
if (any(indice == 27) || (indice == 31) || (indice == 32)) {
  nc.Hubert <- 0
  indice.Hubert <- 0
  par(mfrow = c(1, 2))
  plot(x_axis, res[, 27], tck = 0, type = "b", col = "red",
        xlab = expression(paste("Number of clusters")),
        ylab = expression(paste("Hubert Statistic values")))
  plot(DiffLev[, 1], DiffLev[, 10], tck = 0, type = "b",
        col = "blue", xlab = expression(paste("Number of clusters")),
        ylab = expression(paste("Hubert statistic second differences")))
  cat(paste("**** : The Hubert index is a graphical method of determining the number of clusters\n
            In the plot of Hubert index, we seek a significant knee that corresponds to a\n
            significant increase of the value of the measure i.e the significant peak in Hubert\n
            index second differences plot.",
            "\n", "\n"))
}
nc.sdindex <- indice.sdindex <- 0
if (any(indice == 28) || (indice == 31) || (indice == 32)) {

```

RStudio

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Go to file/function Addins

Source

Console Terminal Background Jobs

```

R 4.3.0 · ~/
  "\n", "\n"))
}
nc.sdindex <- indice.sdindex <- 0
if (any(indice == 28) || (indice == 31) || (indice == 32)) {
  nc.sdindex <- (min_nc:max_nc)[which.min(res[, 28])]
  indice.sdindex <- min(res[, 28], na.rm = TRUE)
  best.nc <- nc.sdindex
}
nc.Dindex <- indice.Dindex <- 0
if (any(indice == 29) || (indice == 31) || (indice == 32)) {
  nc.Dindex <- 0
  indice.Dindex <- 0
  par(mfrow = c(1, 2))
  plot(x_axis, res[, 29], tck = 0, type = "b", col = "red",
        xlab = expression(paste("Number of clusters ")),
        ylab = expression(paste("Dindex Values")))
  plot(DiffLev[, 1], DiffLev[, 12], tck = 0, type = "b",
        col = "blue", xlab = expression(paste("Number of clusters ")),
        ylab = expression(paste("Second differences Dindex Values")))
  cat(paste("*** : The D index is a graphical method of determining the number of clusters.
              In the plot of D index, we seek a significant knee (the significant peak in Dinde
              second differences plot) that corresponds to a significant increase of the value
              the measure.",
              "\n", "\n"))
}
nc.SDbw <- indice.SDbw <- 0
if (any(indice == 30) || (indice == 31) || (indice == 32)) {
  nc.SDbw <- (min_nc:max_nc)[which.min(res[, 30])]
  indice.SDbw <- min(res[, 30], na.rm = TRUE)
  best.nc <- nc.SDbw
}
if (indice < 31) {
  res <- res[, c(indice)]
  if (indice == 14) {
    resCritical <- resCritical[, 1]
  }
  if (indice == 15) {
    resCritical <- resCritical[, 2]
  }
  if (indice == 16) {
    resCritical <- resCritical[, 3]
  }
}

```

RStudio

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Go to file/function

Addins

Source

Console Terminal x Background Jobs x

R 4.3.0 · ~/

```

    }
  }
  if (indice == 31) {
    res <- res[, c(1:19, 21:22, 26:30)]
    resCritical <- resCritical[, c(1:3)]
  }
  if (any(indice == 20) || (indice == 23) || (indice == 24) ||
      (indice == 25) || (indice == 32)) {
    results <- c(nc.KL, indice.KL, nc.CH, indice.CH, nc.Hartigan,
                 indice.Hartigan, nc.CCC, indice.CCC, nc.Scott, indice.Scott,
                 nc.Marriot, indice.Marriot, nc.TrCovW, indice.TrCovW,
                 nc.TraceW, indice.TraceW, nc.Friedman, indice.Friedman,
                 nc.Rubin, indice.Rubin, nc.cindex, indice.cindex,
                 nc.DB, indice.DB, nc.Silhouette, indice.Silhouette,
                 nc.Duda, indice.Duda, nc.Pseudo, indice.Pseudo, nc.Beale,
                 indice.Beale, nc.Ratkowsky, indice.Ratkowsky, nc.Ball,
                 indice.Ball, nc.ptbiserial, indice.ptbiserial, nc.Gap,
                 indice.Gap, nc.Frey, indice.Frey, nc.McClain, indice.McClain,
                 nc.Gamma, indice.Gamma, nc.Gplus, indice.Gplus, nc.Tau,
                 indice.Tau, nc.Dunn, indice.Dunn, nc.Hubert, indice.Hubert,
                 nc.sdindex, indice.sdindex, nc.Dindex, indice.Dindex,
                 nc.SDbw, indice.SDbw)
    results1 <- matrix(c(results), nrow = 2, ncol = 30)
    resultats <- matrix(c(results), nrow = 2, ncol = 30,
                        dimnames = list(c("Number_clusters", "Value_Index"),
                                         c("KL", "CH", "Hartigan", "CCC", "Scott", "Marriot",
                                           "TrCovW", "TraceW", "Friedman", "Rubin", "Cindex",
                                           "DB", "Silhouette", "Duda", "PseudoT2", "Beale",
                                           "Ratkowsky", "Ball", "PtBiserial", "Gap", "Frey",
                                           "McClain", "Gamma", "Gplus", "Tau", "Dunn",
                                           "Hubert", "SDindex", "Dindex", "SDbw"))))
  }
  else {
    results <- c(nc.KL, indice.KL, nc.CH, indice.CH, nc.Hartigan,
                 indice.Hartigan, nc.CCC, indice.CCC, nc.Scott, indice.Scott,
                 nc.Marriot, indice.Marriot, nc.TrCovW, indice.TrCovW,
                 nc.TraceW, indice.TraceW, nc.Friedman, indice.Friedman,
                 nc.Rubin, indice.Rubin, nc.cindex, indice.cindex,
                 nc.DB, indice.DB, nc.Silhouette, indice.Silhouette,
                 nc.Duda, indice.Duda, nc.Pseudo, indice.Pseudo, nc.Beale,
                 indice.Beale, nc.Ratkowsky, indice.Ratkowsky, nc.Ball,
                 indice.Ball, nc.ptbiserial, indice.ptbiserial, nc.Frey,
                 indice.Frey, nc.McClain, indice.McClain, nc.Gamma, indice.Gamma,
                 nc.Gplus, indice.Gplus, nc.Tau, indice.Tau, nc.Dunn, indice.Dunn,
                 nc.Hubert, indice.Hubert, nc.sdindex, indice.sdindex, nc.Dindex,
                 indice.Dindex, nc.SDbw, indice.SDbw)
    results1 <- matrix(c(results), nrow = 2, ncol = 30)
    resultats <- matrix(c(results), nrow = 2, ncol = 30,
                        dimnames = list(c("Number_clusters", "Value_Index"),
                                         c("KL", "CH", "Hartigan", "CCC", "Scott", "Marriot",
                                           "TrCovW", "TraceW", "Friedman", "Rubin", "Cindex",
                                           "DB", "Silhouette", "Duda", "PseudoT2", "Beale",
                                           "Ratkowsky", "Ball", "PtBiserial", "Gap", "Frey",
                                           "McClain", "Gamma", "Gplus", "Tau", "Dunn",
                                           "Hubert", "SDindex", "Dindex", "SDbw"))))
  }
}

```

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```

    indice.sdindex, nc.Dindex, indice.Dindex, nc.SDbw,
    indice.SDbw)
results1 <- matrix(c(results), nrow = 2, ncol = 26)
resultats <- matrix(c(results), nrow = 2, ncol = 26,
  dimnames = list(c("Number_clusters", "Value_Index"),
    c("KL", "CH", "Hartigan", "CCC", "Scott", "Marriot",
      "TrCovW", "TraceW", "Friedman", "Rubin", "Cindex",
      "DB", "Silhouette", "Duda", "PseudoT2", "Beale",
      "Ratkowsky", "Ball", "PtBiserial", "Frey",
      "McClain", "Dunn", "Hubert", "SDindex", "Dindex",
      "SDbw")))
}
if (any(indice <= 20) || (indice == 23) || (indice == 24) ||
  (indice == 25)) {
  resultats <- resultats[, c(indice)]
}
if (any(indice == 21) || (indice == 22)) {
  indice3 <- indice - 1
  resultats <- resultats[, c(indice3)]
}
if (any(indice == 26) || (indice == 27) || (indice == 28) ||
  (indice == 29) || (indice == 30)) {
  indice4 <- indice - 4
  resultats <- resultats[, c(indice4)]
}
resultats <- round(resultats, digits = 4)
res <- round(res, digits = 4)
resCritical <- round(resCritical, digits = 4)
if (any(indice == 31) || (indice == 32)) {
  cat("*****",
    "\n")
  cat("* Among all indices: ",
    "\n")
  BestCluster <- results1[1, ]
  c = 0
  for (i in min.nc:max.nc) {
    vect <- which(BestCluster == i)
    if (length(vect) > 0)
      cat("*", length(vect), "proposed", i, "as the best number of clusters",
        "\n")
    if (c < length(vect)) {

