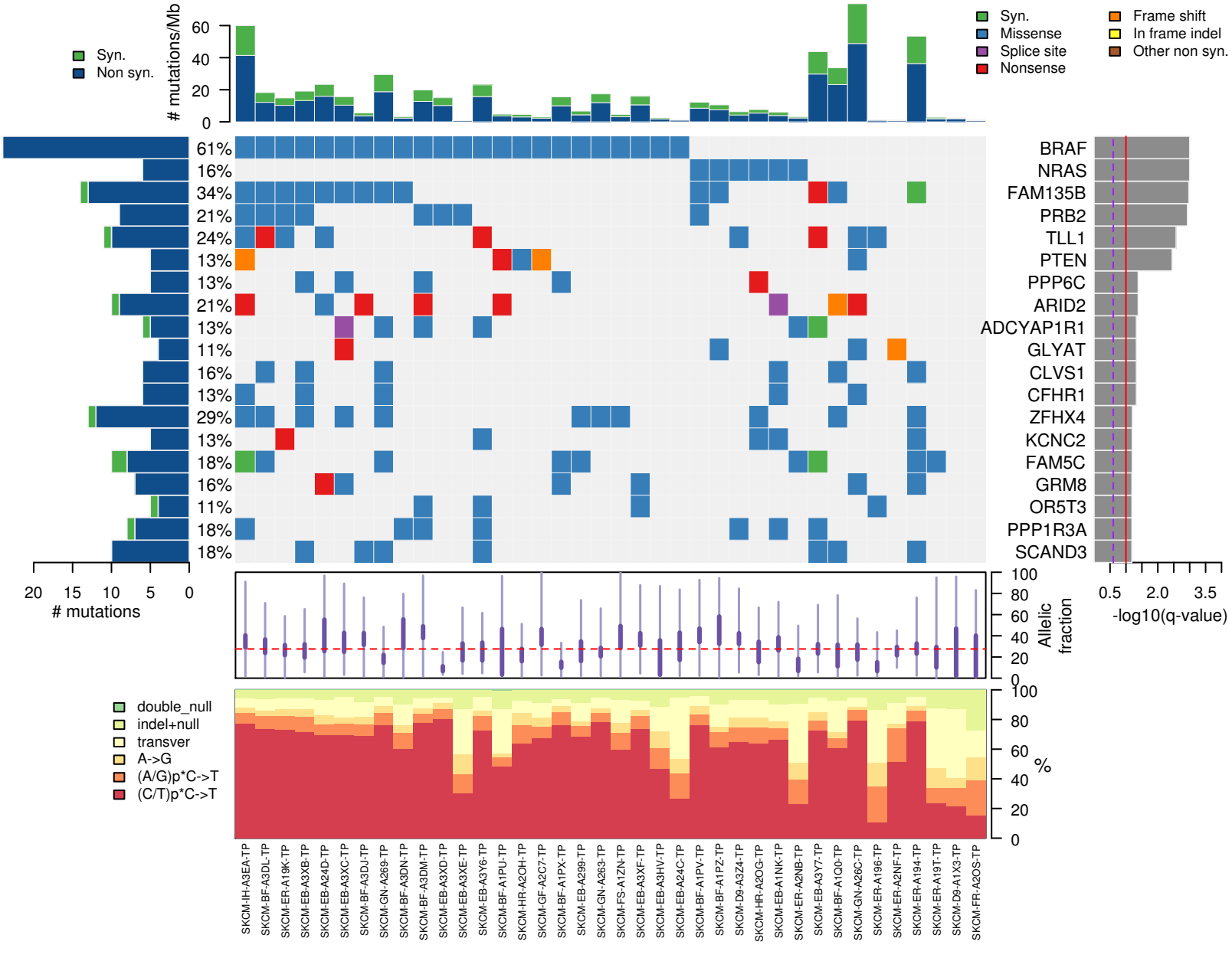
Syn.
Non syn.
mutations/Mb
60
40
20
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
61%
16%
34%
21%
24%
13%
13%
21%
13%
11%
16%
13%
29%
13%
18%
16%
11%
18%
18%
BRAF
NRAS
FAM135B
PRB2
TLL1
PTEN
PPP6C
ARID2
ADCYAP1R1
GLYAT
CLVS1
CFHR1
ZFHX4
KCNC2
FAM5C
GRM8
OR5T3
PPP1R3A
SCAND3
20 15 10 5 0
mutations
0.5 2.0 3.5
-log10(q-value)
Allelic fraction
double_null
indel+null
transver
A->G
(A/G)p*C->T
(C/T)p*C->T
%
SKCM-IH-A3EA-TP
SKCM-BF-A3DL-TP
SKCM-ER-A19K-TP
SKCM-EB-A3XB-TP
SKCM-EB-A24D-TP
SKCM-EB-A3XC-TP
SKCM-BF-A3DJ-TP
SKCM-GN-A269-TP
SKCM-BF-A3DN-TP
SKCM-BF-A3DM-TP
SKCM-EB-A3XD-TP
SKCM-EB-A3XE-TP
SKCM-EB-A3Y6-TP
SKCM-BF-A1PU-TP
SKCM-HR-A2OH-TP
SKCM-GF-A2C7-TP
SKCM-BF-A1PX-TP
SKCM-EB-A299-TP
SKCM-GN-A263-TP
SKCM-FS-A1ZN-TP
SKCM-EB-A3XF-TP
SKCM-EB-A3HV-TP
SKCM-EB-A24C-TP
SKCM-BF-A1PV-TP
SKCM-BF-A1PZ-TP
SKCM-D9-A3Z4-TP
SKCM-HR-A2OG-TP
SKCM-EB-A1NK-TP
SKCM-ER-A2NB-TP
SKCM-EB-A3Y7-TP
SKCM-BF-A1Q0-TP
SKCM-GN-A26C-TP
SKCM-ER-A196-TP
SKCM-ER-A2NF-TP
SKCM-ER-A194-TP
SKCM-ER-A19T-TP
SKCM-D9-A1X3-TP
SKCM-FR-A2OS-TP