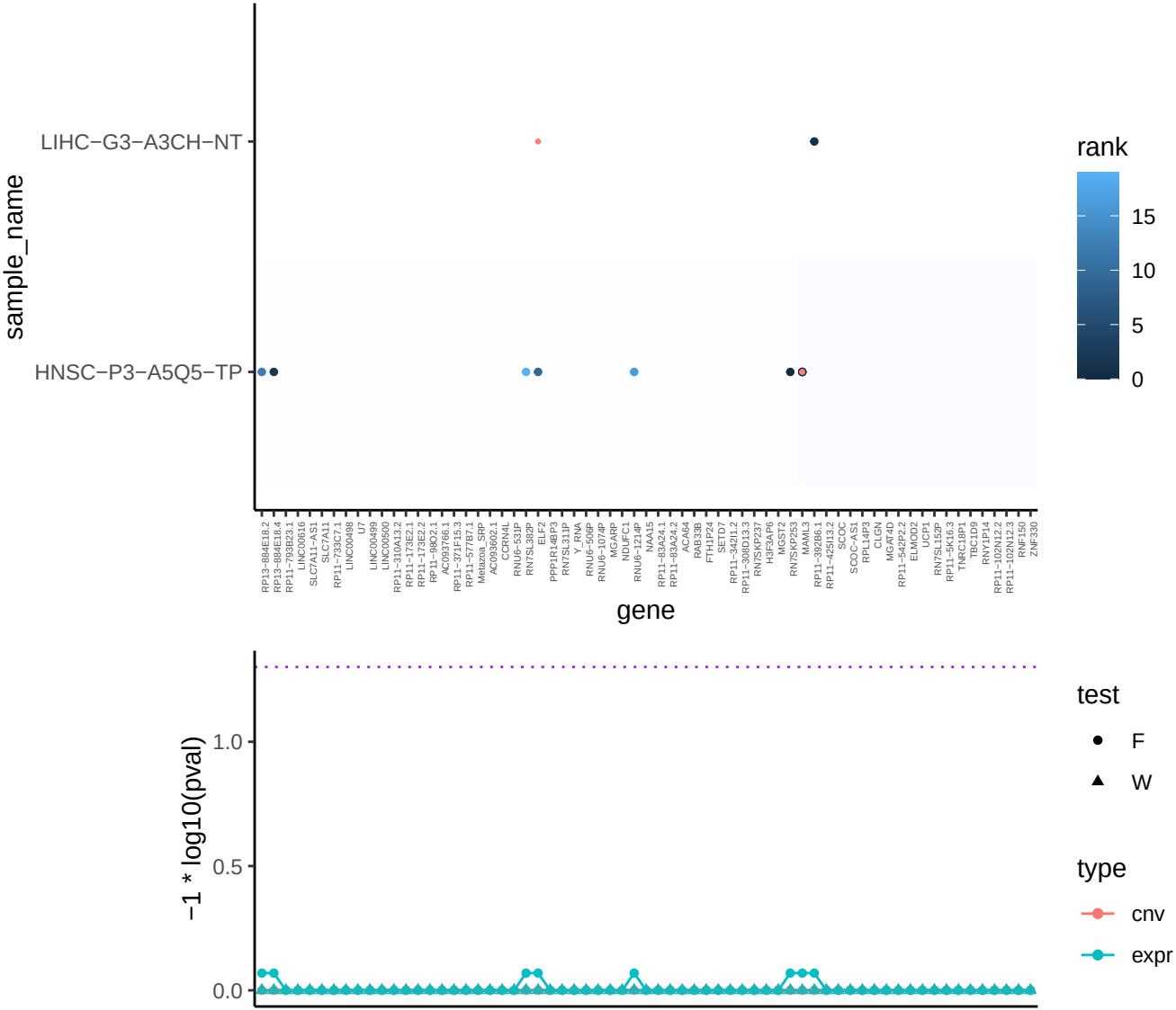
sample_name
LIHC−G3−A3CH−NT
HNSC−P3−A5Q5−TP
rank
15
10
5
0
RP13−884E18.2
RP13−884E18.4
RP11−793B23.1
LINC00616
SLC7A11−AS1
SLC7A11
RP11−733C7.1
LINC00498
U7
LINC00499
LINC00500
RP11−310A13.2
RP11−173E2.1
RP11−173E2.2
RP11−98O2.1
AC093766.1
RP11−371F15.3
RP11−577B7.1
Metazoa_SRP
ACOX3P1
CCRN4L
RNU6−531P
RN7SL382P
ELF2
PPP1R14BP3
RN7SL311P
Y_RNA
RNU6−506P
RNU6−1074P
MGAFP
NDUFC1
RNU6−1214P
NAA15
RP11−83A24.1
RP11−83A24.2
ACA64
RAB33B
FTH1P24
SETD7
RP11−34O11.2
RP11−309O13.3
RN7SKP237
H3F3AP6
MGST2
RN7SKP253
MAML3
RP11−392B6.1
RP11−425I13.2
SCOC
SCOC−AS1
RPL14P3
CLGN
MGAT4D
RP11−542P2.2
ELMOD2
UCP1
RN7SL152P
RP11−5K16.3
TNRC18P1
TBC1D9
RNY1P14
RP11−102O12.2
RP11−102N12.3
RNF150
ZNF330
gene
−1 * log10(pval)
1.0
0.5
0.0
test
F
W
type
cnv
expr