```

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```

    }
    cat("\n", "                ***** Conclusion *****",
        "\n", "\n")
    cat("* According to the majority rule, the best number of clusters is ",
        j, "\n", "\n", "\n")
    cat("*****",
        "\n")
    if (any(method == 1) || (method == 2) || (method == 3) ||
        (method == 4) || (method == 5) || (method == 6) ||
        (method == 7) || (method == 9))
      partition <- cutree(hc, k = j)
    else {
      set.seed(1)
      partition <- kmeans(jeu, j)$cluster
    }
  }
  if (any(indice == 1) || (indice == 2) || (indice == 3) ||
      (indice == 4) || (indice == 5) || (indice == 6) || (indice ==
      7) || (indice == 8) || (indice == 9) || (indice == 10) ||
      (indice == 11) || (indice == 12) || (indice == 13) ||
      (indice == 14) || (indice == 15) || (indice == 16) ||
      (indice == 17) || (indice == 18) || (indice == 19) ||
      (indice == 20) || (indice == 21) || (indice == 22) ||
      (indice == 23) || (indice == 24) || (indice == 25) ||
      (indice == 26) || (indice == 28) || (indice == 30)) {
    if (any(method == 1) || (method == 2) || (method == 3) ||
        (method == 4) || (method == 5) || (method == 6) ||
        (method == 7) || (method == 9))
      partition <- cutree(hc, k = best.nc)
    else {
      set.seed(1)
      partition <- kmeans(jeu, best.nc)$cluster
    }
  }
  if ((indice == 14) || (indice == 15) || (indice == 16) ||
      (indice == 20) || (indice == 31) || (indice == 32)) {
    results.final <- list(All.index = res, All.CriticalValues = resCritical,
      Best.nc = resultats, Best.partition = partition)
  }
  if ((indice == 27) || (indice == 29))
    results.final <- list(All.index = res)

```


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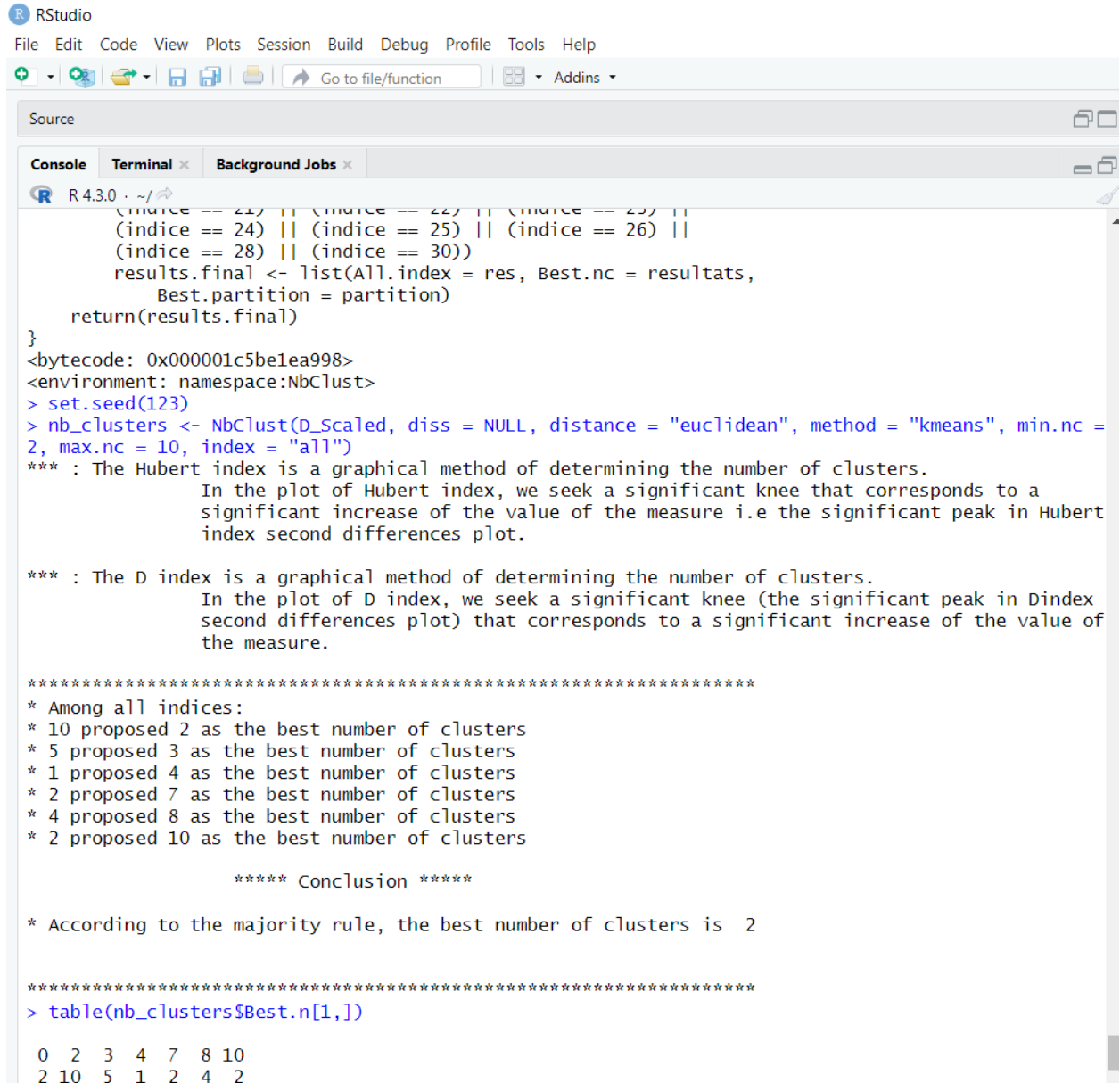
Source

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```

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}
if (any(indice == 1) || (indice == 2) || (indice == 3) ||
    (indice == 4) || (indice == 5) || (indice == 6) || (indice ==
7) || (indice == 8) || (indice == 9) || (indice == 10) ||
    (indice == 11) || (indice == 12) || (indice == 13) ||
    (indice == 14) || (indice == 15) || (indice == 16) ||
    (indice == 17) || (indice == 18) || (indice == 19) ||
    (indice == 20) || (indice == 21) || (indice == 22) ||
    (indice == 23) || (indice == 24) || (indice == 25) ||
    (indice == 26) || (indice == 28) || (indice == 30)) {
  if (any(method == 1) || (method == 2) || (method == 3) ||
      (method == 4) || (method == 5) || (method == 6) ||
      (method == 7) || (method == 9))
    partition <- cutree(hc, k = best.nc)
  else {
    set.seed(1)
    partition <- kmeans(jeu, best.nc)$cluster
  }
}
if ((indice == 14) || (indice == 15) || (indice == 16) ||
    (indice == 20) || (indice == 31) || (indice == 32)) {
  results.final <- list(All.index = res, All.CriticalValues = resCritical,
    Best.nc = resultats, Best.partition = partition)
}
if ((indice == 27) || (indice == 29))
  results.final <- list(All.index = res)
if (any(indice == 1) || (indice == 2) || (indice == 3) ||
    (indice == 4) || (indice == 5) || (indice == 6) || (indice ==
7) || (indice == 8) || (indice == 9) || (indice == 10) ||
    (indice == 11) || (indice == 12) || (indice == 13) ||
    (indice == 17) || (indice == 18) || (indice == 19) ||
    (indice == 21) || (indice == 22) || (indice == 23) ||
    (indice == 24) || (indice == 25) || (indice == 26) ||
    (indice == 28) || (indice == 30))
  results.final <- list(All.index = res, Best.nc = resultats,
    Best.partition = partition)
return(results.final)
}
<bytecode: 0x000001c5be1ea998>
<environment: namespace:NbClust>
> |

```



The screenshot shows the RStudio environment. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. Below the menu is a toolbar with icons for saving, opening, and navigating files. The main window is divided into three panes: Source, Console, and Background Jobs. The Source pane shows a script with R code for clustering analysis. The Console pane shows the output of the script, including comments about the Hubert and D indices, a list of proposed cluster numbers for various indices, a conclusion, and a table of results.

```

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  (indice == 21) || (indice == 22) || (indice == 23) ||
  (indice == 24) || (indice == 25) || (indice == 26) ||
  (indice == 28) || (indice == 30))
  results.final <- list(All.index = res, Best.nc = resultats,
    Best.partition = partition)
  return(results.final)
}
<bytecode: 0x000001c5be1ea998>
<environment: namespace:NbClust>
> set.seed(123)
> nb_clusters <- NbClust(D_Scaled, diss = NULL, distance = "euclidean", method = "kmeans", min.nc =
2, max.nc = 10, index = "all")
*** : The Hubert index is a graphical method of determining the number of clusters.
      In the plot of Hubert index, we seek a significant knee that corresponds to a
      significant increase of the value of the measure i.e the significant peak in Hubert
      index second differences plot.

*** : The D index is a graphical method of determining the number of clusters.
      In the plot of D index, we seek a significant knee (the significant peak in Dindex
      second differences plot) that corresponds to a significant increase of the value of
      the measure.

*****
* Among all indices:
* 10 proposed 2 as the best number of clusters
* 5 proposed 3 as the best number of clusters
* 1 proposed 4 as the best number of clusters
* 2 proposed 7 as the best number of clusters
* 4 proposed 8 as the best number of clusters
* 2 proposed 10 as the best number of clusters

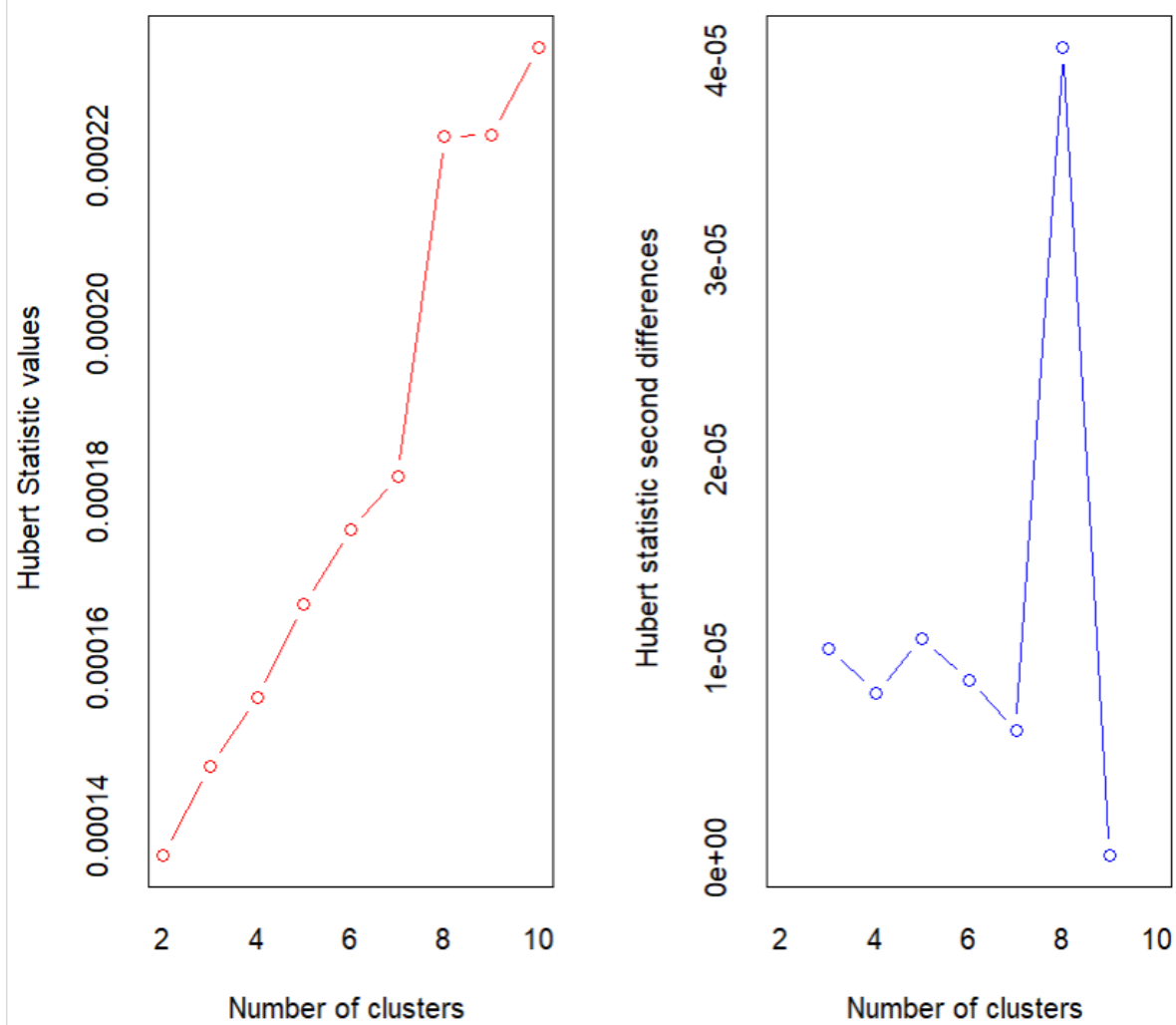
      ***** Conclusion *****

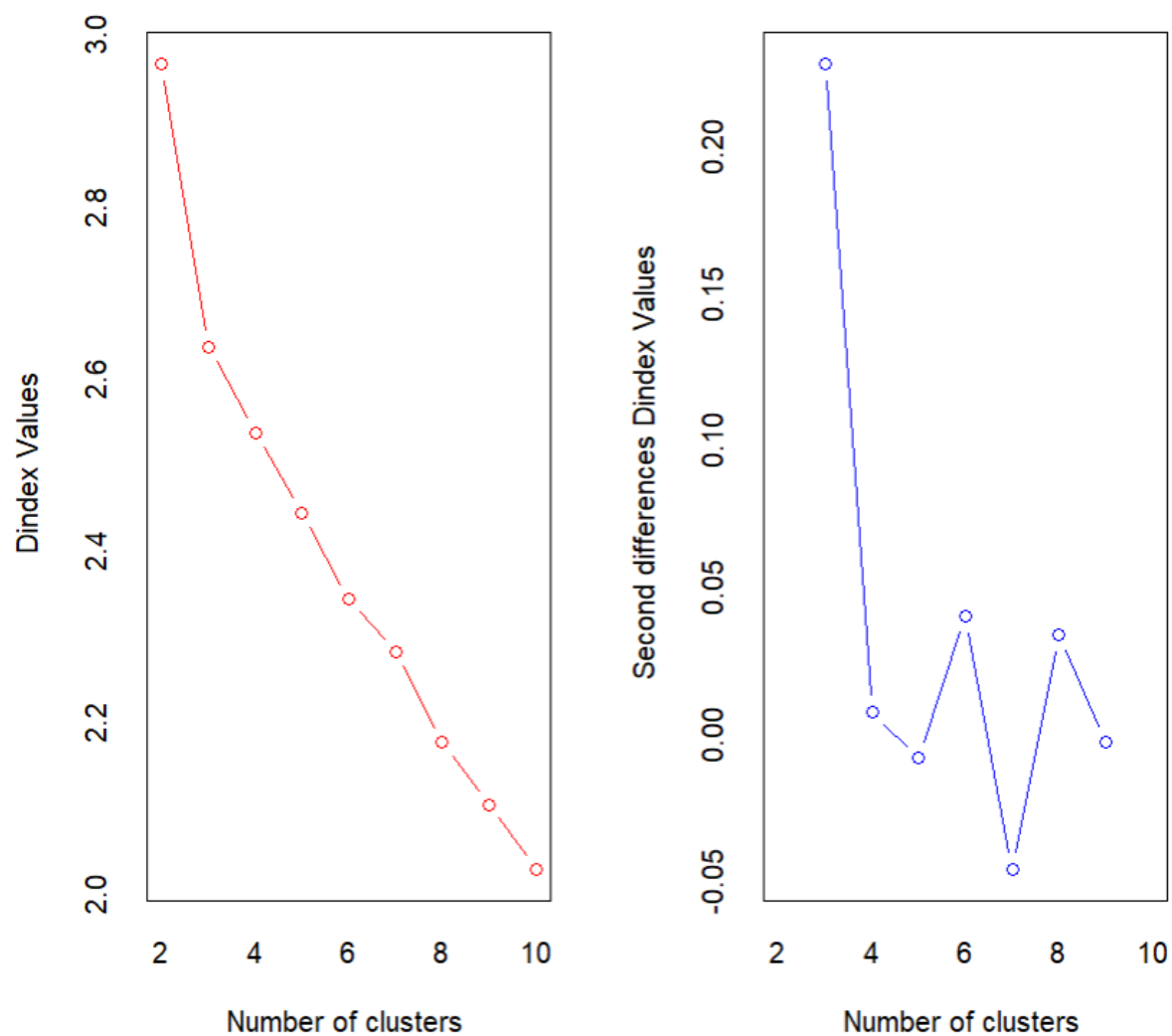
* According to the majority rule, the best number of clusters is 2

*****
> table(nb_clusters$Best.n[,1])

0 2 3 4 7 8 10
2 10 5 1 2 4 2

