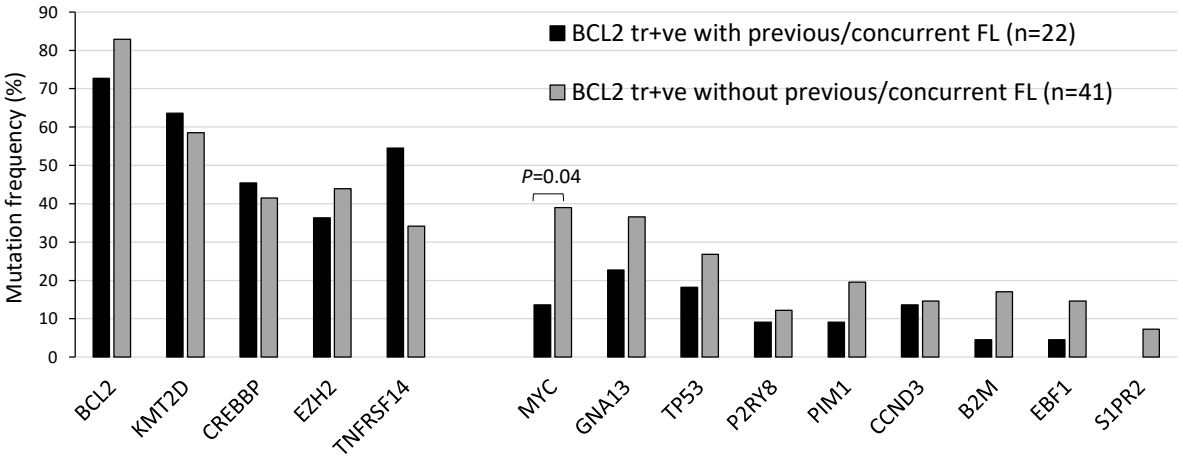
Figure S3


Mutation frequency (%)
90
80
70
60
50
40
30
20
10
0
BCL2 tr+ve with previous/concurrent FL (n=22)
BCL2 tr+ve without previous/concurrent FL (n=41)
P=0.04
BCL2
KMT2D
CREBBP
EZH2
TNFRSF14
MYC
GNA13
TP53
P2RY8
PIM1
CCND3
B2M
EBF1
S1PR2