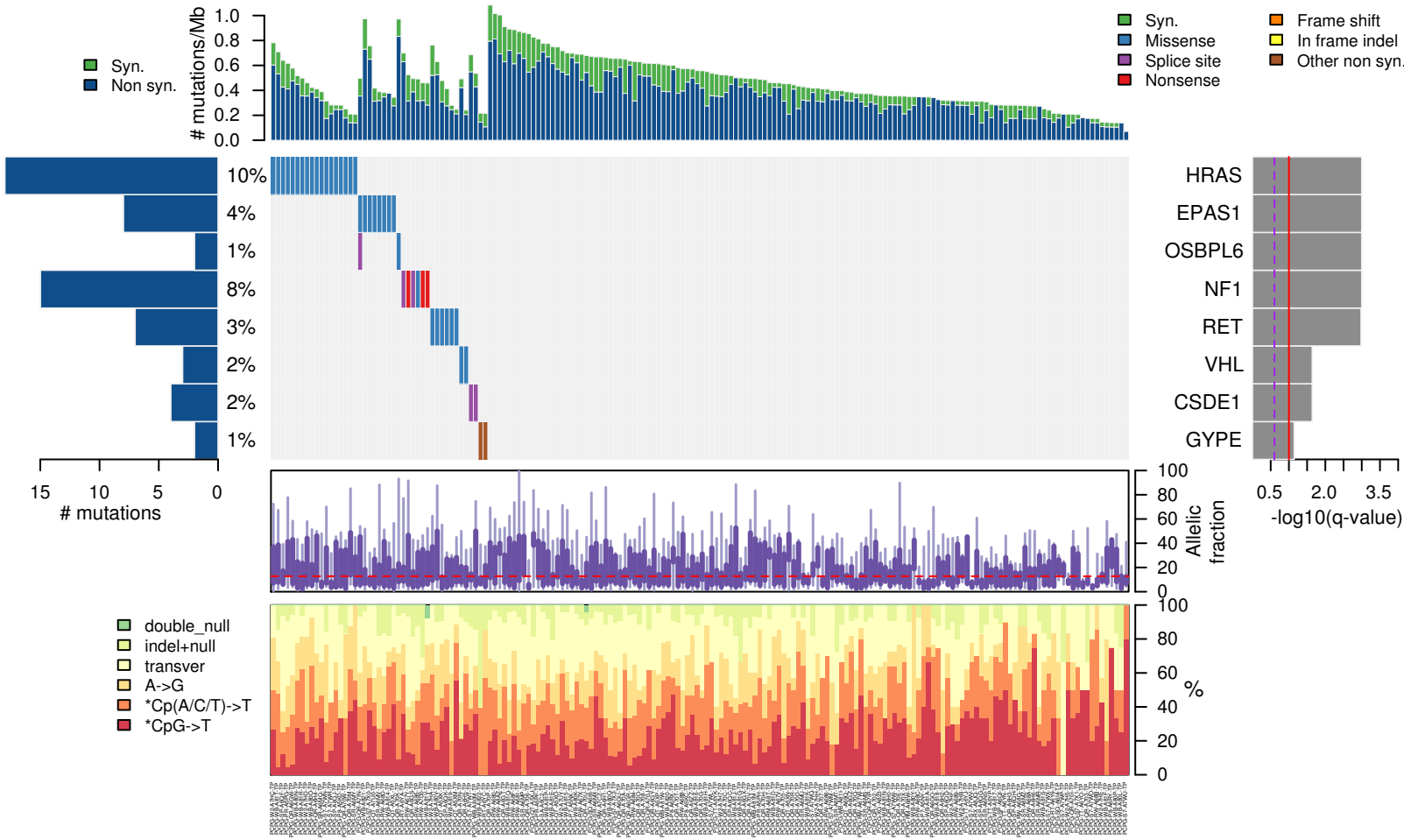
Syn.
Non syn.
mutations/Mb
1.0
0.8
0.6
0.4
0.2
0.0
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
10%
4%
1%
8%
3%
2%
2%
1%
15
10
5
0
mutations
HRAS
EPAS1
OSBPL6
NF1
RET
VHL
CSDE1
GYPE
0.5
2.0
3.5
-log10(q-value)
100
80
60
40
20
0
Allelic fraction
double_null
indel+null
transver
A->G
*Cp(A/C/T)->T
*CpG->T
100
80
60
40
20
0
%