```





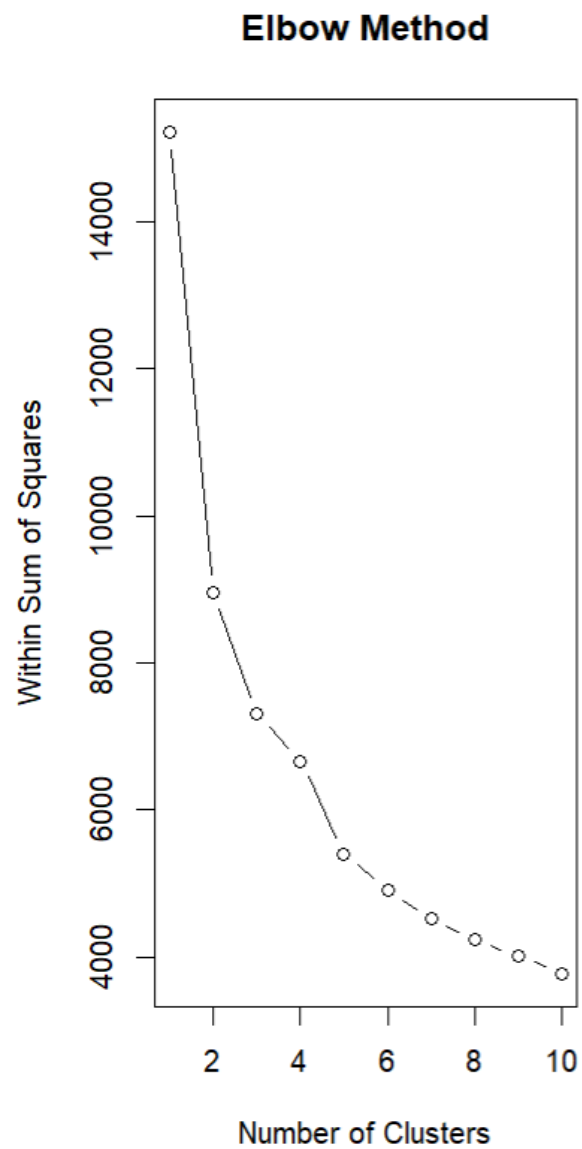
Elbow method

The elbow method is a method used to calculate the ideal number of clusters for a given dataset in clustering algorithms like K-means. Finding the elbow point in the graph, when more clusters do not significantly increase the explained variance, involves graphing the explained variation versus the number of clusters. When the number of clusters is unknown beforehand, this technique aids in balancing the quality of fit and complexity of the model and finding the right number.

```

> table(nb_clusters$Best.n[1,])
 0  2  3  4  7  8 10
2 10  5  1  2  4  2
> # Elbow method
> wss_ratio <- c()
> for (i in 1:10) {
+   set.seed(123)
+   kmeans_clusters <- kmeans(D_Scaled, centers = i, nstart = 10)
+   wss_ratio[i] <- kmeans_clusters$tot.withinss
+ }
> plot(1:10, wss_ratio, type = "b", main = "Elbow Method", xlab = "Number of Clusters", ylab = "Within Sum of Squares")
> |

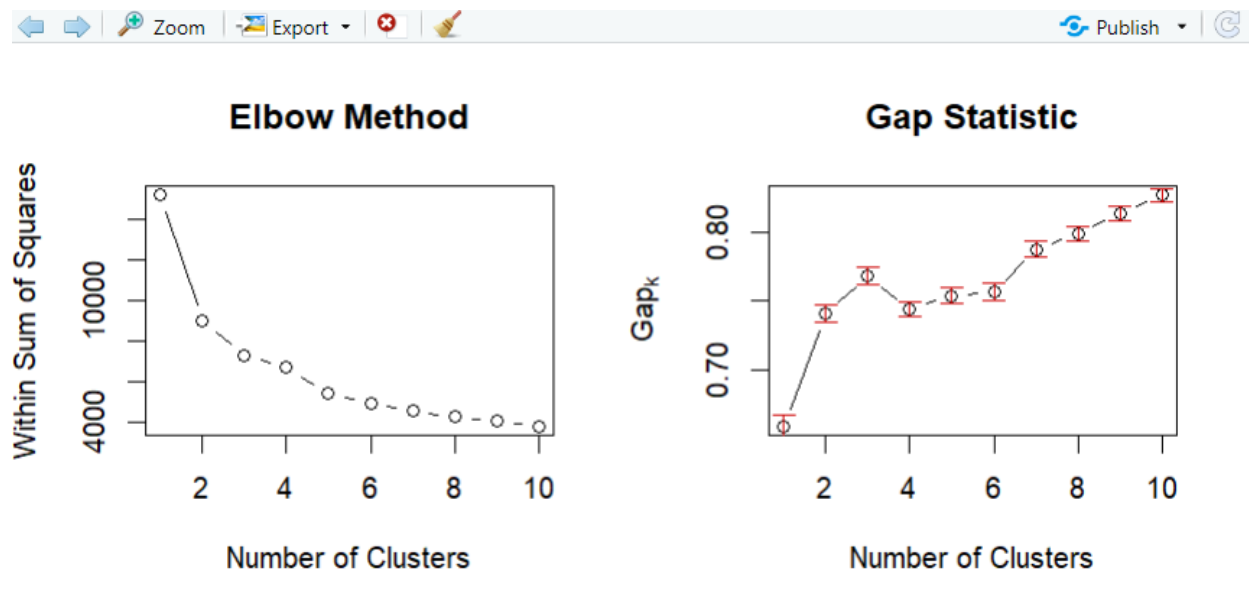
```



Gap statics Method

The ideal number of clusters in a dataset for clustering algorithms can be found using gap statistics, a statistical technique. It contrasts the actual data's within-cluster variation with the variation anticipated by a null reference distribution. Applying a clustering technique on the dataset for a range of cluster numbers, computing the within-cluster variation, and creating a set of reference datasets with a random distribution are the steps required to employ this method. The gap statistic is then derived as the difference between the predicted variance of the reference datasets and the logarithm of the within-cluster variation of the actual data. The number of clusters that maximizes the gap statistic is considered to be the ideal number of clusters.

```
> # Gap statistics
> set.seed(123)
> gap <- clusGap(D_Scaled, FUN = kmeans, nstart = 2, K.max = 10, B = 100)
Clustering k = 1,2,..., K.max (= 10): .. done
Bootstrapping, b = 1,2,..., B (= 100) [one "." per sample]:
..... 50
..... 100
> plot(gap, main = "Gap Statistic", xlab = "Number of Clusters")
>
```

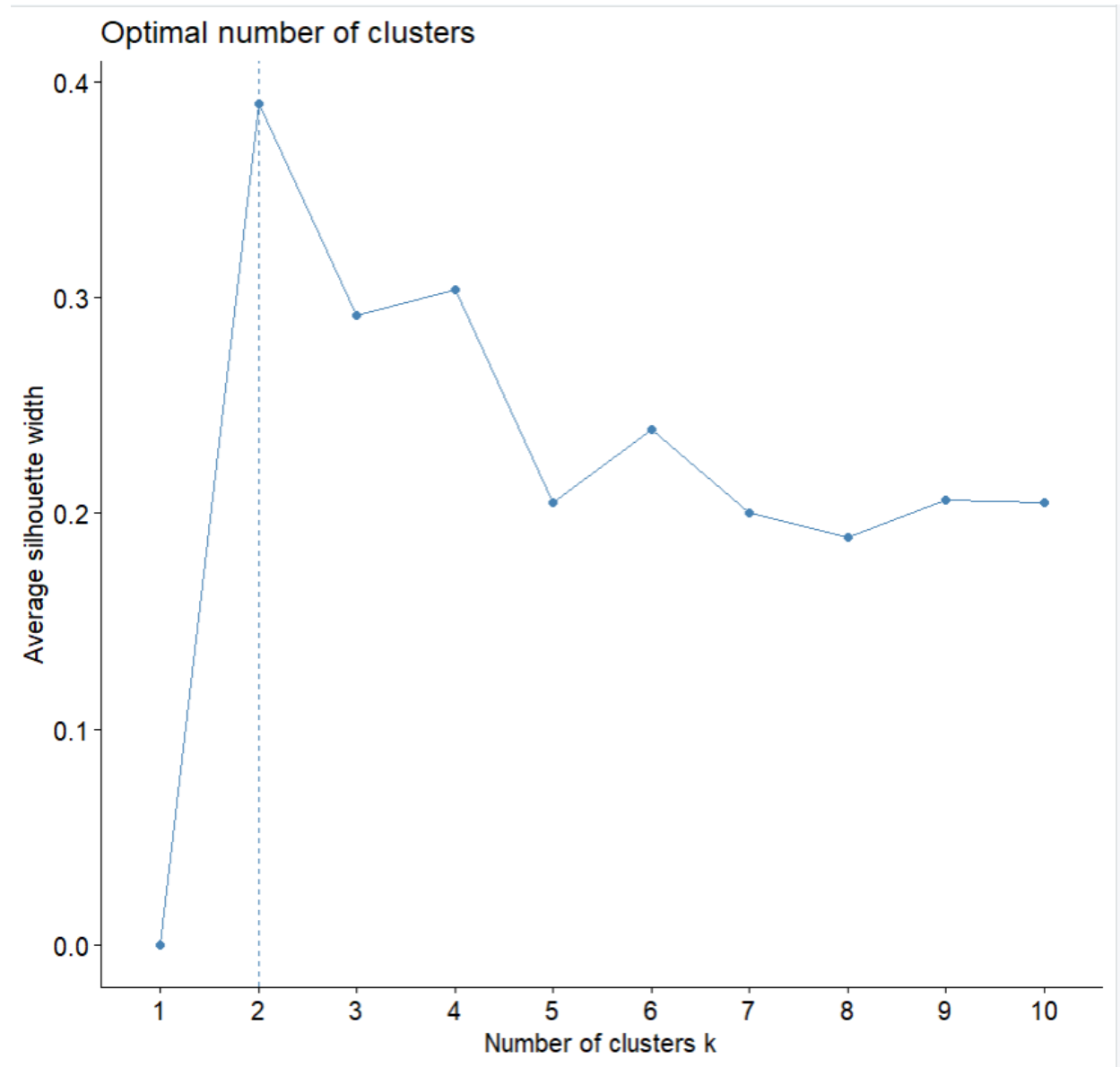


Silhouette method

A approach for assessing the quality of clusters in a dataset is the Silhouette method. For each object, it determines a Silhouette coefficient, which assesses how well the object fits into its designated cluster in relation to other clusters. The ideal number of clusters is then determined by calculating the average Silhouette coefficient for all items for each cluster number. This

approach can identify overlapping or improperly separated clusters and offers a more thorough evaluation of cluster quality.

```
> # Silhouette method  
> sil_ratio <- c()  
> fviz_nbclust(D_Scaled, kmeans, method = 'silhouette')  
> |
```



optimal number of clusters


```
> set.seed(123)
> kmeans_clusters <- kmeans(D_Scaled, centers = 2, nstart = 25)
> kmeans_summary <- as.data.frame(kmeans_clusters$centers)
> kmeans_summary$cluster <- c("Cluster 1", "Cluster 2")
> D_clustered <- datasheet %>% mutate(cluster = as.factor(kmeans_clusters$cluster))
> |
```

