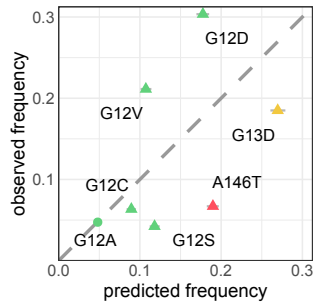


a

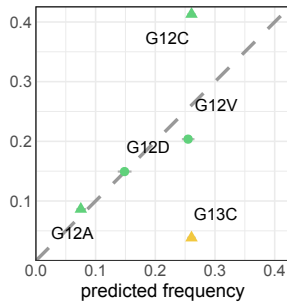
COAD

Pearson correlation: 0.454



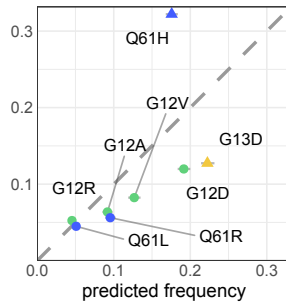
LUAD

Pearson correlation: 0.409



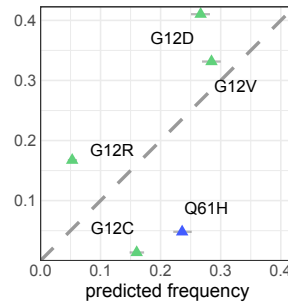
MM

Pearson correlation: 0.607

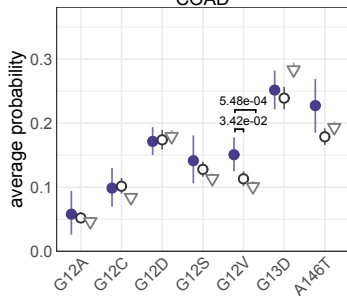


PAAD

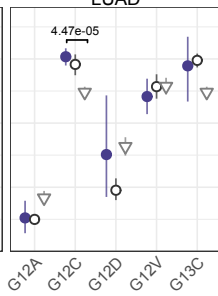
Pearson correlation: 0.484

**b**

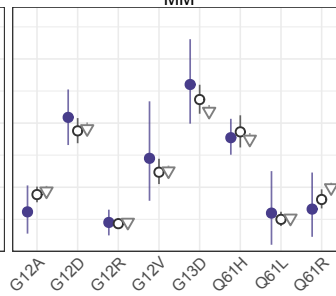
COAD



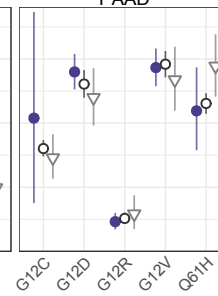
LUAD



MM



PAAD



tumor samples with

- the KRAS allele
- another KRAS mutation
- KRAS WT