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1 C:\Users\Sapan\Downloads\4p76Assign1\assign1\venv\Scripts\python.exe C:/Users/Sapan/
  Downloads/4p76Assign1/assign1/main.py
2 Enter 1 for Digits classification, Enter 2 for Cancer classification: 2
3 Enter number of tests to compare (only works on 3), if don't want to run tests just
  single run enter 1: 3
4 Give name to your Test for files: cancer_momentum
5
6 Test no.: 0
7 Enter number of neurons for hidden layer: 3
8 Enter number of epochs: 8
9 Enter 1 for sigmoid or 2 for tanh activation function: 1
10 Enter learning rate: 0.5
11 Enter Momentum: 0.15
12 Running-----
13 Fold no.: 0
14 Accumulated sum error over epochs: [80.97455485285855, 38.455472904258905, 15.
  01162027698563, 10.143407275794551, 8.575864325917275, 7.849296238622715, 7.
  433883744617162, 7.160871903166746]
15
16 Correct classification validation set: 123
17
18 Correct classification test set: 200
19 end of fold
20
21 Running-----
22 Fold no.: 1
23 Accumulated sum error over epochs: [75.25165328125382, 31.618668439448058, 14.
  068342988409727, 10.386338973861454, 9.076958340580896, 8.436470124644032, 8.
  061934715520552, 7.815529919571838]
24
25 Correct classification validation set: 123
26
27 Correct classification test set: 200
28 end of fold
29
30 Running-----
31 Fold no.: 2
32 Accumulated sum error over epochs: [72.37047759517374, 28.812977778046342, 12.
  717966671932446, 9.157992127845096, 7.832599403943147, 7.178706120387782, 6.
  80196479545976, 6.561491055933503]
33
34 Correct classification validation set: 123
35
36 Correct classification test set: 200
37 end of fold
38
39 Running-----
40 Fold no.: 3
41 Accumulated sum error over epochs: [73.69933297541613, 24.252115310929852, 10.
  938703993417564, 8.214601988269425, 7.158607014493312, 6.602313705523263, 6.
  25388092728742, 6.009248341966572]
42
43 Correct classification validation set: 123
44
45 Correct classification test set: 200
46 end of fold
47
48 Fold test accuracies%: [100.0, 100.0, 100.0, 100.0]
49 Fold validation accuracies%: [100.0, 100.0, 100.0, 100.0]
50 Mean fold validation accuracies%: 100.0
51
52 Test no.: 1
53 Enter number of neurons for hidden layer: 3
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```
54 Enter number of epochs: 8
55 Enter 1 for sigmoid or 2 for tanh activation function: 1
56 Enter learning rate: 0.5
57 Enter Momentum: 0.5
58 Running-----
59 Fold no.: 0
60 Accumulated sum error over epochs: [83.29601145122218, 49.80375582537235, 17.
    89419677724818, 11.724022125627114, 10.103484679640117, 9.425726700111468, 9.
    07095553097434, 8.859372997869253]
61
62 Correct classification validation set: 123
63
64 Correct classification test set: 200
65 end of fold
66
67 Running-----
68 Fold no.: 1
69 Accumulated sum error over epochs: [77.26616568239608, 28.13108761964464, 11.
    111122895811757, 7.788699665828535, 6.5650146108755, 5.939148569334366, 5.
    556002527758465, 5.291148725314111]
70
71 Correct classification validation set: 123
72
73 Correct classification test set: 200
74 end of fold
75
76 Running-----
77 Fold no.: 2
78 Accumulated sum error over epochs: [75.61648306821414, 35.03838659576255, 14.
    310663758507372, 10.730540928109953, 9.616654763680417, 9.075421382782007, 8.
    736339315454806, 8.489986646871268]
79
80 Correct classification validation set: 123
81
82 Correct classification test set: 200
83 end of fold
84
85 Running-----
86 Fold no.: 3
87 Accumulated sum error over epochs: [78.86379401650046, 32.761895013425814, 14.
    451747599483001, 11.247793076470115, 10.185726547131411, 9.665888895989736, 9.