Evaluation metrics

```
> # Evaluation metrics
> BSS <- sum(kmeans_clusters$betweenss)
> TSS <- sum(kmeans_clusters$totss)
> WSS <- sum(kmeans_clusters$tot.withinss)
> cat("Ratio of BSS to TSS:", BSS/TSS, "\n")
Ratio of BSS to TSS: 0.4110022
> cat("BSS:", BSS, "\n")
BSS: 6251.343
> cat("WSS:", WSS, "\n")
WSS: 8958.657
> |
```

W1870590

Silhouette plot

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```

BSS: 6251.343
> cat("WSS:", WSS, "\n")
WSS: 8958.657
> # Silhouette plot
> silhouette(kmeans_clusters$cluster, dist(D_Scaled))
  cluster neighbor  sil_width
[1,]      1        2 0.140451608
[2,]      1        2 0.461810445
[3,]      2        1 0.488575080
[4,]      1        2 0.394383678
[5,]      1        2 0.014286921
[6,]      2        1 0.403942965
[7,]      1        2 0.282578045
[8,]      1        2 0.465562093
[9,]      1        2 0.457076992
[10,]     2        1 0.078164075
[11,]     1        2 0.510528647
[12,]     1        2 0.462731287
[13,]     1        2 0.365459685
[14,]     1        2 0.453210784
[15,]     2        1 0.076725970
[16,]     2        1 0.462599423
[17,]     1        2 0.488233428
[18,]     1        2 0.110178692
[19,]     2        1 0.536817836
[20,]     2        1 0.513287875
[21,]     1        2 0.399807555
[22,]     1        2 0.486713032
[23,]     1        2 0.334102310
[24,]     1        2 0.508220687
[25,]     2        1 0.474723652
[26,]     1        2 0.426657813
[27,]     1        2 0.493044366
[28,]     2        1 0.513509277
[29,]     2        1 0.176125423
[30,]     1        2 0.467719919
[31,]     1        2 0.386814008
[32,]     1        2 0.434675999
[33,]     1        2 0.474225224
[34,]     2        1 0.420942638
[35,]     1        2 0.303538231

```

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[33,]	1	2	0.474225224
[34,]	2	1	0.420942638
[35,]	1	2	0.303538231
[36,]	1	2	0.146280592
[37,]	1	2	0.417900152
[38,]	2	1	0.045901447
[39,]	2	1	0.485242692
[40,]	1	2	0.380059164
[41,]	2	1	0.419682851
[42,]	1	2	0.464757500
[43,]	1	2	0.232666538
[44,]	1	2	0.369805428
[45,]	2	1	0.426181704
[46,]	1	2	0.272956175
[47,]	1	2	0.509552601
[48,]	1	2	0.433305435
[49,]	1	2	0.301998607
[50,]	1	2	0.191579099
[51,]	1	2	0.488565708
[52,]	1	2	0.430824119
[53,]	2	1	0.492728492
[54,]	1	2	0.302208052
[55,]	2	1	0.415657057
[56,]	1	2	0.424013817
[57,]	2	1	0.109870699
[58,]	1	2	0.338346445
[59,]	2	1	0.472630851
[60,]	1	2	0.524002017
[61,]	2	1	0.365142555
[62,]	1	2	0.487105264
[63,]	1	2	0.461804470
[64,]	1	2	0.463898179
[65,]	2	1	0.078192682
[66,]	1	2	0.419741482
[67,]	1	2	0.443536158
[68,]	2	1	0.543569786
[69,]	1	2	0.433911094
[70,]	2	1	0.292438715
[71,]	2	1	0.423666231
[72,]	2	1	0.502793261
[73,]	1	2	0.161015619
[74,]	1	2	0.460889467

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[71,]	2	1	0.423666231
[72,]	2	1	0.502793261
[73,]	1	2	0.161015619
[74,]	1	2	0.460889467
[75,]	1	2	0.378094323
[76,]	2	1	0.445340604
[77,]	1	2	0.301219528
[78,]	1	2	0.465171488
[79,]	2	1	0.458852688
[80,]	1	2	0.379879286
[81,]	1	2	0.400101889
[82,]	2	1	0.448969592
[83,]	1	2	0.484386713
[84,]	1	2	0.436690449
[85,]	1	2	0.395254261
[86,]	2	1	0.416592527
[87,]	1	2	0.454464016
[88,]	1	2	0.525811771
[89,]	1	2	0.167909334
[90,]	1	2	0.491322800
[91,]	2	1	0.552114318
[92,]	1	2	0.329263238
[93,]	2	1	0.478089834
[94,]	1	2	0.203998020
[95,]	1	2	0.463642158
[96,]	2	1	0.493051447
[97,]	1	2	0.413445337
[98,]	1	2	0.500004362
[99,]	2	1	0.333117951
[100,]	1	2	0.428610439
[101,]	1	2	0.004580152
[102,]	1	2	0.334162411
[103,]	1	2	0.468550756
[104,]	1	2	0.492972916
[105,]	2	1	0.177570099
[106,]	2	1	0.559430635
[107,]	2	1	0.275413411
[108,]	1	2	0.489059854
[109,]	1	2	0.460591024
[110,]	2	1	0.150164485
[111,]	1	2	0.471485350

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[108,]	1	2	0.489059854
[109,]	1	2	0.460591024
[110,]	2	1	0.150164485
[111,]	1	2	0.471485350
[112,]	1	2	0.285264041
[113,]	1	2	0.429546062
[114,]	1	2	0.431323682
[115,]	1	2	0.369682289
[116,]	1	2	0.189240997
[117,]	2	1	0.376243772
[118,]	2	1	0.517302123
[119,]	1	2	0.456003282
[120,]	2	1	0.019086890
[121,]	1	2	0.469132105
[122,]	1	2	0.296708488
[123,]	1	2	0.381283992
[124,]	1	2	0.419695384
[125,]	1	2	0.464565423
[126,]	1	2	0.274978822
[127,]	1	2	0.496683749
[128,]	1	2	0.362202451
[129,]	1	2	0.232026068
[130,]	2	1	0.409326346
[131,]	2	1	0.355879125
[132,]	1	2	0.387138379
[133,]	2	1	0.094622896
[134,]	1	2	0.469437249
[135,]	2	1	0.435815372
[136,]	2	1	0.044946700
[137,]	1	2	0.322199616
[138,]	1	2	0.491429289
[139,]	1	2	0.494669955
[140,]	1	2	0.448852828
[141,]	1	2	0.331289100
[142,]	1	2	0.442184213
[143,]	2	1	0.370048359
[144,]	1	2	0.433721682
[145,]	1	2	0.155550893
[146,]	2	1	0.525594947
[147,]	1	2	0.122254252
[148,]	1	2	0.456242279
[149,]	1	2	0.276582438

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[146,]	2	1	0.525594947
[147,]	1	2	0.122254252
[148,]	1	2	0.456242279
[149,]	1	2	0.276582438
[150,]	1	2	0.424262592
[151,]	2	1	0.343680514
[152,]	2	1	0.075498621
[153,]	1	2	0.331205293
[154,]	2	1	0.469452083
[155,]	1	2	0.504191996
[156,]	2	1	0.440397581
[157,]	1	2	0.425403218
[158,]	1	2	0.518724039
[159,]	1	2	0.415487660
[160,]	1	2	0.348510777
[161,]	1	2	0.454084816
[162,]	2	1	0.394378728
[163,]	1	2	0.241499338
[164,]	1	2	0.541857399
[165,]	2	1	0.477263928
[166,]	2	1	0.439851339
[167,]	1	2	0.340956674
[168,]	2	1	0.423592387
[169,]	1	2	0.424777936
[170,]	1	2	0.477253114
[171,]	2	1	0.487139609
[172,]	2	1	0.384637548
[173,]	1	2	0.345200401
[174,]	2	1	0.393065445
[175,]	1	2	0.235571935
[176,]	1	2	0.357558274
[177,]	1	2	0.172617800
[178,]	1	2	0.433508496
[179,]	1	2	0.389664149
[180,]	1	2	0.489103350
[181,]	2	1	0.495778234
[182,]	1	2	0.491251577
[183,]	1	2	0.374673773
[184,]	1	2	0.477128876
[185,]	2	1	0.448241783
[186,]	1	2	0.084308552

RStudio

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Source

Console Terminal Background Jobs

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[183,]	1	2	0.374673773
[184,]	1	2	0.477128876
[185,]	2	1	0.448241783
[186,]	1	2	0.084308552
[187,]	1	2	0.216363384
[188,]	1	2	0.261491324
[189,]	2	1	0.520327400
[190,]	1	2	0.408165141
[191,]	2	1	0.164845086
[192,]	1	2	0.453800682
[193,]	1	2	0.395642539
[194,]	2	1	0.453357549
[195,]	1	2	0.391921475
[196,]	1	2	0.460164189
[197,]	2	1	0.482655250
[198,]	1	2	0.418018168
[199,]	1	2	0.409295225
[200,]	1	2	0.448006071
[201,]	1	2	0.264847658
[202,]	1	2	0.391236808
[203,]	2	1	0.452889420
[204,]	2	1	0.540261835
[205,]	1	2	0.463660143
[206,]	1	2	0.391735434
[207,]	1	2	0.266028846
[208,]	1	2	0.435468428
[209,]	1	2	0.442222712
[210,]	2	1	0.440375686
[211,]	1	2	0.165037730
[212,]	1	2	0.529413108
[213,]	1	2	0.452408524
[214,]	2	1	0.342907322
[215,]	1	2	0.513390329
[216,]	1	2	0.356841329
[217,]	1	2	0.439943322
[218,]	2	1	0.525104005
[219,]	1	2	0.468255634
[220,]	1	2	0.444937319
[221,]	2	1	0.499102761
[222,]	1	2	0.356988401
[223,]	2	1	0.246551205
[224,]	1	2	0.427629913

RStudio

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Go to file/function Addins

Source

Console Terminal x Background Jobs x

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[221,]	2	1	0.499102761
[222,]	1	2	0.356988401
[223,]	2	1	0.246551205
[224,]	1	2	0.427629913
[225,]	1	2	0.540339103
[226,]	1	2	0.434217442
[227,]	2	1	0.492249936
[228,]	1	2	0.485716700
[229,]	2	1	0.515214600
[230,]	1	2	0.485147639
[231,]	1	2	0.463130430
[232,]	1	2	0.436980095
[233,]	1	2	0.417894380
[234,]	2	1	0.478227879
[235,]	1	2	0.476581580
[236,]	1	2	0.275473611
[237,]	1	2	0.281102133
[238,]	1	2	0.426663548
[239,]	2	1	0.542894269
[240,]	1	2	0.400204515
[241,]	1	2	0.386822002
[242,]	1	2	0.281142168
[243,]	1	2	0.450901436
[244,]	2	1	0.539761476
[245,]	1	2	0.350704691
[246,]	1	2	0.489289116
[247,]	1	2	0.329806180
[248,]	1	2	0.307689268
[249,]	2	1	0.551347522
[250,]	1	2	0.502776398
[251,]	1	2	0.481886928
[252,]	2	1	0.526828587
[253,]	1	2	0.390958752
[254,]	1	2	0.391671963
[255,]	1	2	0.278598249
[256,]	2	1	0.437145691
[257,]	1	2	0.434920078
[258,]	1	2	0.352803563
[259,]	2	1	0.417743180
[260,]	2	1	0.498398484
[261,]	1	2	0.503301414

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Go to file/function Addins

Source

Console	Terminal x	Background Jobs x
R 4.3.0 · ~/		
[258,]	1	2 0.352803563
[259,]	2	1 0.417743180
[260,]	2	1 0.498398484
[261,]	1	2 0.503301414
[262,]	1	2 0.200108255
[263,]	1	2 0.505291922
[264,]	1	2 0.459729742
[265,]	2	1 0.274366733
[266,]	1	2 0.447904278
[267,]	1	2 0.435832636
[268,]	1	2 0.055512964
[269,]	1	2 0.460515258
[270,]	1	2 0.500299997
[271,]	1	2 0.408723872
[272,]	2	1 0.054383234
[273,]	1	2 0.296245126
[274,]	1	2 0.337553646
[275,]	2	1 0.424302188
[276,]	1	2 0.373227204
[277,]	1	2 0.452064044
[278,]	1	2 0.514519117
[279,]	2	1 0.503506108
[280,]	1	2 0.370580996
[281,]	2	1 0.078812427
[282,]	1	2 0.480591666
[283,]	1	2 0.410125453
[284,]	2	1 0.469347935
[285,]	1	2 0.352702746
[286,]	1	2 0.067632704
[287,]	1	2 0.524061513
[288,]	1	2 0.420278234
[289,]	2	1 0.376155681
[290,]	1	2 0.452031221
[291,]	1	2 0.505806264
[292,]	2	1 0.012467862
[293,]	1	2 0.460333897
[294,]	1	2 0.471458283
[295,]	2	1 0.423246900
[296,]	1	2 0.419768539
[297,]	2	1 0.352535417
[298,]	1	2 0.476349782
[299,]	1	2 0.059712917

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

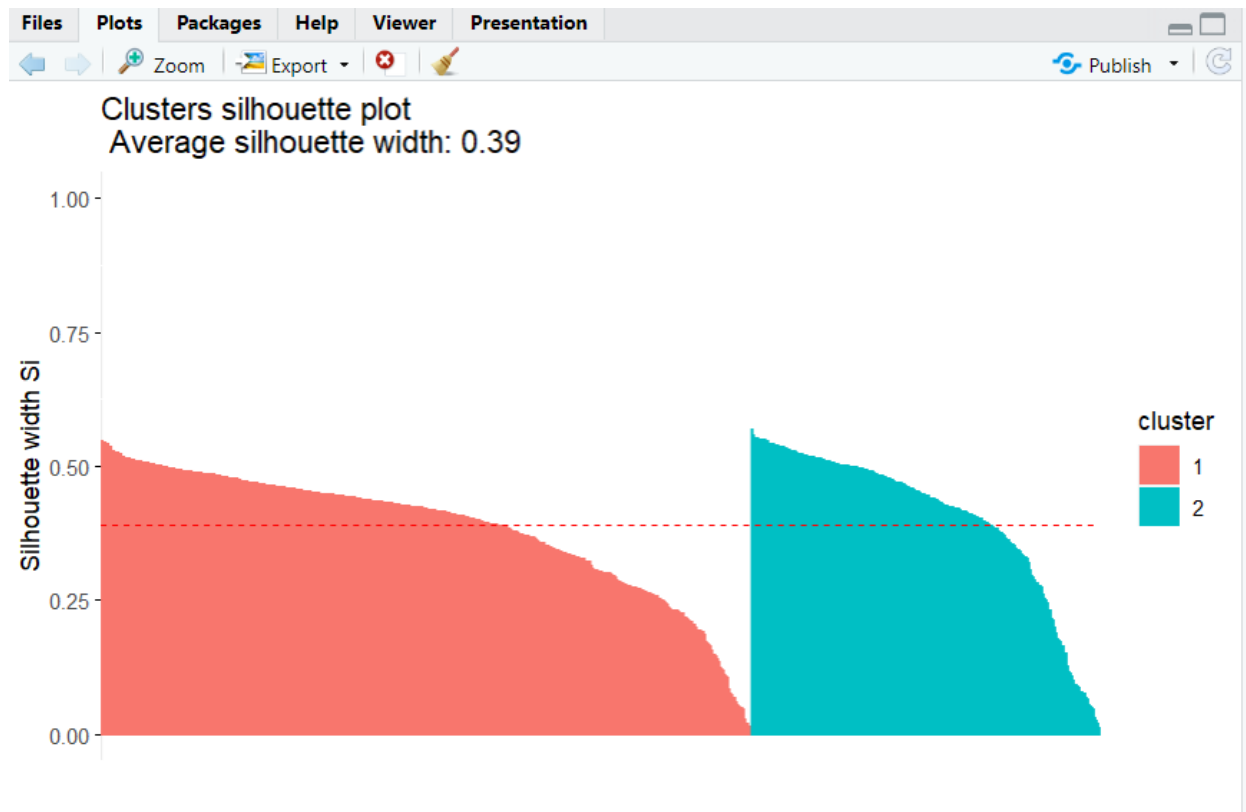
Go to file/function Addins

Source

Console Terminal x Background Jobs x

R 4.3.0 · ~/

