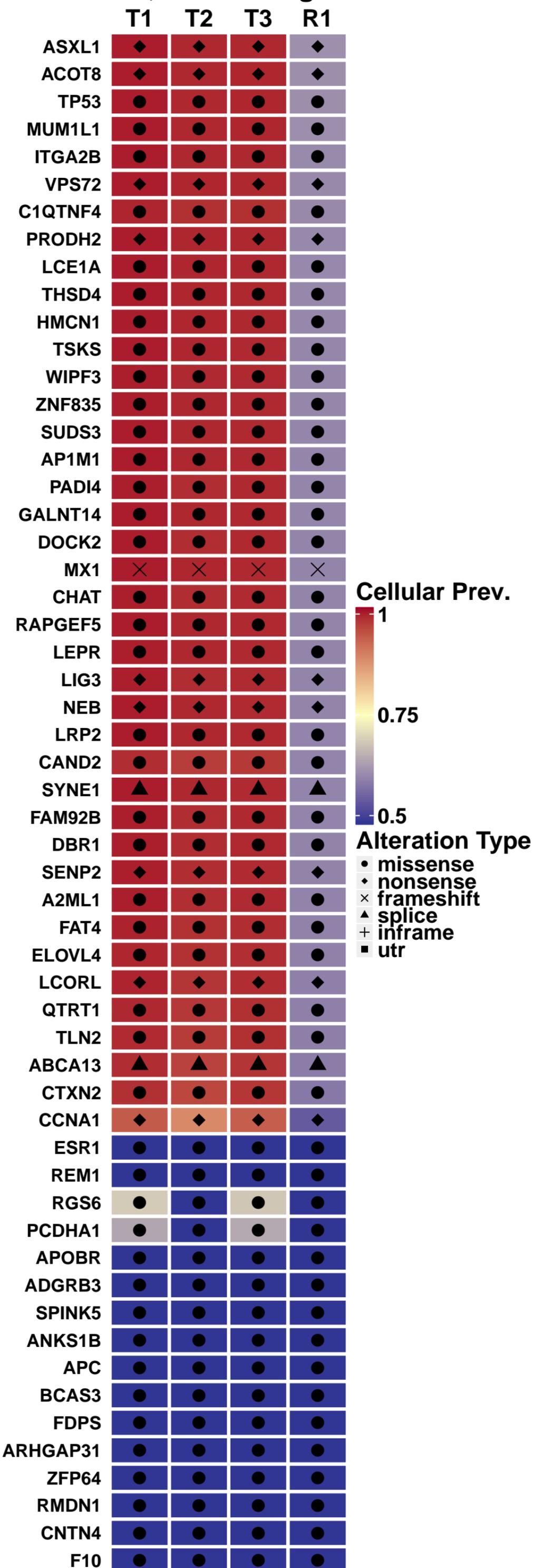
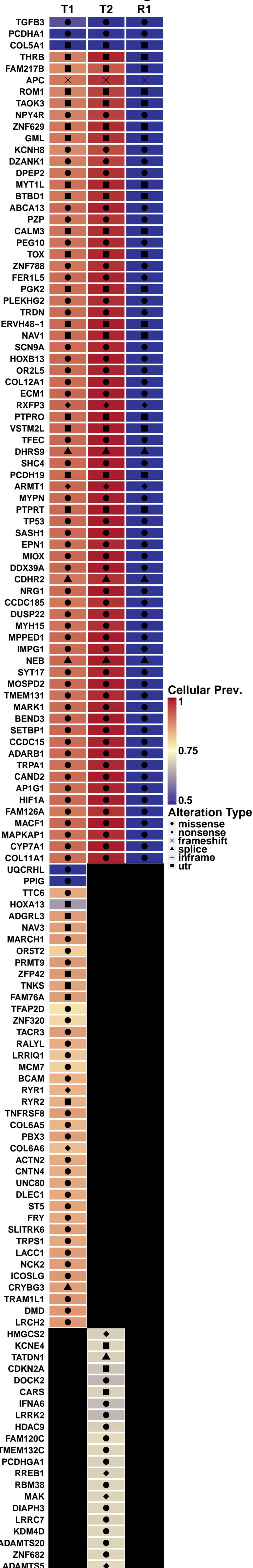


