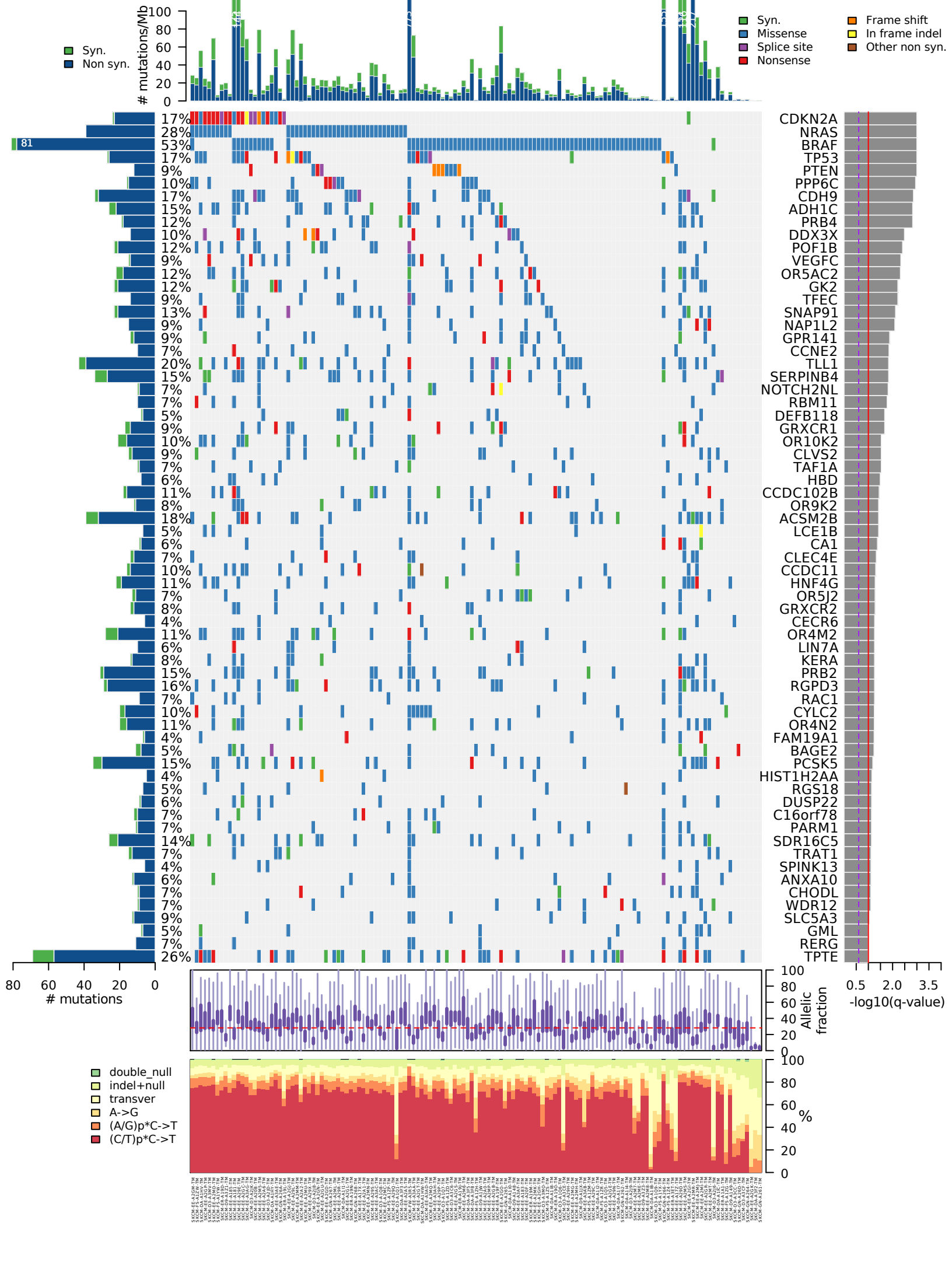
Syn.
Non syn.
mutations/Mb
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
CDKN2A
NRAS
BRAF
TP53
PTEN
PPP6C
CDH9
ADH1C
PRB4
DDX3X
POF1B
VEGFC
OR5AC2
GK2
TFEC
SNAP91
NAP1L2
GPR141
CCNE2
TLL1
SERPINB4
NOTCH2NL
RBM11
DEFB118
GRXCR1
OR10K2
CLVS2
TAF1A
HBD
CCDC102B
OR9K2
ACSM2B
LCE1B
CA1
CLEC4E
CCDC11
HNF4G
OR5J2
GRXCR2
CECR6
OR4M2
LIN7A
KERA
PRB2
RGPD3
RAC1
CYLC2
OR4N2
FAM19A1
BAGE2
PCSK5
HIST1H2AA
RGS18
DUSP22
C16orf78
PARM1
SDR16C5
TRAT1
SPINK13
ANXA10
CHODL
WDR12
SLC5A3
GML
RERG
TPTE
mutations
-log10(q-value)
Allelic fraction
double_null
indel+null
transver
A->G
(A/G)p*C->T
(C/T)p*C->T
%