```
[296,]      1      2 0.419768539
[297,]      2      1 0.352535417
[298,]      1      2 0.476349782
[299,]      1      2 0.059712917
[300,]      1      2 0.217806077
[301,]      2      1 0.551678228
[302,]      1      2 0.352908642
[303,]      1      2 0.323180082
[304,]      1      2 0.435537031
[305,]      1      2 0.439116337
[306,]      1      2 0.454712579
[307,]      2      1 0.470196604
[308,]      2      1 0.483642191
[309,]      2      1 0.362875139
[310,]      2      1 0.396967645
[311,]      2      1 0.231127743
[312,]      1      2 0.280059611
[313,]      1      2 0.510664672
[314,]      2      1 0.465244748
[315,]      1      2 0.435847723
[316,]      1      2 0.484276554
[317,]      1      2 0.528094314
[318,]      2      1 0.172197021
[319,]      1      2 0.487526213
[320,]      2      1 0.451245756
[321,]      2      1 0.527940508
[322,]      2      1 0.424123328
[323,]      1      2 0.441558414
[324,]      2      1 0.357725846
[325,]      1      2 0.227051629
[326,]      1      2 0.428409377
[327,]      2      1 0.203832082
[328,]      1      2 0.445141045
[329,]      1      2 0.338752856
[330,]      1      2 0.421301411
[331,]      1      2 0.268790899
[332,]      2      1 0.378905084
[333,]      2      1 0.537032725
[ reached getOption("max.print") -- omitted 513 rows ]
attr(,"Ordered")
[1] FALSE
```



Dimensionality Reduction with PCA

