

Number of co-occurring mutations

40

30

20

10

0

$p = 2e-06$

$p = 1$

$p = 0.00015$

$p = 0.58$

observed  
expected

BRAF.Class.I  
and  
NF1.LOF

BRAF.Class.II  
and  
NF1.LOF

BRAF.Class.III  
and  
NF1.LOF

NRAS.G12.G13.Q61  
and  
BRAF.Class.III

