

of unique mutations

20
15
10
5
0

Inframe deletion
Missense variant
Splice variant
Truncating variant

DNMT3A
TET2
BCOR
CHEK2
TP53
CBL
ZNF318
EP300
PPM1D
SRCAP
ARID2
ASXL1
BRAF
BRCC3
CEBPA
CUX1
NF2
PHF6
PHIP
SETD2
SH2B3
WT1
ZBTB33
ZNF234
ZRSR2