    348268167142358, 9.124307780161491]
88
89 Correct classification validation set: 123
90
91 Correct classification test set: 200
92 end of fold
93
94 Fold test accuracies%: [100.0, 100.0, 100.0, 100.0]
95 Fold validation accuracies%: [100.0, 100.0, 100.0, 100.0]
96 Mean fold validation accuracies%: 100.0
97
98 Test no.: 2
99 Enter number of neurons for hidden layer: 3
100 Enter number of epochs: 8
101 Enter 1 for sigmoid or 2 for tanh activation function: 1
102 Enter learning rate: 0.5
103 Enter Momentum: 0.85
104 Running-----
105 Fold no.: 0
106 Accumulated sum error over epochs: [83.03496877134717, 53.66671221085658, 18.
    051366430453676, 10.455913671931722, 8.571736737451669, 7.805080147601388, 7.
    401445090702632, 7.153980171626269]
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107
108 Correct classification validation set: 123
109
110 Correct classification test set: 200
111 end of fold
112
113 Running-----
114 Fold no.: 1
115 Accumulated sum error over epochs: [83.6807234316516, 59.91799223821635, 20.
    52724891932206, 10.902594659151891, 8.798678413600294, 7.9889153090419525, 7.
    560985976587155, 7.28828890698281]
116
117 Correct classification validation set: 123
118
119 Correct classification test set: 200
120 end of fold
121
122 Running-----
123 Fold no.: 2
124 Accumulated sum error over epochs: [68.60003649470478, 21.8132726527921, 12.
    471786513412033, 10.594669136886852, 9.85895765732044, 9.442893398294428, 9.
    15010002762886, 8.917526857547546]
125
126 Correct classification validation set: 123
127
128 Correct classification test set: 200
129 end of fold
130
131 Running-----
132 Fold no.: 3
133 Accumulated sum error over epochs: [71.46772009445702, 22.12349928523018, 11.
    108382261031519, 9.274345648607383, 8.666168785385686, 8.38598622142792, 8.
    22940584210338, 8.128672086872955]
134
135 Correct classification validation set: 123
136
137 Correct classification test set: 200
138 end of fold
139
140 Fold test accuracies%: [100.0, 100.0, 100.0, 100.0]
141 Fold validation accuracies%: [100.0, 100.0, 100.0, 100.0]
142 Mean fold validation accuracies%: 100.0
143
144 Test results:
145     test0  test1  test2
146 0  100.0  100.0  100.0
147 1  100.0  100.0  100.0
148 2  100.0  100.0  100.0
149 3  100.0  100.0  100.0
150 Test stats:
151     test0  test1  test2
152 count    4.0    4.0    4.0
153 mean   100.0  100.0  100.0
154 std     0.0    0.0    0.0
155 min   100.0  100.0  100.0
156 25%   100.0  100.0  100.0
157 50%   100.0  100.0  100.0
158 75%   100.0  100.0  100.0
159 max   100.0  100.0  100.0
160
161 Validation results:
162     test0  test1  test2
163 0  100.0  100.0  100.0
```

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164 1 100.0 100.0 100.0
165 2 100.0 100.0 100.0
166 3 100.0 100.0 100.0
167 Valid stats:
168      test0  test1  test2
169 count    4.0    4.0    4.0
170 mean   100.0  100.0  100.0
171 std     0.0    0.0    0.0
172 min   100.0  100.0  100.0
173 25%   100.0  100.0  100.0
174 50%   100.0  100.0  100.0
175 75%   100.0  100.0  100.0
176 max   100.0  100.0  100.0
177 C:\Users\Sapan\Downloads\4p76Assign1\assign1\venv\lib\site-packages\scipy\stats\
    morestats.py:1678: UserWarning: Input data for shapiro has range zero. The results may
    not be accurate.
178     warnings.warn("Input data for shapiro has range zero. The results "
179
180 shapiro test for testing accuracies
181 test0 stats=1.0000, p=1.0000
182 test1 stats=1.0000, p=1.0000
183 test2 stats=1.0000, p=1.0000
184
185 shapiro test for validation accuracies
186 test0 stats=1.0000, p=1.0000
187 test1 stats=1.0000, p=1.0000
188 test2 stats=1.0000, p=1.0000
189
190 ANOVA test
191
192      sum_sq    df      F      PR(>F)
192 df_test.iloc[:, 1]  4.000000e+04  1.0  3.713820e+31  9.744049e-48
193 df_test.iloc[:, 2]  4.000000e+04  1.0  3.713820e+31  9.744049e-48
194 Residual          3.231174e-27  3.0           NaN           NaN
195
196 t-tests:
197 Ttest_indResult(statistic=nan, pvalue=nan)
198 Ttest_indResult(statistic=nan, pvalue=nan)
199 Ttest_indResult(statistic=nan, pvalue=nan)
200
201 No statistically better mean
202
203 Process finished with exit code 0
204
```