```

R 4.3.0 · ~/
> # Perform PCA
> D_quality <- Data_N_outliers
> D_pca <- prcomp(D_Scaled)
> summary(D_pca)
Importance of components:
      PC1      PC2      PC3      PC4      PC5      PC6      PC7      PC8      PC9
Standard deviation 3.0705 1.7386 1.3779 1.08728 0.95400 0.73033 0.59669 0.46989 0.39755
Proportion of Variance 0.5238 0.1679 0.1055 0.06568 0.05056 0.02963 0.01978 0.01227 0.00878
Cumulative Proportion 0.5238 0.6917 0.7972 0.86287 0.91344 0.94307 0.96285 0.97511 0.98389
      PC10     PC11     PC12     PC13     PC14     PC15     PC16     PC17     PC18
Standard deviation 0.30218 0.2508 0.20940 0.18721 0.14610 0.12646 0.11355 0.07870 0.01904
Proportion of Variance 0.00507 0.0035 0.00244 0.00195 0.00119 0.00089 0.00072 0.00034 0.00002
Cumulative Proportion 0.98897 0.9925 0.99490 0.99685 0.99803 0.99892 0.99964 0.99998 1.00000
> D_pca.transform <- as.data.frame(-D_pca$x[, 1:10])
> D_pca.transform
      PC1      PC2      PC3      PC4      PC5      PC6      PC7
1  0.33008671 -0.214358296 -0.996691160 0.17188815 0.082605091 -0.726465844 -0.91461297
2 -1.59322354 -0.421883157 0.368657675 0.23214677 0.691903545 -0.528331092 0.35646058
3  3.75905753 0.187893015 -0.088626688 1.20046225 0.718805359 0.704361537 -0.01030921
4 -1.74165971 -2.821701010 -0.112458943 0.37385893 -0.368148386 -0.506501111 0.45118004
5  0.55099709 4.765977863 -11.679347547 0.16643526 3.242122566 -0.298604457 2.66105177
6  6.39853584 3.925694674 2.067951954 -0.38404114 -0.571226699 0.899195611 0.77730594
7 -0.77669551 -2.203235795 0.123030908 2.10964871 -0.212144346 0.777780808 -0.12203096
8 -2.14105741 -1.166269567 -0.651455750 0.81825415 -1.242264794 -0.657561654 -0.16034312
9 -4.45658043 -3.086803124 -0.104909036 -0.55053903 -0.571267072 -0.304295061 0.26871082
10 1.17001772 -1.996724845 -0.175795805 -0.67893639 -0.445301995 -0.140904054 0.48855979
11 -3.49477750 -1.758952296 -0.291257851 -0.42223407 -0.684765229 -0.324841271 0.47611904
12 -4.38320711 -2.424415301 0.774102875 -1.30578452 0.850273985 -0.053265511 0.32667541
13 -0.82545521 0.029209348 -0.445542089 0.29574946 -0.570224147 -0.031753070 -1.31812479
14 -1.41227892 -0.018430581 0.119183043 0.14372362 0.505877154 -0.642632229 0.26870471
15 1.19863657 -0.861076457 -0.536371241 1.18511362 -1.399177178 0.828170988 -1.17255668
16 3.79704179 1.295440361 -0.172146344 1.02505361 -1.119105668 -0.132572652 -0.70562160
17 -5.27747851 1.737605710 0.549555431 -0.86021046 0.158967860 -0.104805385 0.78667883
18 0.34588877 -1.593432266 -0.500656283 -0.40875915 -0.798262118 1.697273095 0.25736716
19 4.15858451 1.155933921 0.405031590 0.11053613 -0.489808556 -0.528537997 0.20508500
20 4.21076898 1.059629390 -0.593377459 0.72816543 -0.863905083 0.111037589 -0.42879745
21 -1.37701833 1.188201912 -0.328298736 0.06760925 -1.663956716 -0.240602060 -0.80455415
22 -5.10479548 2.260031736 -0.070717396 0.58683746 -0.075240822 0.895736601 0.29751049
23 -1.15871891 -2.050370801 -0.595831487 1.70168523 -0.770586984 0.556565573 -0.42291384
24 -2.75681565 -0.853744312 -0.146857237 0.07871385 -1.501249949 0.016529997 0.34957582
25 3.85713290 -0.401090743 -0.555210410 -0.00318032 -1.797477794 -0.586226026 -0.24779055

```

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22	-5.10479548	2.260031736	-0.070717396	0.58683746	-0.075240822	0.895736601	0.29751049
23	-1.15871891	-2.050370801	-0.595831487	1.70168523	-0.770586984	0.556565573	-0.42291384
24	-2.75681565	-0.853744312	-0.146857237	0.07871385	-1.501249949	0.016529997	0.34957582
25	3.85713290	-0.401090743	-0.555210410	-0.00318032	-1.797477794	-0.586226026	-0.24779055
26	-1.44672712	-0.016461828	-0.607235427	0.25049711	-1.383243726	-0.900679099	-0.38963485
27	-4.82856111	2.351690378	0.041113514	0.22011134	-0.053902843	0.755735582	0.44069768
28	4.59574320	0.140063696	-0.018160246	-1.69690838	0.029185716	-0.662545258	-0.15978039
29	1.54126791	-0.290080767	0.089666219	-0.01503813	-0.626356138	1.754587627	0.65063381
30	-3.21085890	2.204527239	0.323430804	-0.79119315	1.408821394	0.478766777	-0.25139872
31	-0.89165586	0.306095790	-0.676669881	0.21776713	-0.256248534	-0.777402054	-0.08048598
32	-2.01951090	-2.173003600	0.430173510	0.96629311	0.220384861	-0.626962133	-0.17906560
33	-3.01178099	-2.404288303	0.469415437	-0.38692089	-0.241362103	0.550823147	0.62307919
34	3.91370687	-1.114955675	-0.287043371	-2.29904555	1.255243956	0.007728381	-0.79591582
35	-1.16809525	-2.771683406	0.099808788	-2.05155331	-0.158977981	0.132409835	0.03348118
36	0.16272752	-1.205959591	-0.082277260	2.20822795	1.069596502	0.046329835	-0.21946839
37	-2.24245205	2.912416275	0.548085272	0.11086620	-0.995190764	-0.493626740	0.20942363
38	3.05995499	2.233875180	-13.750911492	-0.88247688	1.419253060	0.098634574	0.66414896
39	4.02169457	1.022518756	0.931121336	-0.28752496	-1.168966043	-0.725503471	0.95967244
40	-1.57144876	1.362050724	-1.132814744	1.12867923	-0.870948672	0.574469589	-1.51989870
41	2.95681526	0.642281618	-0.502465105	-0.32278971	-0.592938009	0.843267642	-0.20773379
42	-5.32641418	2.399977400	0.650299672	-0.35624969	1.276910832	0.342270933	0.54301642
43	-0.21797670	1.405048146	0.375257980	0.78975425	-0.442687716	-1.595287861	0.58752127
44	-1.10340896	-1.214077014	-0.234077691	-0.36146137	-0.743673233	1.030031494	0.68796843
45	5.23303708	-1.012502417	0.297129596	3.06444310	0.914277904	0.491685928	0.80860658
46	-0.39967124	1.464123755	-0.215137888	0.38641024	-0.928216071	-1.215021580	0.49150793
47	-2.76556752	0.007220952	0.384999187	0.06867854	-0.948463332	-1.067350852	0.38433741
48	-2.57854760	3.113261106	0.428419644	-0.41196628	0.557721669	0.231714003	-0.08065196
49	-0.48881672	1.186709672	-0.573122060	0.35013351	0.158593994	-1.305313378	-0.51150869
50	-0.13789740	-2.308104548	0.157813233	-0.18906932	-1.549805820	-0.913210015	-0.09373896
51	-3.85656480	2.099877910	0.764985351	-0.44175241	-0.822830104	0.117635631	0.42745909
52	-1.79957607	-0.916236926	-0.375539120	-1.56615340	0.495838111	0.712755576	-0.14948658
53	4.52401983	0.591050179	-0.732275575	0.09792532	-1.748703054	-0.011290308	-0.96589101
54	-1.11048194	-3.050015888	-0.121515624	0.72346585	-0.449537427	0.518720546	-0.17764643
55	6.03158515	3.302342949	1.811665156	-1.93081742	0.542330414	0.742002904	0.06914396
56	-3.10554406	-3.214139749	0.084175871	1.24413524	-0.351468069	0.158282871	0.90541923
57	1.20555089	-1.057956986	-0.005386846	-0.26526507	-0.813771679	0.418833766	0.53224650
58	-0.83818681	1.472776142	0.304435226	0.35397832	-0.150071093	-1.621870681	0.28661878
59	4.16787231	-0.639576293	-0.369489903	0.29930007	-2.094560207	-0.190331897	0.53877507
60	-4.02339031	1.397150443	1.006422303	0.37592917	0.328092146	-0.118339042	0.24666807
61	2.81501708	-1.895783141	0.104495261	0.67678642	-1.227315676	-0.024223560	0.34241859
62	-4.19759767	1.199638415	0.251300681	-0.15382439	1.893740600	0.471896506	-0.96468929
63	-3.03331630	1.778701635	0.908529352	0.26521327	1.230880571	-0.003856559	1.20789033

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66	-2.22085752	1.104624631	0.648634279	1.76364252	1.524861523	0.778923932	0.29410453
67	-2.40270691	2.713388023	0.086259703	-0.60599310	0.756494408	-0.463466705	-0.52520280
68	4.90976395	0.808828345	-0.575225081	-0.07168510	-0.308766165	-0.201262609	-0.61504679
69	-1.59876007	-0.707764659	-1.008164157	-0.17924636	-0.966867569	-0.855189542	-0.79905018
70	2.28994955	-0.352886739	-1.040916047	0.31311230	-1.203818894	1.172899890	-0.91260299
71	3.59690171	1.949081382	0.906397048	0.20266625	-1.093589232	-1.022034861	0.37242155
72	5.49324077	0.282838004	-0.360273159	-0.13751982	0.860814076	-0.990898835	-1.49246777
73	0.09878185	-1.524073223	0.065141386	-1.72915890	0.101631400	0.441430330	0.06649150
74	-5.27030626	2.760873997	0.188779779	-1.24916793	-0.007170271	-0.396001063	-0.45776668
75	-1.16510620	-0.405484304	-0.120519669	-0.17609162	1.646936376	-0.861766181	-0.18608130
76	3.02744946	-0.299999575	0.113507326	0.71785593	-1.091370898	-0.306891956	-0.10004913
77	-0.63482241	-0.018371674	-0.008139357	-1.24077517	-0.998037958	0.706607832	0.70721815
78	-2.67999494	2.364083891	-0.459893688	-0.13333343	-0.881151553	0.667303743	0.03541521
79	3.72504704	0.555840978	0.389952118	1.25426491	-0.380326512	1.563782077	-0.32655804
80	-2.15450951	3.557778097	0.744276184	0.48006264	-0.004762879	-0.174856548	0.48520682
81	-1.37238585	1.055198125	-0.425374477	0.64552976	-1.611948328	0.078648883	-0.78191495
82	3.37635619	1.058748050	0.274331638	-0.21352922	-0.718524450	1.288261979	-0.52670223
83	-3.83606125	-1.273252347	0.659376088	0.08861005	-1.302099094	-0.072939616	1.50673279
84	-2.64501635	0.865119627	0.605703814	-1.58725541	1.774483566	-0.538524839	-0.60664314
85	-1.53373322	-1.096394171	-1.564344846	-0.19406853	-0.912492694	1.052606945	-0.64458896
86	6.76702609	3.667714342	2.219229021	-0.17235008	0.246150690	1.061355556	0.68734665
87	-1.56911901	-0.343999529	0.173045700	-0.66651314	0.200129324	-0.944361241	0.07991113
88	-4.86546477	0.512946298	0.691108443	-0.23607198	-0.612290982	-0.311392987	0.41066857
89	0.05027942	-1.832498751	-1.140863209	-0.54855870	-0.938666173	1.216253049	-0.89267101
90	-4.98265862	2.299489520	0.272050225	0.28523393	0.381047960	0.490568076	0.31565849
91	4.95427838	0.202624867	-0.515643197	0.04640846	-1.032716695	-0.416902950	-0.26729233
92	-0.83209894	-0.605699876	0.702429753	-0.40671715	0.728234559	0.687389924	0.94516029
93	3.35707158	-0.055363119	0.715399574	0.96228200	0.050905565	-0.064463549	-0.01964542
94	0.06151440	0.790955692	-0.476361026	-0.09142817	-0.054420715	-1.163917489	-0.39380543
95	-2.07767545	1.306014243	-0.381956564	0.32313804	-1.323338519	-0.015416223	-0.85538971
96	3.99869127	-0.011584744	-0.004362434	-1.38395733	0.959126309	0.058899592	-0.76115842
97	-1.82752543	0.663529572	0.369236763	1.23893022	1.736389416	0.597026452	-0.64425562
98	-1.85177326	0.599374706	-0.221160988	-0.04954649	-0.545926744	0.158206871	-0.33164709
99	3.01390385	-1.320592191	-0.287997344	-2.35785450	1.197497615	0.109995104	-1.18602002
100	-2.40048449	3.163859481	0.269833393	-0.40680567	0.090163300	0.065347115	-0.07917751
PC8 PC9 PC10							
1	-0.400214338	-0.874065893	-0.097489202				
2	0.251006040	0.129983525	0.089554116				
3	0.466635984	-0.317045735	0.470495718				
4	-0.018486013	-0.035221243	-0.459144754				
5	-0.404142379	2.100507107	0.362529057				
6	0.184222225	0.068502826	0.426800687				

```

R 4.3.0 ~ /
2 0.21100000 0.12550000 0.00000000
3 0.466635984 -0.317045735 0.470495718
4 -0.018486013 -0.035221243 -0.459144754
5 -0.404142379 2.100507107 0.362529057
6 -0.184232335 0.968592826 0.426800687
7 -0.406425878 0.341578800 0.066457130
8 -0.423144489 0.004864284 0.059177489
9 -0.378198246 0.418893513 -0.206113855
10 0.992707363 0.193445702 -0.273979262
11 0.283220367 0.139816267 -0.025911626
12 -0.398428701 0.098556980 0.505846716
13 -0.201106667 -0.054330477 0.246228901
14 0.538564946 0.067047703 0.080985202
15 -0.506002548 0.394362031 0.004514205
16 0.483798790 0.201751661 0.550348072
17 -0.871223642 -0.598050702 0.043534610
18 -0.472160626 0.165223037 -0.344163719
19 -0.214810847 -0.174878534 -0.110745245
20 0.074761317 -0.435128171 -0.032971009
21 0.199559980 0.081197653 0.090651008
22 0.031802073 -0.644474478 -0.124904179
23 -0.678632926 0.486455260 0.012001102
24 0.223424299 0.155785147 -0.136415456
25 0.303315616 0.004822282 0.472309690
26 0.334984698 0.092030653 0.040924559
27 0.212316913 -0.358126325 -0.069027920
28 -0.819700206 -0.319421184 0.125335986
29 -0.372387379 0.212633648 -0.390817374
30 0.477125268 0.510855102 -0.070138785
31 0.506387389 -0.338862133 0.089266293
32 0.108719009 0.680084894 0.094872195
33 -0.143697282 0.021638624 -0.107231526
34 0.001553432 -0.144653925 0.436100119
35 0.706439989 0.599239225 -0.096617135
36 0.029517444 -0.709400987 0.300762015
37 0.389670053 0.552012682 -0.088999445
38 -0.874189275 0.643071178 0.567636806
39 -0.462071711 0.338696315 0.127473882
40 0.558913123 -0.347353822 -0.237484481
41 1.029713484 -0.173474234 0.243754429
42 -0.881897166 -0.485915652 -0.207248608
43 -0.045511305 -0.134901889 0.060133331

```



