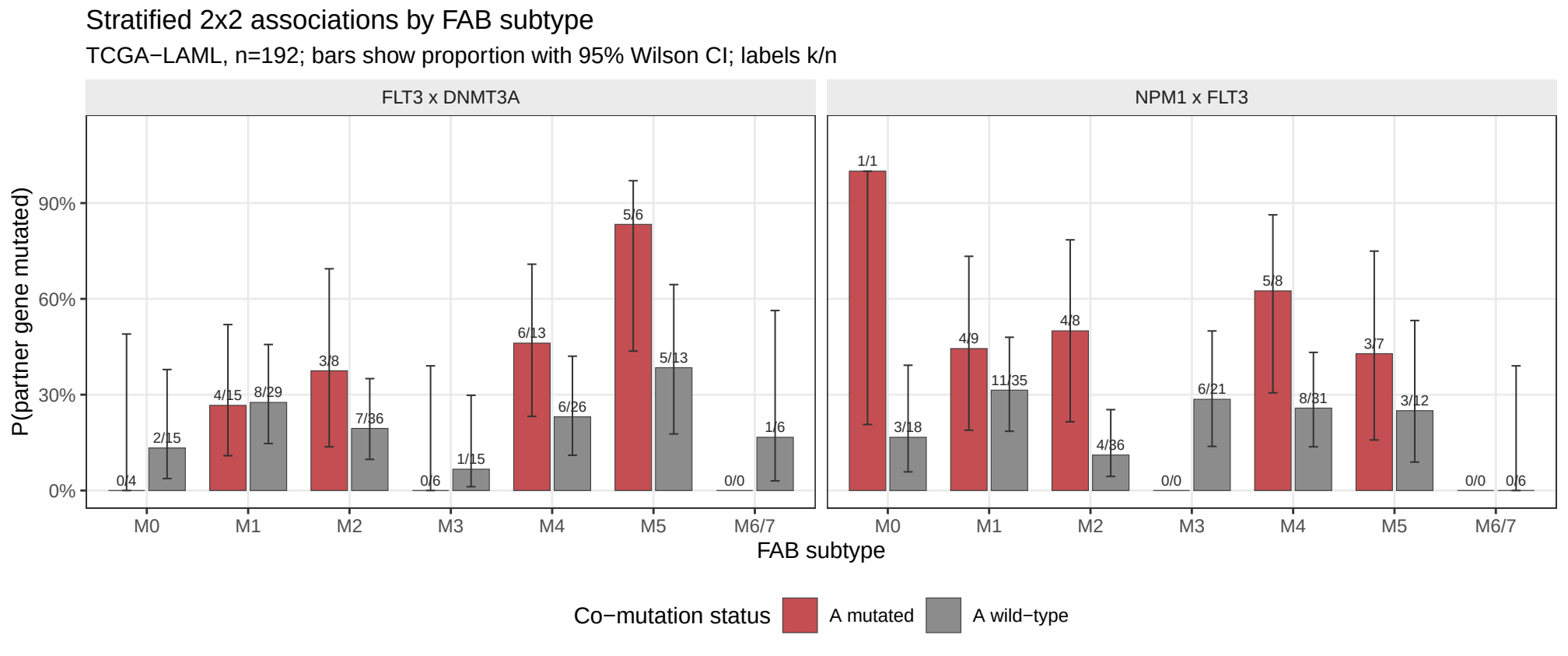

Stratified 2x2 associations by FAB subtype
TCGA−LAML, n=192; bars show proportion with 95% Wilson CI; labels k/n
FLT3 x DNMT3A
NPM1 x FLT3
P(partner gene mutated)
90%
60%
30%
0%
0/4
2/15
4/15
8/29
3/8
7/86
0/6
1/15
6/13
6/26
5/6
5/13
0/0
1/6
1/1
3/18
4/9
11/35
4/8
4/36
0/0
6/21
5/8
8/31
3/7
3/12
0/0
0/6
M0
M1
M2
M3
M4
M5
M6/7
FAB subtype
Co−mutation status
A mutated
A wild−type