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1 C:\Users\jiepeng\anaconda3\python.exe "
  C:/Users/jiepeng/OneDrive - Yaojushi/
  CAU-No.455-Lab/case_study.py"
2 #####
3 =====
4 #####sos
5 GORK#####Top20#####
6 protein          score
7 AKT1             9.2040
8 HKT1             8.7653
9 TPC1             7.7803
10 KT2/3           7.6536
11 HAK5            7.2092
12 SOS3            6.2685
13 CBL1            6.1215
14 NHX1            6.0660
15 Name: GORK, dtype: float32
16 #####
17 =====
18 #####col
19 GORK#####Top20#####
20 protein          score
21 AKT1             9.0849
22 HKT1             8.3984
23 TPC1             7.5799
24 KT2/3           7.3682
25 HAK5            7.2506
26 SOS3            6.0570
27 CBL1            5.9816
28 NHX1            5.9456
29 Name: GORK, dtype: float32

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30 GORKEcol_sos=====32,
   result
31
32 =====
33
34 sos
35 CNGC10Top20
36 protein          score
37 HKT1             7.0292
38 TPC1             6.9503
39 AKT1             6.6732
40 KT2/3           5.7522
41 HAK5             5.6962
42 NHX1            4.9432
43 NHX2            4.7601
44 KAT3            4.5895
45 Name: CNGC10, dtype: float32
46 =====
47
48 col
49 CNGC10Top20
50 protein          score
51 TPC1             7.0753
52 HKT1             6.8784
53 AKT1             6.8378
54 HAK5             5.9408
55 KT2/3           5.6645
56 NHX1            4.9916
57 NHX2            4.8038
58 KAT3            4.5126
59 Name: CNGC10, dtype: float32

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60 CNGC10colsos
   result
61
62
63 =====
64 sos
65 IMPA-2Top20
66 protein      score
67 AT3G08943    3.7334
68 AT5G53480    3.7077
69 AT2G46520    3.5515
70 AT3G08947    3.3563
71 ATARCA       3.3182
72 ABH1          3.2917
73 AT4G27640    2.8851
74 NRPA2         2.8428
75 Name: IMPA-2, dtype: float32
76
77 =====
78 col
79 IMPA-2Top20
80 protein      score
81 AT3G08943    3.9597
82 AT5G53480    3.8942
83 AT2G46520    3.7686
84 AT3G08947    3.5878
85 ABH1          3.3543
86 NOP56         3.2228
87 ATARCA       3.1932
88 AT4G27640    3.0813
89 Name: IMPA-2, dtype: float32

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90 IMPA-2colsos
91 result
92
93 =====
94 sos
95 GLR2.7Top20
96 protein      score
97 TPC1         8.0439
98 GLUR2        4.6981
99 GLR1.4        4.6343
100 GLR3.4       4.4036
101 GLR1.1       4.1878
102 GLR1.3       4.1353
103 GLR2.1       4.1193
104 AKT1         3.9402
105 Name: GLR2.7, dtype: float32
106
107 =====
108 col
109 GLR2.7Top20
110 protein      score
111 TPC1         8.1149
112 GLUR2        4.7950
113 GLR1.4        4.6639
114 GLR3.4       4.5912
115 GLR1.1       4.2980
116 GLR1.3       4.2623
117 GLR2.3       4.1437
118 GLR2.1       4.0590
119 Name: GLR2.7, dtype: float32

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120 GLR2.7colsos=====28,
    result
121
122 =====
123
124 sos
125 CIB5Top20
126 protein          score
127 CRY1          3.4410
128 HY5           3.3518
129 LKP2          3.3274
130 CRY2          3.3143
131 HFR1          3.2853
132 ZTL           3.2339
133 SPA1          3.2183
134 CIB5          3.1019
135 Name: CIB5, dtype: float32
136 =====
137
138 col
139 CIB5Top20
140 protein          score
141 LKP2          3.2349
142 CRY1          3.2289
143 CRY2          3.2103
144 HY5           3.1986
145 SPA1          3.0902
146 HFR1          3.0746
147 ZTL           2.9987
148 PIF4          2.8844
149 Name: CIB5, dtype: float32

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150 CIB5colsos
151 result
152
153 =====
154 sos
155 ZIP7Top20
156 protein          score
157 IAR1              5.3129
158 ZTP29             4.8613
159 IRT1              4.7432
160 AT3G08650        4.4206
161 NRAMP4            3.9439
162 FR02             3.8514
163 ZAT               3.7658
164 VIT1             3.7347
165 Name: ZIP7, dtype: float32
166
167 =====
168 col
169 ZIP7Top20
170 protein          score
171 IAR1              5.2238
172 ZTP29             4.7942
173 IRT1              4.6931
174 AT3G08650        4.3546
175 NRAMP4            3.9003
176 FR02             3.7860
177 ZAT               3.7285
178 VIT1             3.6364
179 Name: ZIP7, dtype: float32

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180 ZIP7colsos00000000000000000018,  
    0000result000
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181
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182
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183 Process finished with exit code 0
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184
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