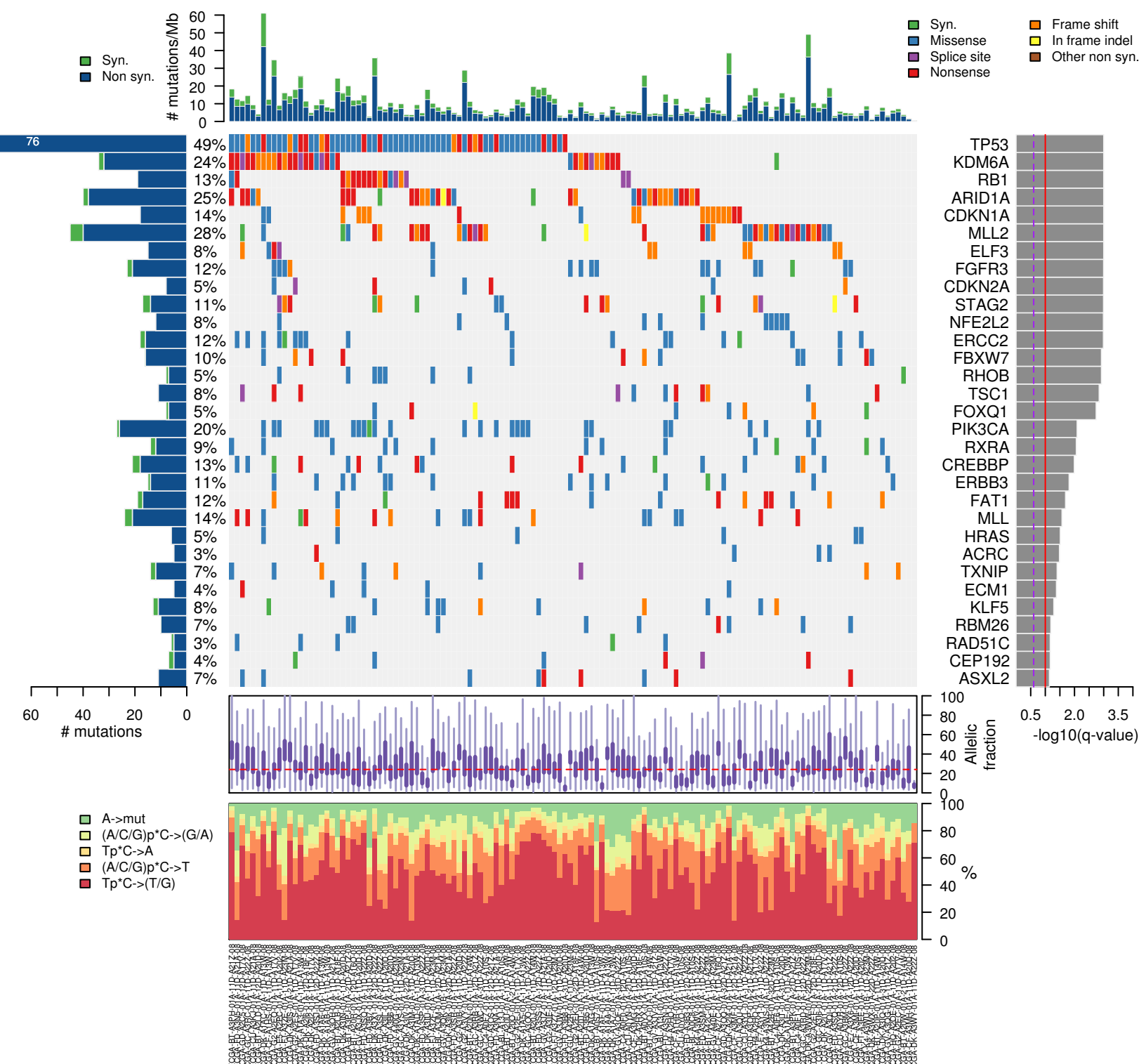
Syn.
Non syn.
# mutations/Mb
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
76
49%
24%
13%
25%
14%
28%
8%
12%
5%
11%
8%
12%
10%
5%
8%
5%
20%
9%
13%
11%
12%
14%
5%
3%
7%
4%
8%
7%
3%
4%
7%
TP53
KDM6A
RB1
ARID1A
CDKN1A
MLL2
ELF3
FGFR3
CDKN2A
STAG2
NFE2L2
ERCC2
FBXW7
RHOB
TSC1
FOXQ1
PIK3CA
RXRA
CREBBP
ERBB3
FAT1
MLL
HRAS
ACRC
TXNIP
ECM1
KLF5
RBM26
RAD51C
CEP192
ASXL2
# mutations
Allelic fraction
-log10(q-value)
A->mut
(A/C/G)p*C->(G/A)
Tp*C->A
(A/C/G)p*C->T
Tp*C->(T/G)
%