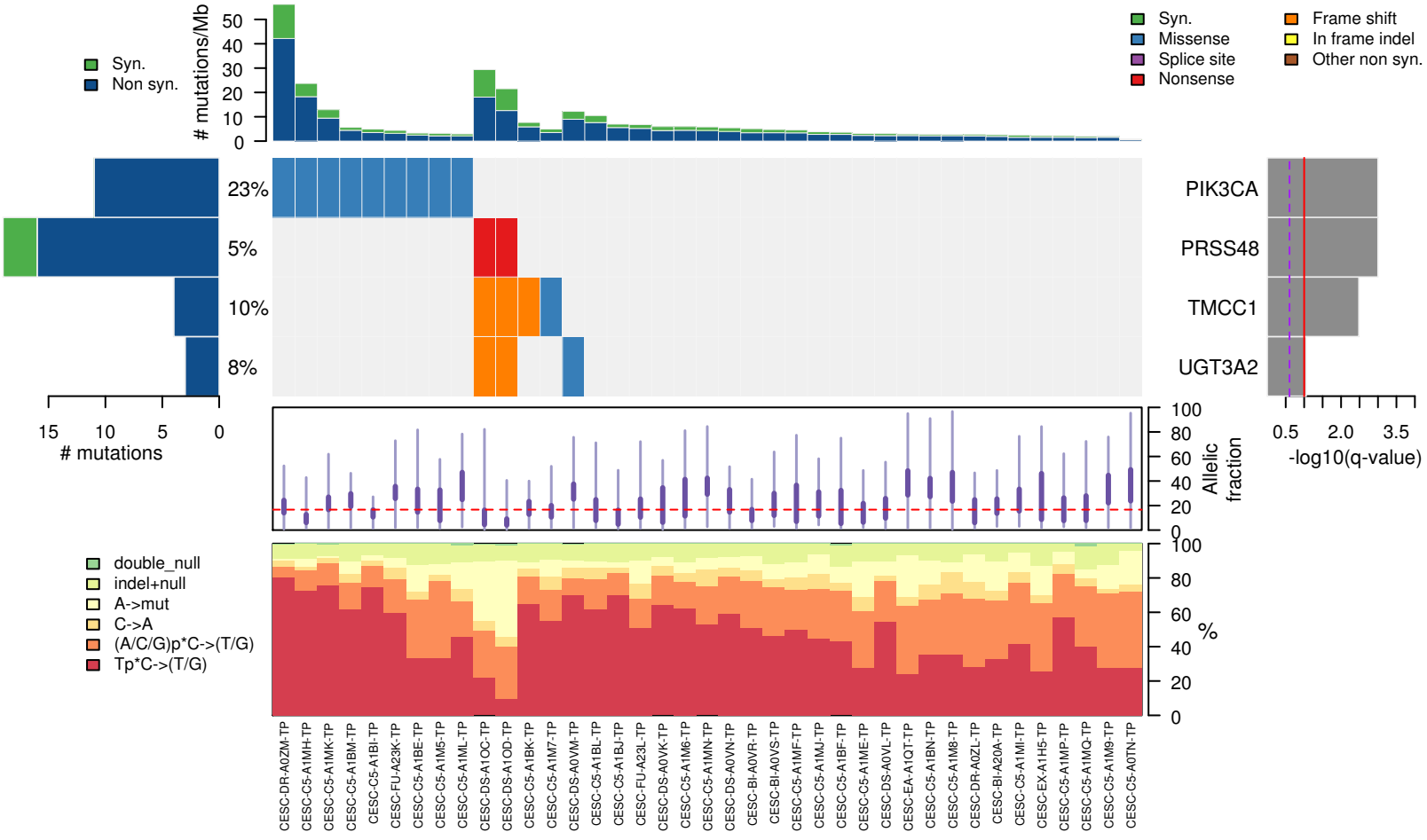
Syn.
Non syn.
mutations/Mb
50
40
30
20
10
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
23%
5%
10%
8%
15
10
5
0
mutations
PIK3CA
PRSS48
TMCC1
UGT3A2
0.5
2.0
3.5
-log10(q-value)
100
80
60
40
20
0
Allelic fraction
double_null
indel+null
A->mut
C->A
(A/C/G)p*C->(T/G)
Tp*C->(T/G)
%
CESC-DR-A0ZM-TP
CESC-C5-A1MH-TP
CESC-C5-A1MK-TP
CESC-C5-A1BM-TP
CESC-C5-A1BI-TP
CESC-FU-A23K-TP
CESC-C5-A1BE-TP
CESC-C5-A1M5-TP
CESC-C5-A1ML-TP
CESC-DS-A1OC-TP
CESC-DS-A1OD-TP
CESC-C5-A1BK-TP
CESC-C5-A1M7-TP
CESC-DS-A0VM-TP
CESC-C5-A1BL-TP
CESC-C5-A1BJ-TP
CESC-FU-A23L-TP
CESC-DS-A0VK-TP
CESC-C5-A1M6-TP
CESC-C5-A1MN-TP
CESC-DS-A0VN-TP
CESC-BI-A0VR-TP
CESC-BI-A0VS-TP
CESC-C5-A1MF-TP
CESC-C5-A1MJ-TP
CESC-C5-A1BF-TP
CESC-C5-A1ME-TP
CESC-DS-A0VL-TP
CESC-EA-A1QT-TP
CESC-C5-A1BN-TP
CESC-C5-A1M8-TP
CESC-DR-A0ZL-TP
CESC-BI-A20A-TP
CESC-C5-A1MI-TP
CESC-EX-A1H5-TP
CESC-C5-A1MP-TP
CESC-C5-A1MQ-TP
CESC-C5-A1M9-TP
CESC-C5-A0TN-TP