NP01, 56 variant genes



NP04, 134 variant genes

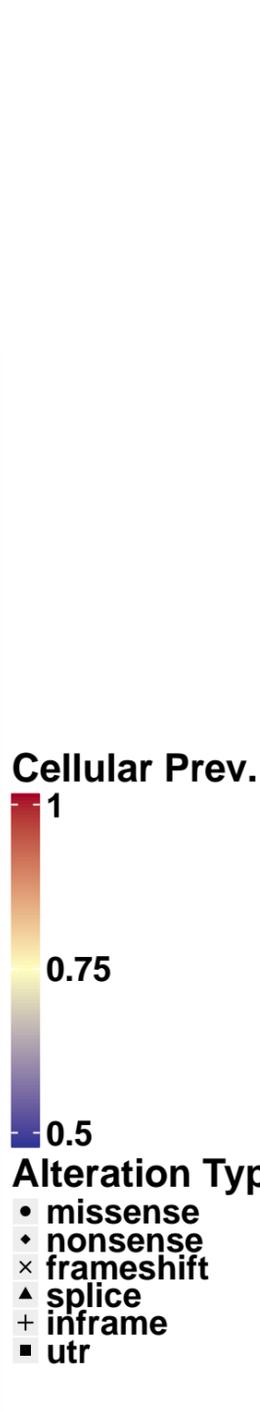
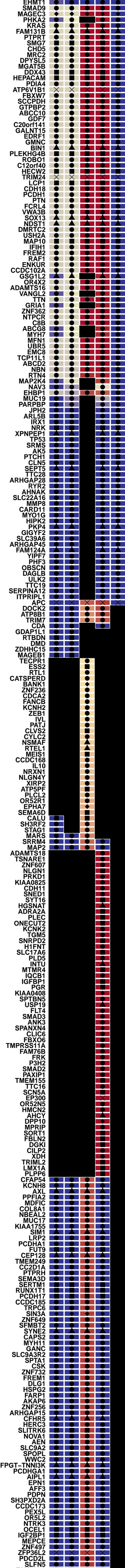


Cellular Prev.
1
0.75
0.5

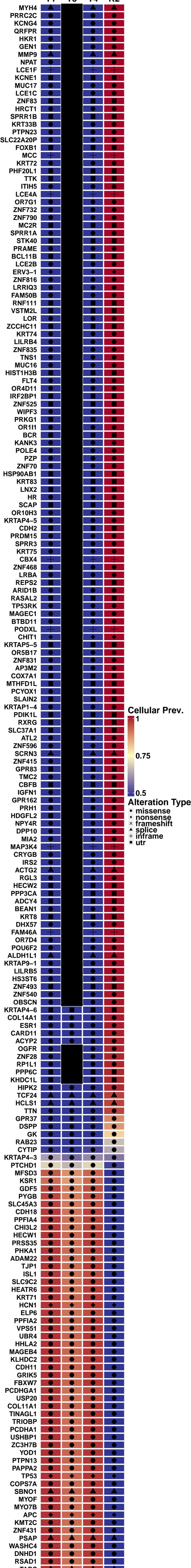
Alteration Type
● missense
◆ nonsense
× frameshift
▲ splice
+ inframe
■ utr

NP10, 272 variant genes

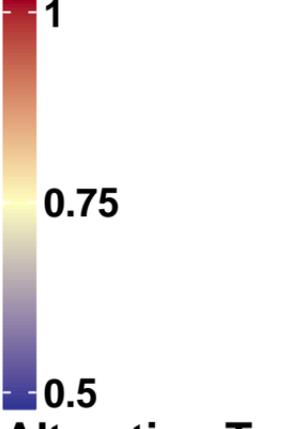
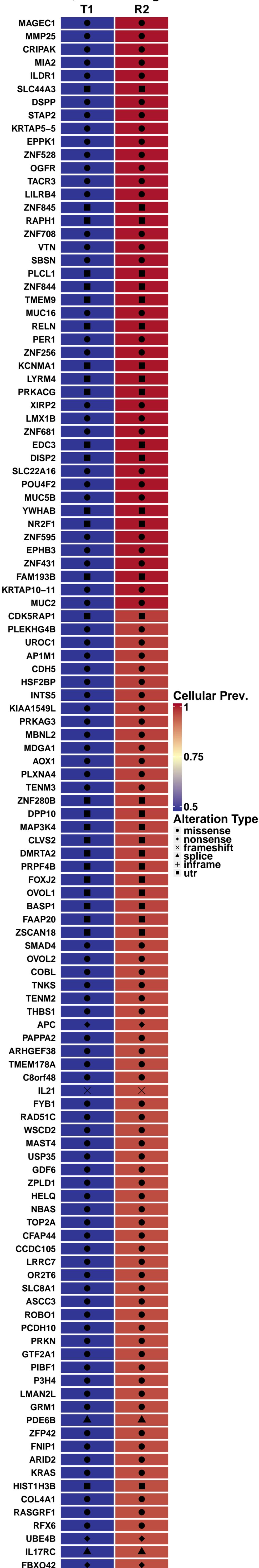
T1 T2 T3 T4 R1



