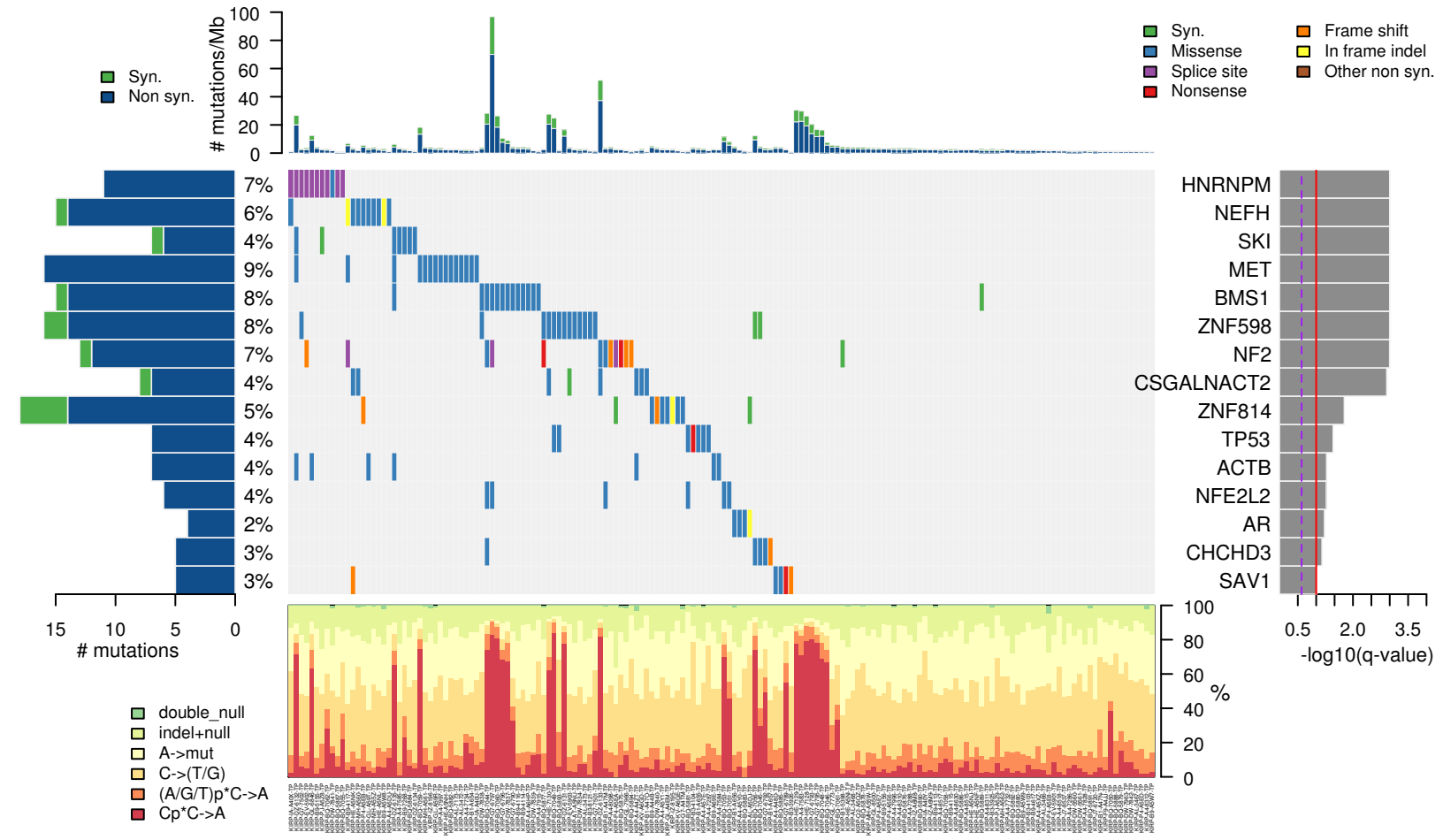
Syn.
Non syn.
mutations/Mb
100
80
60
40
20
0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
7%
6%
4%
9%
8%
8%
7%
4%
5%
4%
4%
4%
2%
3%
3%
HNRNPM
NEFH
SKI
MET
BMS1
ZNF598
NF2
CSGALNACT2
ZNF814
TP53
ACTB
NFE2L2
AR
CHCHD3
SAV1
15
10
5
0
mutations
0.5
2.0
3.5
-log10(q-value)
double_null
indel+null
A->mut
C->(T/G)
(A/G/T)p*C->A
Cp*C->A
100
80
60
40
20
0
%