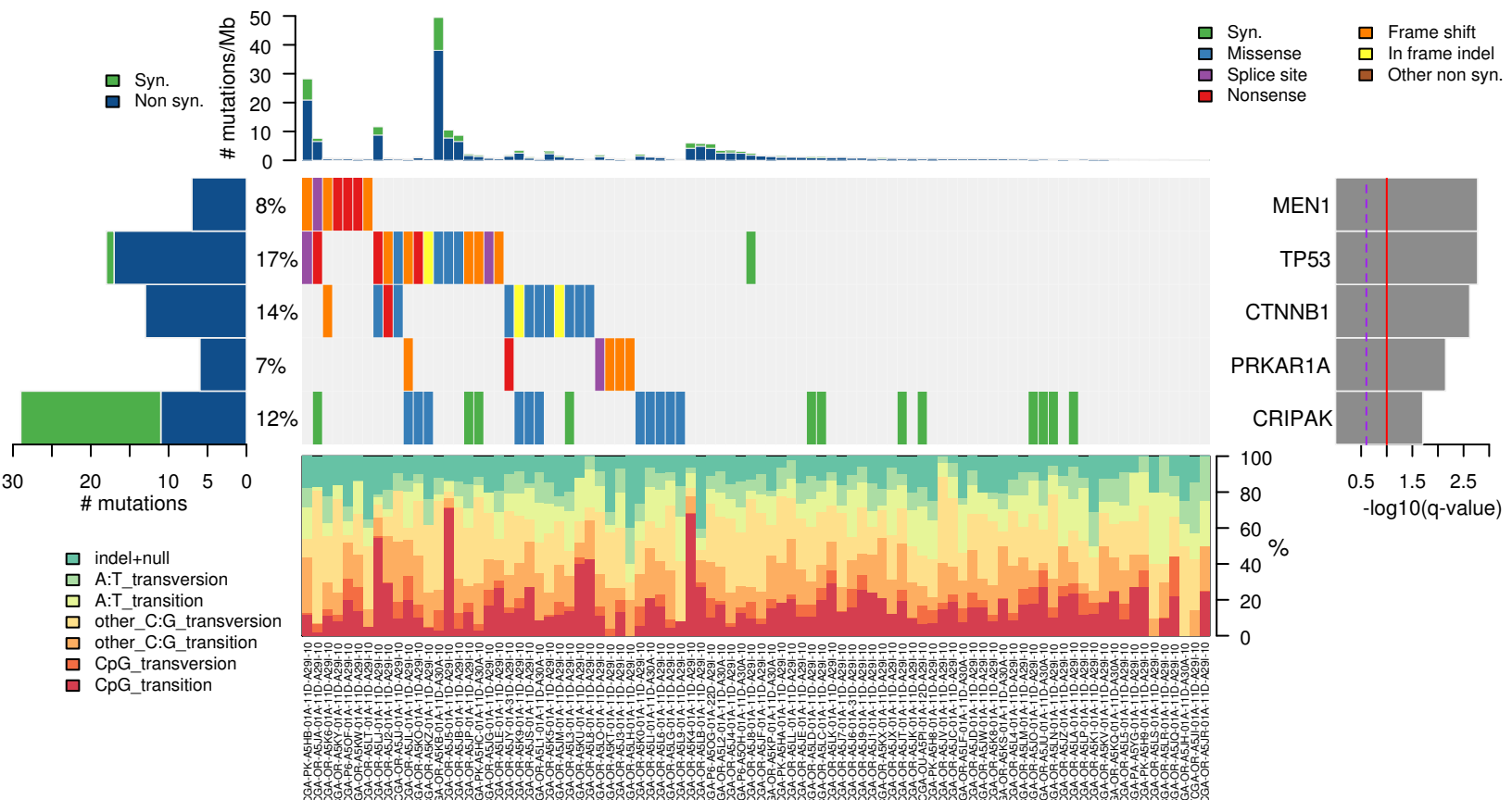
Syn.
Non syn.
mutations/Mb
50
40
30
20
10
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
8%
17%
14%
7%
12%
30
20
10
5
0
mutations
MEN1
TP53
CTNNB1
PRKAR1A
CRIPAK
0.5
1.5
2.5
-log10(q-value)
indel+null
A:T_transversion
A:T_transition
other_C:G_transversion
other_C:G_transition
CpG_transversion
CpG_transition
100
80
60
40
20
0
%