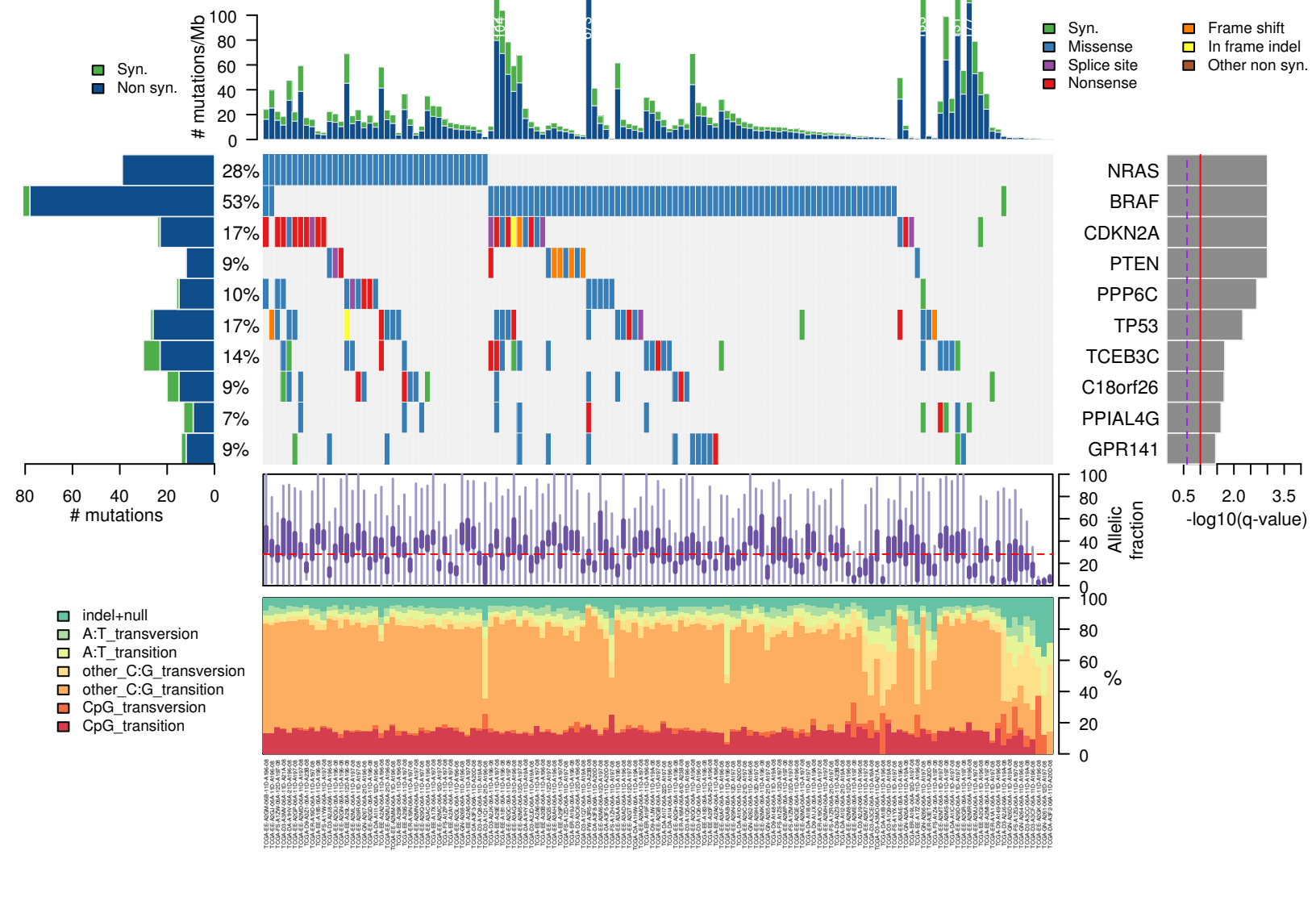
Syn.
Non syn.
mutations/Mb
100
80
60
40
20
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
28%
53%
17%
9%
10%
17%
14%
9%
7%
9%
80
60
40
20
0
mutations
NRAS
BRAF
CDKN2A
PTEN
PPP6C
TP53
TCEB3C
C18orf26
PPIAL4G
GPR141
0.5
2.0
3.5
-log10(q-value)
Allelic fraction
indel+null
A:T_transversion
A:T_transition
other_C:G_transversion
other_C:G_transition
CpG_transversion
CpG_transition
%