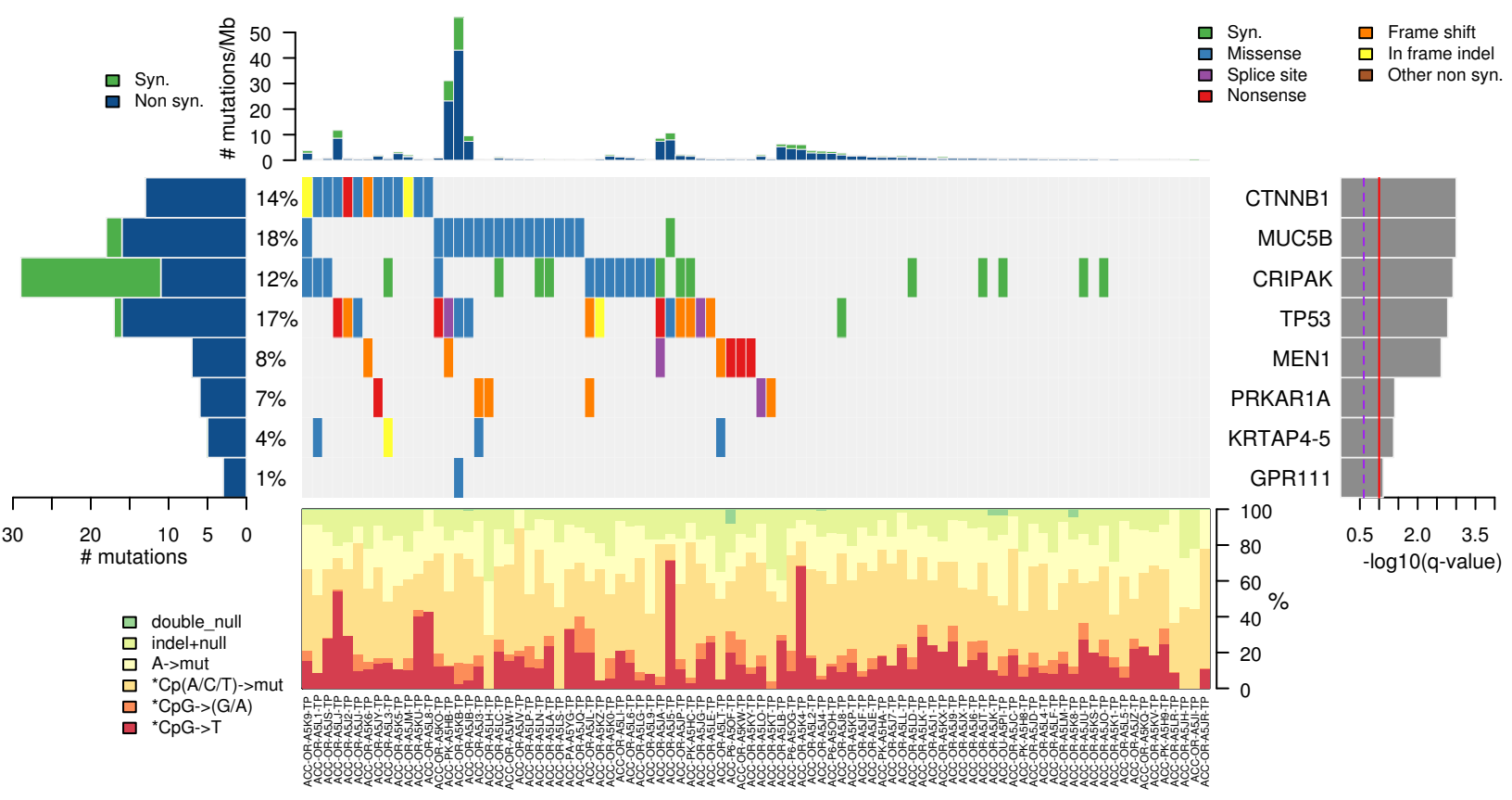
Syn.
Non syn.
mutations/Mb
50
40
30
20
10
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
14%
18%
12%
17%
8%
7%
4%
1%
30
20
10
5
0
mutations
CTNNB1
MUC5B
CRIPAK
TP53
MEN1
PRKAR1A
KRTAP4-5
GPR111
0.5
2.0
3.5
-log10(q-value)
double_null
indel+null
A->mut
*Cp(A/C/T)->mut
*CpG->(G/A)
*CpG->T
100
80
60
40
20
0
%