NP11, 202 variant genes



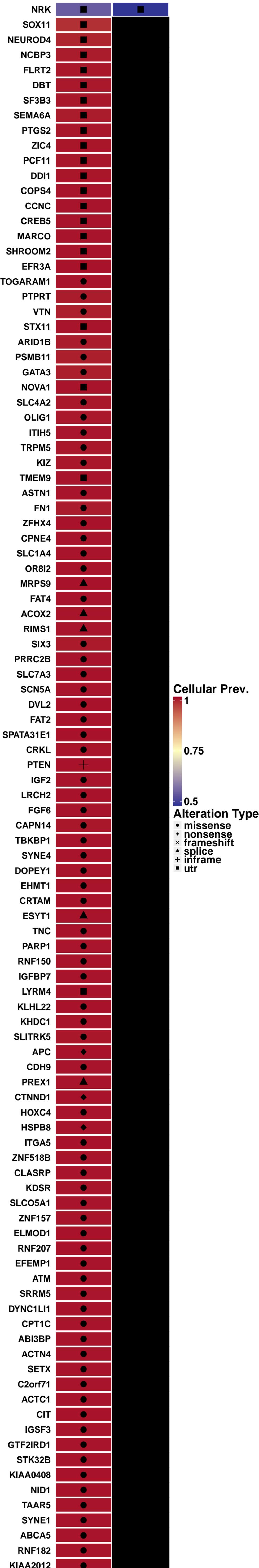
NP12, 119 variant genes



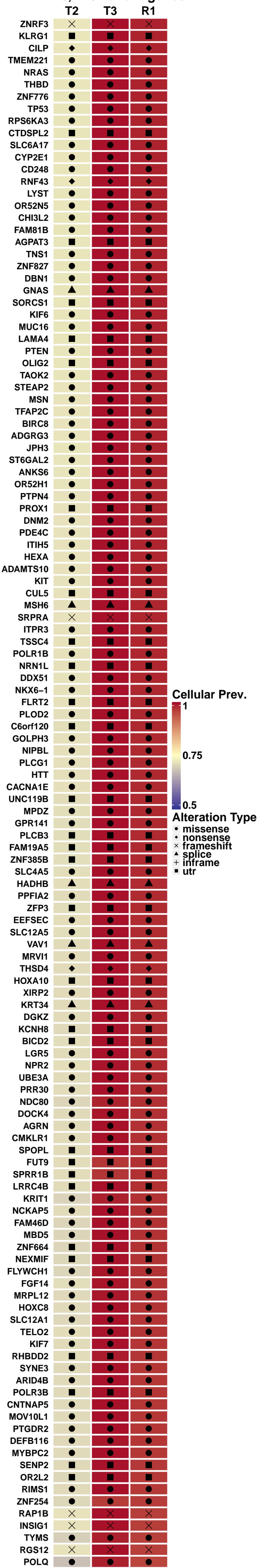
- Alteration Type**
- missense
 - ◆ nonsense
 - × frameshift
 - ▲ splice
 - + inframe
 - utr

NP14, 105 variant genes

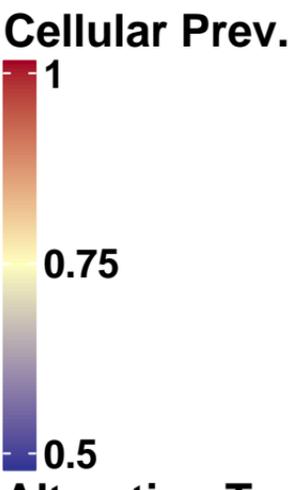
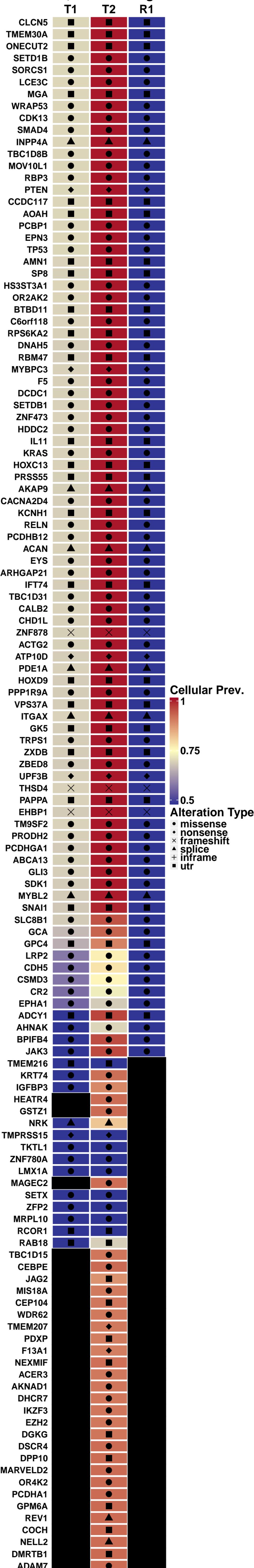
T1 R2



