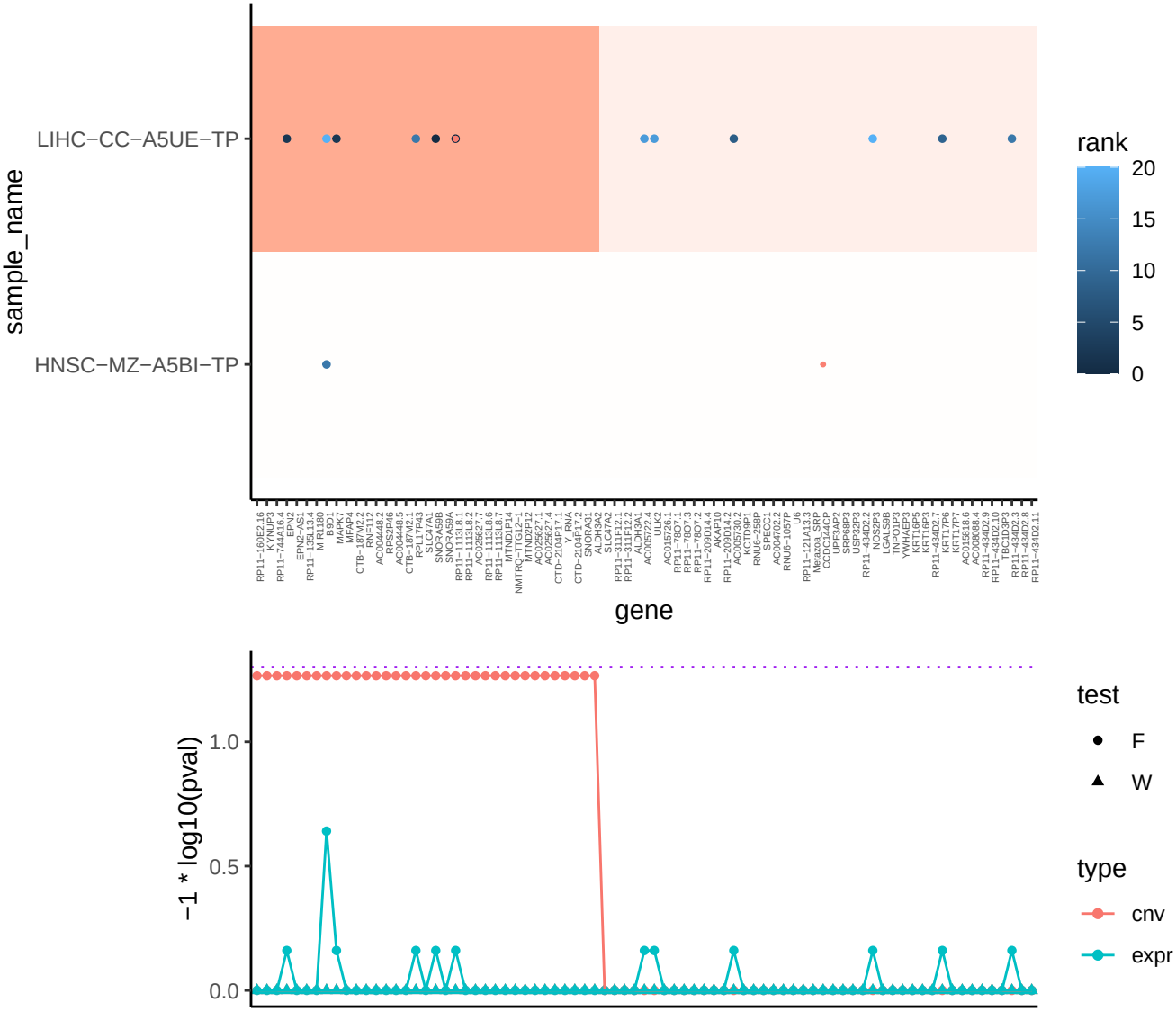
sample_name
LIHC−CC−A5UE−TP
HNSC−MZ−A5BI−TP
rank
20
15
10
5
0
RP11−160E2.16
KYNU P3
RP11−74A4.5
EBF4
EPN2
EPN2−AS1
RP11−135L13.4
MIR1180
ZBP1
MAPK7
MFAP4
CTB−187M2.2
RNF112
AC004448.2
RPS2P46
AC004448.5
CTB−187M2.1
RPL17P43
SLC7A1
SNORA59B
SNORA59A
RP11−1113L8.1
RP11−1113L8.7
AC025627
RP11−1113L8.6
RP11−1113L8.7
MTND1P14
NMTRK2 TTC22.1
MTND2P12
AC025627.1
AC025627.4
CTD−2104P17.1
CTD−2104P17.2
SNORA31
ALDH3A2
SLC7A2
RP11−311F12.1
RP11−311F12.2
ALDH3A1
AC005722.4
LUM
AC015726.1
RP11−78O7.1
RP11−78O7.3
RP11−78O7.2
RP11−209D14.1
RP11−209D14.2
AKAP10
RP11−209D14.2
AC005730.2
KIF19P1
RNU6−1299P
SPECC1
AC004702.2
RNU6−1057P
U6
RP11−121A13.3
Metazoa_SRP
CCDC144CP
UPF3AP2
SSR3P2
USP32P3
RP11−434D2.2
NOS2P3
LGALS9B
TNNC2P3
YWHAEP3
KRT16P5
KRT16P3
RP11−434D2.7
KRT17P6
KRT17P7
AC015918.6
AC008088.4
RP11−434D2.9
RP11−434D2.10
TBC1D3P3
RP11−434D2.3
RP11−434D2.8
RP11−434D2.11
gene
−1 * log10(pval)
1.0
0.5
0.0
test
F
W
type
cnv
expr