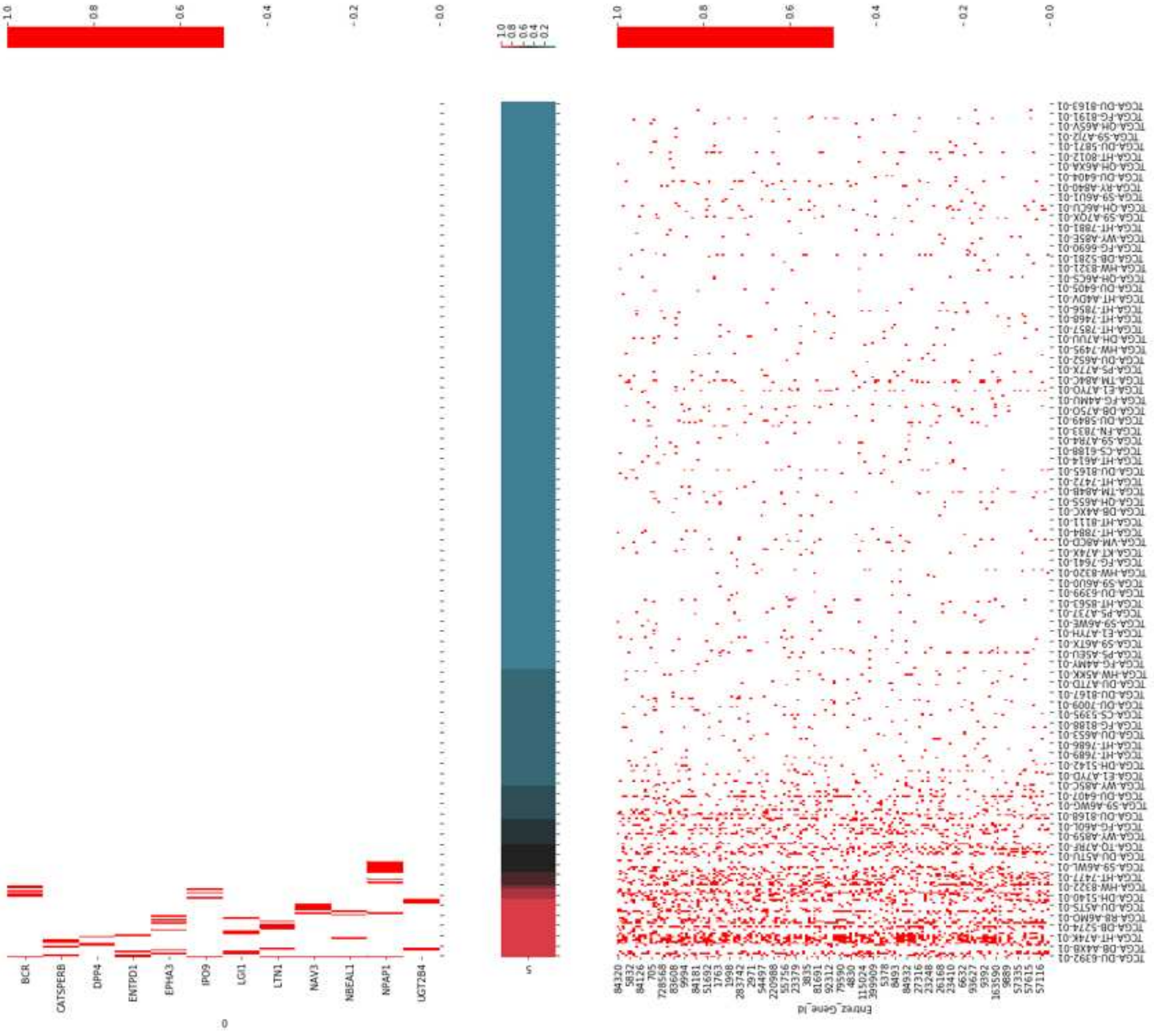
1.0
0.8
0.6
0.4
0.2
0.0
n
BCR
CATSPERB
DPP4
ENTPD1
EPHA3
IPO9
LGI1
LTN1
NAV3
NBEAL1
NPAP1
UGT2B4
0
Entrez_Gene_Id
84320
5832
84126
705
728568
83608
9994
84181
51692
1763
1998
283742
2971
54497
220988
55756
23379
3835
81691
92312
79590
4830
115024
399909
5378
8493
84932
27316
23248
26168
23410
6632
93627
9392
163590
9889
57335
57615
57116
TCGA-DU-8163-01
TCGA-FG-8191-01
TCGA-OH-A65V-01
TCGA-59-A7J2-01
TCGA-DU-5871-01
TCGA-HT-8012-01
TCGA-OH-A6XA-01
TCGA-DU-6404-01
TCGA-RY-A840-01
TCGA-S9-A6U1-01
TCGA-OH-A6CU-01
TCGA-S9-A7QX-01
TCGA-HT-7881-01
TCGA-WY-A85E-01
TCGA-FG-6690-01
TCGA-DB-5281-01
TCGA-HW-8321-01
TCGA-OH-A6C5-01
TCGA-DU-6405-01
TCGA-HT-A4DV-01
TCGA-HT-7856-01
TCGA-HT-7468-01
TCGA-HT-7857-01
TCGA-DH-A7UU-01
TCGA-HW-7495-01
TCGA-DU-A652-01
TCGA-P5-A77X-01
TCGA-E1-A7YO-01
TCGA-TM-A84C-01
TCGA-FG-A4MU-01
TCGA-DB-A7S0-01
TCGA-DU-5849-01
TCGA-FN-7833-01
TCGA-59-A7R4-01
TCGA-CS-6188-01
TCGA-HT-A614-01
TCGA-DU-8165-01
TCGA-HT-7472-01
TCGA-TM-A84B-01
TCGA-OH-A655-01
TCGA-DB-A4XC-01
TCGA-HT-8111-01
TCGA-HT-7884-01
TCGA-VM-A8CD-01
TCGA-KT-A74X-01
TCGA-FG-7641-01
TCGA-HW-8320-01
TCGA-S9-A6U0-01
TCGA-DU-6399-01
TCGA-HT-8563-01
TCGA-P5-A737-01
TCGA-S9-A6WE-01
TCGA-E1-A7YH-01
TCGA-S9-A6TX-01
TCGA-P5-A5EU-01
TCGA-FG-A4MY-01
TCGA-HW-A5KK-01
TCGA-DU-A7TD-01
TCGA-DU-8167-01
TCGA-DU-7009-01
TCGA-CS-5395-01
TCGA-FG-8188-01
TCGA-DU-A653-01
TCGA-HT-7686-01
TCGA-HT-7689-01
TCGA-DH-5142-01
TCGA-E1-A7YD-01
TCGA-WY-A85C-01
TCGA-DU-6407-01
TCGA-S9-A6WG-01
TCGA-DU-8168-01
TCGA-FG-A60L-01
TCGA-WY-A859-01
TCGA-TQ-A7RF-01
TCGA-DU-A5TU-01
TCGA-S9-A6WL-01
TCGA-HT-7477-01
TCGA-HW-8322-01
TCGA-DH-5140-01
TCGA-DU-A5TS-01
TCGA-RB-A6MO-01
TCGA-DB-5274-01
TCGA-HT-A74X-01
TCGA-DB-A4XB-01
TCGA-DU-6392-01