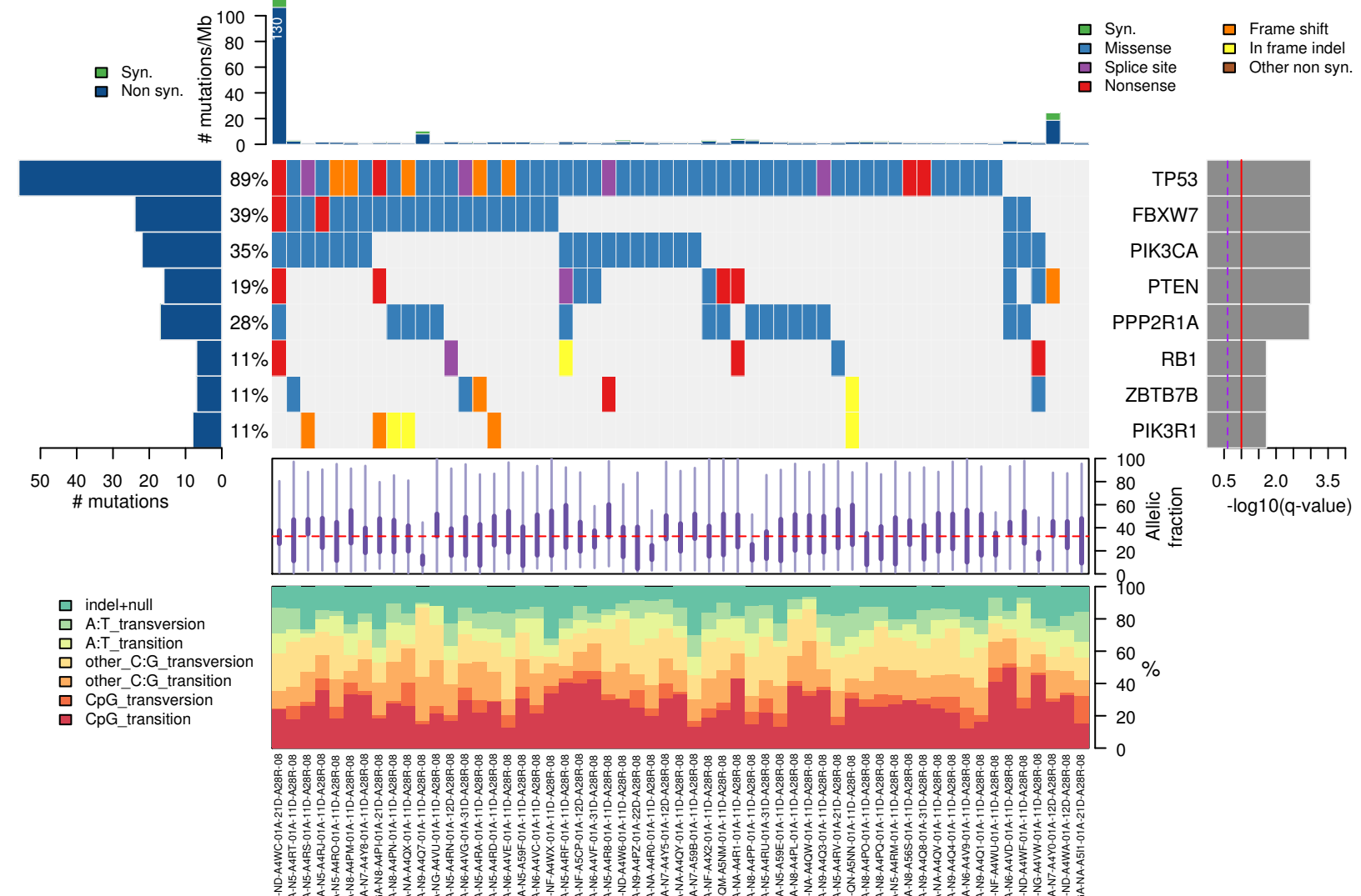
Syn.
Non syn.
mutations/Mb
130
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
89%
39%
35%
19%
28%
11%
11%
11%
TP53
FBXW7
PIK3CA
PTEN
PPP2R1A
RB1
ZBTB7B
PIK3R1
50 40 30 20 10 0
mutations
0.5 2.0 3.5
-log10(q-value)
Allelic fraction
indel+null
A:T_transversion
A:T_transition
other_C:G_transversion
other_C:G_transition
CpG_transversion
CpG_transition
%