```

R 4.3.0 . ~/
46 -0.172235905 -0.085192746 0.106566638
47 -0.614871418 -0.099823738 0.251287892
48 -0.343786616 0.470526138 -0.002872660
49 -0.198231433 -0.178780198 -0.254356164
50 -0.090044196 0.884103999 0.244217106
51 0.941515700 0.430153150 -0.162974762
52 -0.065834996 -0.562713496 -0.226001415
53 0.116647642 0.006827807 0.463357994
54 -1.429132039 0.250817093 0.168469965
55 0.086381832 0.724768890 -0.407824563
56 -0.249828513 0.141460348 -0.350117829
57 0.833972071 -0.360571492 0.076075630
58 0.281308836 -0.165936045 -0.215037718
59 0.034698887 -0.578600513 0.407387723
60 0.307773462 -0.417758393 -0.015768001
61 -0.066027944 0.027758633 -0.055836373
62 0.483978629 0.037787753 -0.456535267
63 -0.553390339 -0.524138453 0.040193319
64 0.388791178 -0.294690415 -0.511054641
65 0.206716909 0.273801694 -0.029185133
66 0.758961352 -0.184745977 -0.088686787
67 0.098973697 0.300476127 -0.239286356
68 0.152687923 -0.069050874 -0.762986930
69 -0.043248769 -0.042145751 0.137750652
70 -0.565053368 0.092007197 0.100469158
71 0.185588382 0.755972574 -0.364935643
72 0.232372142 -0.434973037 -0.979674668
73 0.736098355 -0.006877523 0.042965762
74 0.396826275 0.072582877 -0.135708684
75 0.479727876 -0.028904811 0.089589596
76 0.174185610 -0.264983281 0.363796286
77 0.755237486 0.066882528 0.076253109
78 0.096852945 -0.133757143 0.235845105
79 -0.404773228 0.133134960 -0.078973508
80 -0.437475961 0.457192777 0.230503006
81 0.014497926 -0.199251472 0.239978873
82 -0.363347260 0.400436651 -0.110675665
83 -0.313487367 -0.208073930 0.060690747
84 0.877881780 0.205289689 -0.222131642
85 0.043709677 -0.430906320 -0.205181288
86 -0.267473209 0.564395476 0.200483617
87 0.128099318 0.170973983 0.146380018
88 -0.275688572 -0.415958455 0.039636385
89 0.211610013 -0.121529113 -0.143601982
90 0.573201572 -0.313923339 -0.241439153
91 -0.056699583 -0.365312355 0.184004979
92 0.839280997 0.270990118 0.465741959
93 -0.073716851 0.239172452 0.903717839
94 -0.381188303 -0.565584935 0.032806248
95 0.126580977 -0.319181644 0.020387786
96 0.518848006 0.196283036 0.413013926
97 0.512552468 -0.470989556 0.166488244
98 -0.037498101 0.304608620 -0.148718765
99 0.383769873 0.121240996 0.497759928
100 -0.543689596 0.091948213 -0.131718380
[ reached 'max' / getOption("max.print") -- omitted 746 rows ]
>

```

K2 cluster

```

> #k2 cluster
> k=2
> kmeans_clustered.pca = kmeans(D_pca.transform, centers = k, nstart = 10)
> kmeans_clustered.pca
K-means clustering with 2 clusters of sizes 295, 551

Cluster means:
      PC1      PC2      PC3      PC4      PC5      PC6      PC7
1  3.712440  0.05517768  0.09037270 -0.04544055  0.004748502  0.04182875  0.03018973
2 -1.987604 -0.02954159 -0.04838466  0.02432843 -0.002542301 -0.02239470 -0.01616328
      PC8      PC9      PC10
1 -0.007121053 -0.02886359  0.016831490
2  0.003812542  0.01545328 -0.009011415

Clustering vector:
[1] 2 2 1 2 2 1 2 2 2 1 2 2 2 2 1 1 2 2 2 1 2 2 1 1 2 2 2 2 1 2 2 1 1 2 2 2 2 1 2 2 2 1 2 2 2 1 1 2 1 2 2 2 1
[46] 2 2 2 2 2 2 2 1 2 1 2 1 2 1 2 2 2 1 2 2 1 2 2 1 1 1 2 2 2 1 2 2 1 2 2 1 2 2 2 1 2 2 2 1 2 2 2 1 2 2 2 2
[91] 1 2 1 2 2 1 2 2 1 2 2 2 2 2 1 1 1 2 2 1 2 2 2 2 2 2 1 1 2 1 2 2 2 2 2 2 2 2 2 1 1 2 1 2 1 2 1
[136] 1 2 2 2 2 2 2 1 2 2 1 2 2 2 2 1 2 1 2 1 2 2 2 2 2 1 2 2 1 1 2 1 2 2 1 1 2 1 2 2 2 2 2 2 2 2
[181] 1 2 2 2 1 2 2 2 1 2 1 2 2 1 2 2 2 1 2 2 2 2 2 1 2 2 2 2 2 1 2 2 2 1 2 2 2 1 2 2 1 2 1 2 2 2
[226] 2 1 2 1 2 2 2 2 1 2 2 2 2 1 2 2 2 2 1 2 2 2 2 1 2 2 2 1 2 2 2 1 2 2 2 1 2 2 2 2 1 2 2 2 2 2
[271] 2 1 2 2 1 2 2 2 1 2 1 2 2 1 2 2 2 2 1 2 2 1 2 2 1 2 1 2 2 2 1 2 2 2 2 2 1 1 1 1 1 1 2 2 1 2
[316] 2 2 1 2 1 1 1 2 1 2 2 2 1 2 2 2 2 1 1 2 1 1 2 2 2 2 2 1 1 1 1 2 2 2 1 2 2 2 1 2 2 1 2 2 1 2 1
[361] 1 1 2 2 2 1 2 2 2 2 2 2 2 2 2 1 1 2 2 1 2 1 2 1 2 2 2 1 2 1 1 2 2 2 2 1 2 1 2 2 2 1 2 1 2 1 2 1
[406] 2 2 2 2 2 2 2 2 2 2 1 2 2 2 2 2 1 2 2 2 2 1 2 1 2 1 1 2 2 1 2 2 2 1 1 1 2 2 1 1 2 1 1 2 1 1 2
[451] 2 2 2 2 1 2 2 2 1 2 2 1 2 2 1 2 2 1 1 2 2 1 1 2 1 1 2 1 1 2 1 2 2 2 2 1 2 2 1 1 2 2 1 2 2 1 1
[496] 2 1 2 2 1 1 1 2 2 1 1 1 2 2 1 2 2 1 2 2 2 2 2 1 2 2 2 2 2 1 2 1 1 2 2 1 1 1 2 2 2 1 2 1 1
[541] 2 2 2 2 2 2 2 2 2 1 2 2 2 2 1 1 1 2 2 1 1 2 2 2 1 1 2 2 2 1 1 1 1 2 2 2 2 1 1 1 2 2 2 2 1 1 1
[586] 2 1 2 2 1 2 2 2 2 2 1 2 2 2 2 2 2 1 2 2 1 2 2 2 2 1 2 2 2 2 2 2 2 2 2 1 1 2 2 1 2 1 2 2 1 2 2
[631] 2 1 2 1 2 2 2 2 1 1 1 2 1 2 2 2 2 2 1 2 1 2 2 2 1 2 2 2 2 1 2 1 2 2 2 2 1 1 2 2 1 2 2 1 2 2 2
[676] 1 2 1 2 2 2 2 2 1 2 2 2 1 2 2 1 2 2 1 2 2 2 2 2 1 1 2 2 1 1 2 2 2 2 1 1 1 1 2 1 2 2 1 2 1 1
[721] 2 1 2 1 2 2 1 2 2 1 1 1 2 1 2 2 1 1 1 2 2 1 2 2 2 2 2 1 2 2 2 2 2 2 1 2 2 2 1 2 2 2 1 1 2 1 1
[766] 2 2 2 1 2 2 1 1 2 2 2 1 1 2 1 2 1 1 2 2 1 2 2 2 2 1 1 2 2 2 1 1 2 2 2 2 2 2 1 1 2 2 1 2 2 2
[811] 1 2 2 2 2 2 2 1 1 1 2 1 2 1 1 2 1 1 2 2 2 2 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 1 2 2 2 2 2

```

```

Within cluster sum of squares by cluster:
[1] 3113.806 5678.106
(between_SS / total_SS = 41.6 %)

Available components:

[1] "cluster"      "centers"      "totss"        "withinss"     "tot.withinss" "betweenss"
[7] "size"         "iter"         "ifault"
> |

```

Finding BSS and WSS

```

> #CM
> wss_ratio = kmeans_clustered.pca$tot.withinss
> bss = kmeans_clustered.pca$betweenss
> print(paste("Total within-cluster sum of square is", wss_ratio))
[1] "Total within-cluster sum of square is 8791.91164569894"
> print(paste("Between Sum of Squares is", bss))
[1] "Between Sum of Squares is 6250.27506960953"
> |

```


Energy Forecasting Part

1st Subtask Objectives

A)

Electricity load forecasting is critical to assuring the reliability of power systems. Because of its capacity to describe complicated non-linear interactions between inputs and outputs, MLP models have been frequently employed in this domain. Defining the input vector for MLP models in electrical load forecasting, on the other hand, is a significant undertaking that necessitates careful consideration of several parameters.

The usage of autoregressive models is a standard method for defining the input vector. These models anticipate future load levels by using historical load values as inputs. This method has the benefit of capturing the temporal dependence between load levels, which is necessary for good forecasting. It does, however, presume that load levels follow a linear trend, which is not necessarily the case.

Another method for capturing periodicity in load data is to employ Fourier transformations. This entails breaking down the load data into its constituent frequency components and feeding them into the MLP model. This method has the benefit of capturing seasonal trends and periodic swings in load data. However, it assumes that the load data has a stationary frequency domain, which is not always the case.

Wavelet transformations have also been utilized to capture load data's temporal and frequency dependencies. The load data is decomposed into wavelet coefficients, which are then fed into the MLP model. This method has the benefit of capturing both short-term and long-term dependence in load data. However, it necessitates a significant amount of computation, which may be prohibitively expensive for large datasets.

To minimize the dimensionality of the input vector, principal component analysis (PCA) was also performed. This entails reducing the original input variables to a smaller collection of uncorrelated variables that explain the majority of the data variance. This method has the advantage of reducing the computational complexity of the MLP model and potentially improving its interpretability. It does, however, imply that the input variables are linearly connected, which is not always the case.

To summarize, there are numerous techniques for creating the input vector for MLP models in energy load forecasting, each with its own set of benefits and drawbacks. The technique of choosing should be determined by the unique characteristics of the load data as well as the needs of the forecasting activity. The trade-off between model accuracy and computational complexity, as well as the interpretability of the final model, should be carefully considered. More research is needed to compare the performance of these various approaches on various datasets and under various conditions.

B)

Loading the dataset

```

R 4.3.0 · ~/
> #import libraries
> library(ggpubr)
> library(neuralnet)
> library(readxl)
> library(Metrics)
> Data_load <- read_excel("C:/Users/my pc/Desktop/ML cw/uow_consumption.xlsx")
> Data_load
# A tibble: 470 × 4
  date           `0.75` `0.7916666666666663` `0.8333333333333337`
  <dtm>          <dbl>          <dbl>          <dbl>
1 2018-01-01 00:00:00 38.9            38.9            38.9
2 2018-01-02 00:00:00 42.3            41.9            41.9
3 2018-01-03 00:00:00 40.8            40.5            40.7
4 2018-01-04 00:00:00 42.3            41.9            41.9
5 2018-01-05 00:00:00 44              44.1            44
6 2018-01-06 00:00:00 45.6            44.5            44.3
7 2018-01-07 00:00:00 40.9            41.2            35.7
8 2018-01-08 00:00:00 44.6            44.3            43.6
9 2018-01-09 00:00:00 41.9            42.9            42.5
10 2018-01-10 00:00:00 44.9            44.4            43.9
# i 460 more rows
# i Use `print(n = ...)` to see more rows
> |