NP16, 129 variant genes

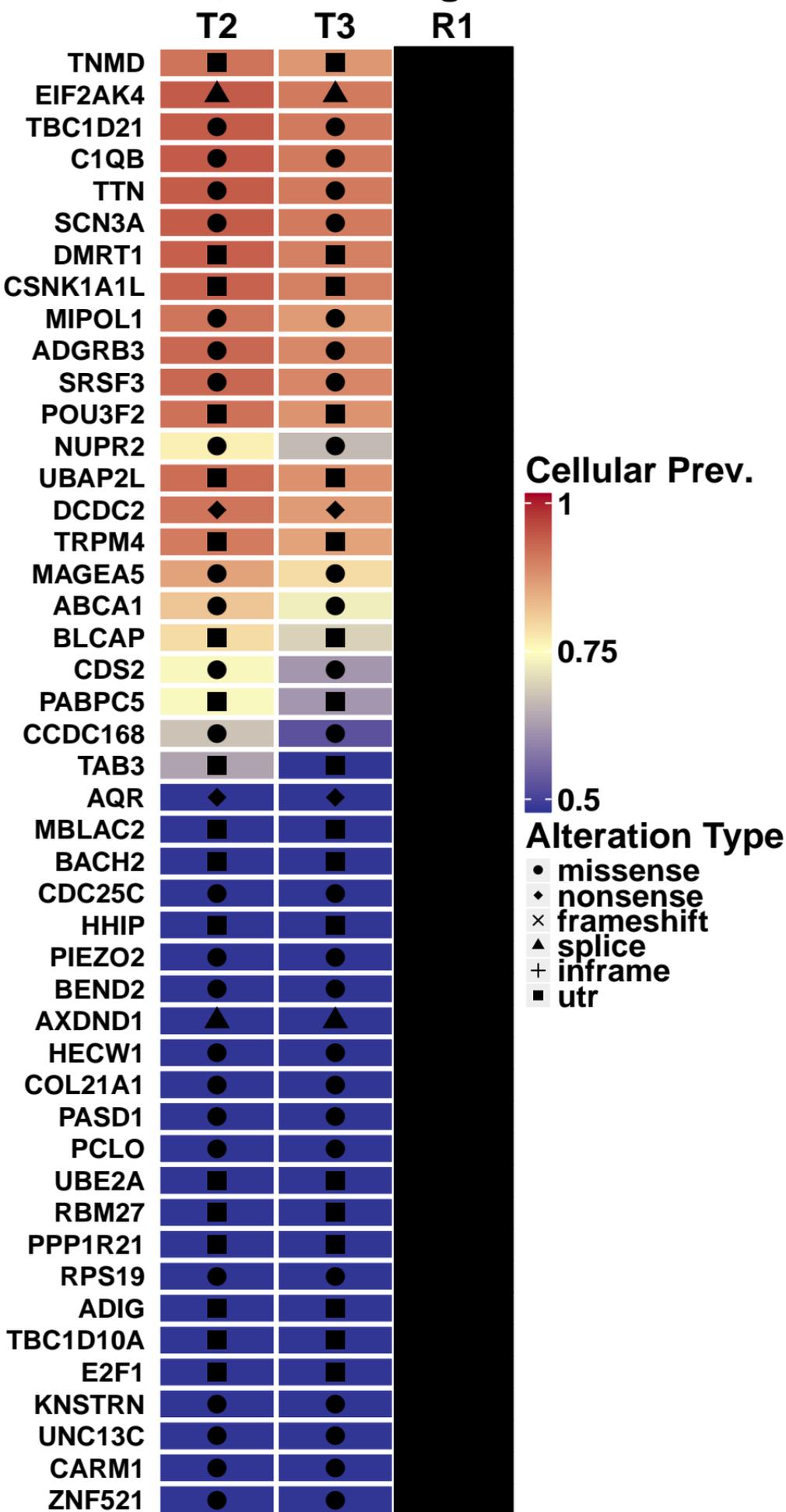


NP17, 130 variant genes



- Alteration Type**
- missense
 - ◆ nonsense
 - × frameshift
 - ▲ splice
 - + inframe
 - utr

NP18, 46 variant genes



NP19, 95 variant genes