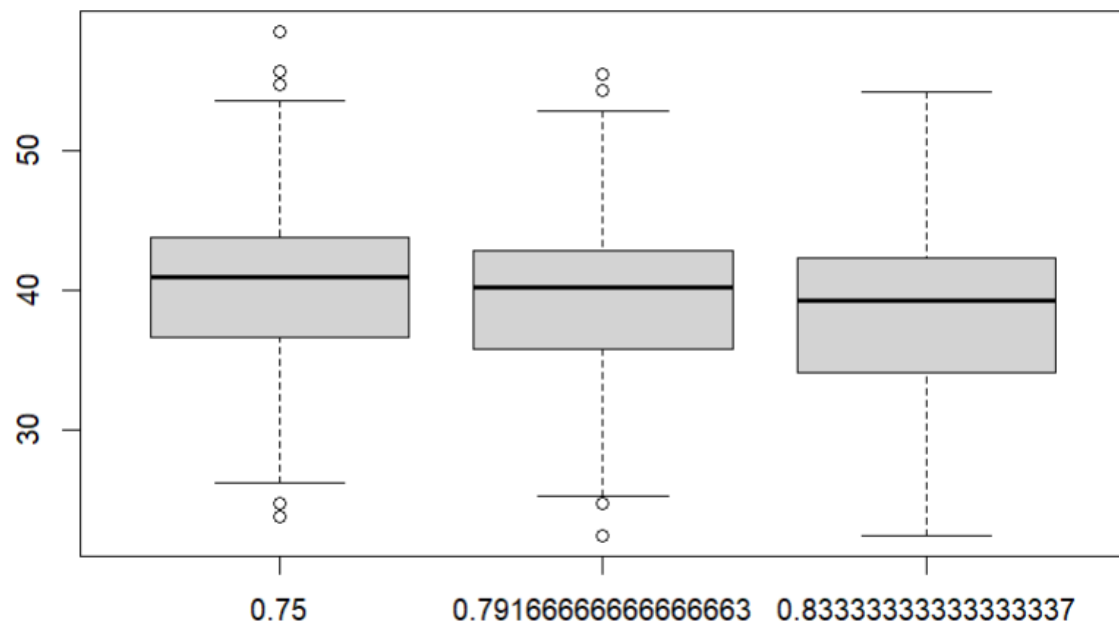
```

Summary of the dataset

```

> summary(Data_load)
      date           0.75      0.7916666666666663 0.8333333333333337
Min.   :2018-01-01 00:00:00 Min.   :23.80 Min.   :22.40 Min.   :22.40
1st Qu.:2018-04-28 06:00:00 1st Qu.:36.60 1st Qu.:35.83 1st Qu.:34.10
Median :2018-08-23 12:00:00 Median :40.90 Median :40.15 Median :39.20
Mean   :2018-08-23 12:00:00 Mean   :40.18 Mean   :39.27 Mean   :38.36
3rd Qu.:2018-12-18 18:00:00 3rd Qu.:43.80 3rd Qu.:42.77 3rd Qu.:42.30
Max.   :2019-04-15 00:00:00 Max.   :58.50 Max.   :55.40 Max.   :54.20
> boxplot(Data_load[, -1]) #plot before normalizing data
> |

```



c)

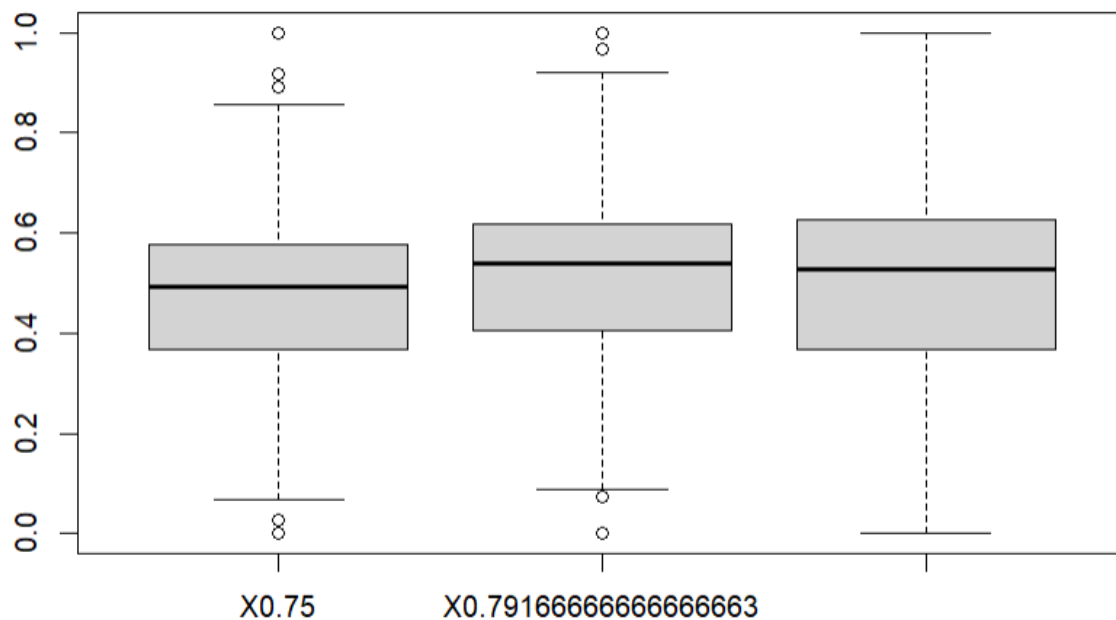
Normalizing dataset

To guarantee that all input variables are on a same scale, data must be normalized before being used in a Multi-Layer Perceptron (MLP) structure. This is significant because, during training, MLPs utilize mathematical functions to adjust the weights of connections between neurons, and these functions might be sensitive to the size of input data. When data is not normalized, variables with greater values might dominate the training process, resulting in poor performance or difficulty with convergence. By normalizing the data, we guarantee that each input variable contributes evenly to the training process, hence improving the model's overall performance.

```

> #normalize function
> normalize <- function(x){
+   return ((x - min(x)) / (max(x) - min(x)))
+ }
> uow_load_norm <- as.data.frame(lapply(Data_load[-1,-1], normalize))# normalized data without
> View(uow_load_norm)
> summary(uow_load_norm)
      X0.75      X0.7916666666666663 X0.8333333333333337
Min.   :0.0000   Min.   :0.0000      Min.   :0.0000
1st Qu.:0.3689   1st Qu.:0.4061      1st Qu.:0.3679
Median :0.4928   Median :0.5394      Median :0.5283
Mean   :0.4721   Mean   :0.5113      Mean   :0.5018
3rd Qu.:0.5764   3rd Qu.:0.6182      3rd Qu.:0.6258
Max.   :1.0000   Max.   :1.0000      Max.   :1.0000
> boxplot(uow_load_norm) #plot after normalization
> |

```



Split dataset

```

> #split data
> traindata <- uow_load_norm[1:400,] #data used for training model
> testdata <- uow_load_norm[400:469,] #data used for testing model
> View(traindata)
> names(traindata)
[1] "X0.75" "X0.7916666666666663" "X0.8333333333333337"
> testdata
      X0.75 X0.7916666666666663 X0.8333333333333337
400 0.4927954 0.5181818 0.4874214
401 0.3861671 0.4545455 0.4402516
402 0.3573487 0.4060606 0.4119497
403 0.3602305 0.3969697 0.4056604
404 0.3429395 0.4151515 0.4213836
405 0.2507205 0.2939394 0.2672956
406 0.4466859 0.4757576 0.4748428
407 0.4063401 0.4666667 0.4465409
408 0.4610951 0.4939394 0.5125786
409 0.3804035 0.4303030 0.4433962
410 0.4466859 0.5060606 0.5251572
411 0.3487032 0.4151515 0.4402516
412 0.3717579 0.4303030 0.3113208
413 0.3804035 0.4393939 0.4371069
414 0.4034582 0.4848485 0.4968553
415 0.3919308 0.4424242 0.4182390
416 0.4553314 0.5090909 0.5062893
417 0.4783862 0.5212121 0.5157233
418 0.5187320 0.5818182 0.5786164
419 0.4236311 0.4787879 0.4245283
420 0.5878963 0.6121212 0.6226415
421 0.6858790 0.7090909 0.6855346
422 0.8126801 0.7606061 0.7421384
423 0.4783862 0.5484848 0.5345912
424 0.5677233 0.5545455 0.5691824
425 0.4149856 0.4666667 0.4937107
426 0.4121037 0.4696970 0.3710692
427 0.5734870 0.5969697 0.5691824
428 0.7089337 0.7212121 0.7389937
429 0.6714697 0.7303030 0.7389937
430 0.6685879 0.7484848 0.7547170
431 0.6195965 0.6878788 0.7389937

```

430	0.6685879	0.7484848	0.7547170
431	0.6195965	0.6878788	0.7389937
432	0.6570605	0.7272727	0.6981132
433	0.5302594	0.5818182	0.4937107
434	0.5734870	0.6848485	0.6949686
435	0.5331412	0.6181818	0.6415094
436	0.6080692	0.6636364	0.6761006
437	0.5936599	0.6363636	0.5849057
438	0.5648415	0.6484848	0.6792453
439	0.5216138	0.5818182	0.6194969
440	0.5331412	0.6000000	0.4371069
441	0.6570605	0.6909091	0.7327044
442	0.6772334	0.7696970	0.7987421
443	0.6167147	0.6939394	0.7295597
444	0.6282421	0.6909091	0.7264151
445	0.5446686	0.6484848	0.6603774
446	0.6080692	0.6666667	0.7264151
447	0.5216138	0.5787879	0.4968553
448	0.6167147	0.6757576	0.6981132
449	0.5590778	0.5787879	0.6006289
450	0.6570605	0.7151515	0.7327044
451	0.6195965	0.6575758	0.6572327
452	0.8069164	0.7272727	0.7201258
453	0.5878963	0.6333333	0.6572327
454	0.5244957	0.6030303	0.5125786
455	0.4985591	0.5363636	0.5377358
456	0.2536023	0.2454545	0.2484277
457	0.5389049	0.6030303	0.5880503
458	0.5763689	0.6363636	0.6572327
459	0.5216138	0.5727273	0.6069182
460	0.5417867	0.6121212	0.6383648
461	0.4755043	0.4151515	0.4213836
462	0.4524496	0.5181818	0.5188679
463	0.2968300	0.3363636	0.3270440
464	0.5216138	0.5484848	0.5440252
465	0.5302594	0.5818182	0.5880503
466	0.5533141	0.6030303	0.6320755
467	0.4524496	0.5303030	0.5314465
468	0.4726225	0.4757576	0.4716981
469	0.5331412	0.5969697	0.6037736

> |

E)

Indices explanation

RMSE

RMSE is an abbreviation for Root Mean Squared Error, which is a popular statistic for determining the accuracy of a prediction model. It computes the square root of the average squared difference between expected and actual values to calculate the difference between predicted and actual values.

The root mean square error (RMSE) is often used in regression analysis to calculate the difference between the anticipated and actual values of the dependent variable. It is chosen over Mean Absolute Error (MAE) because squaring the difference penalizes big mistakes more harshly than tiny errors. A lower RMSE number shows that the model is better fitted to the data, whereas a higher RMSE value indicates that the model has a bigger prediction error. RMSE, on the other hand, should always be viewed in the context of the issue domain and the range of values being forecasted.

MAE

MAE is an abbreviation for Mean Absolute Error, which is a popular statistic for determining the accuracy of a prediction model. It computes the average absolute difference between anticipated and actual values. MAE is frequently used in regression analysis to calculate the difference between the expected and actual values of the dependent variable.

In cases when big mistakes are not necessarily more relevant than tiny errors, it is chosen over Root Mean Squared Error (RMSE). A lower MAE number implies that the model fits the data better, whereas a higher MAE value shows that the model has a bigger prediction error. However, MAE should always be considered in the context of the problem domain and the range of predicted values.

MAPE

MAPE is an abbreviation for Mean Absolute Percentage Error, which is a popular statistic for determining the accuracy of a prediction model. It computes the percentage difference between expected and actual values.

MAPE is a regularly used metric in forecasting and time series analysis to assess the accuracy of predictions made across several time periods. It is an effective indicator for comparing the performance of various forecasting models. A lower MAPE number implies that the model fits the data better, whereas a higher MAPE value shows that the model has a bigger prediction error. MAPE does have certain limitations, such as being sensitive to extreme values or tiny data sets, and it may not be appropriate for instances when the real values are near to zero.

Overall, MAPE should be considered alongside other evaluation metrics and interpreted within the context of the specific problem domain and the range of values being predicted.

symmetric MAPE

The measure Symmetric Mean Absolute Percentage Error (sMAPE) is a variation on the Mean Absolute Percentage Error (MAPE) that tackles some of its shortcomings. sMAPE calculates the percentage difference between expected and real values, but unlike MAPE, it uses the absolute difference between actual and predicted values as the denominator rather than the average of the actual values.

The sMAPE formula is as follows:

$$\text{sMAPE} = (1/n) * (((|A_t| + |F_t|)/2)) * 100\%$$

Where A_t represents the actual value, F_t represents the forecast value, and n is the number of observations.

sMAPE is a number that spans from 0% to 200%, with lower values suggesting higher model accuracy. A number of 0% represents the best possible forecast, while a value of 200% represents the worst possible prediction. sMAPE is widely used in time series forecasting, and it is especially beneficial when the actual and forecasted values have comparable scales. It is particularly important when dealing with intermittent demand since it allows for a more realistic comparison of various forecasting models.

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