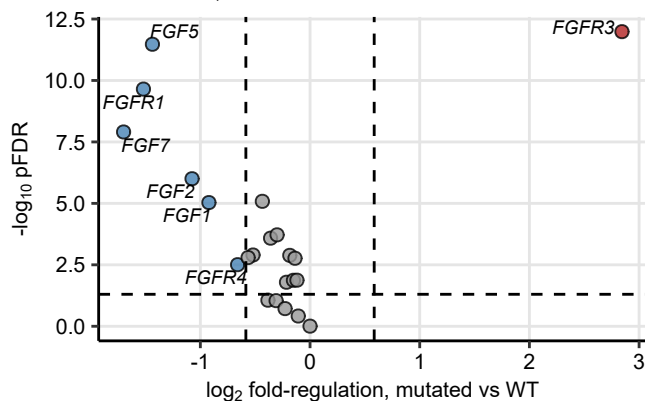
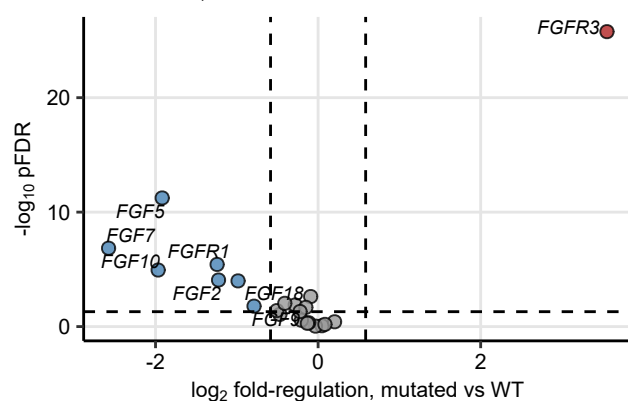
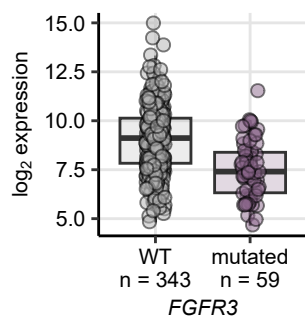
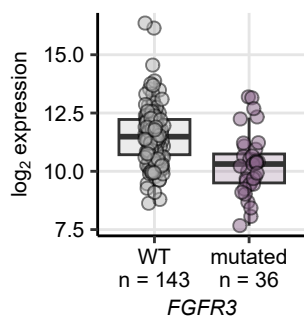
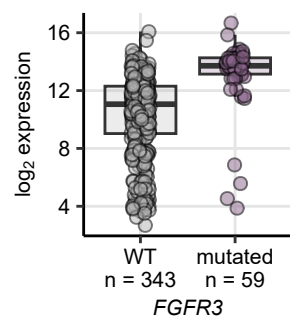
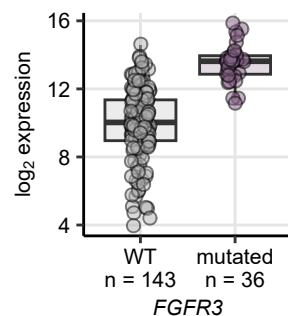
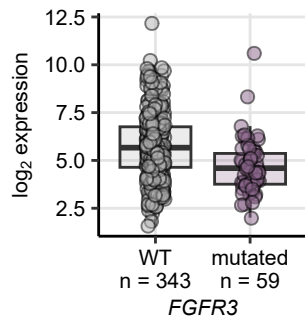
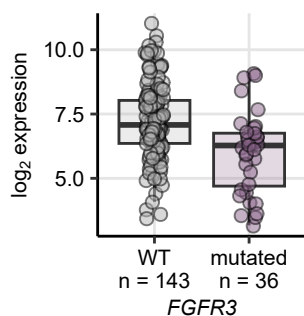
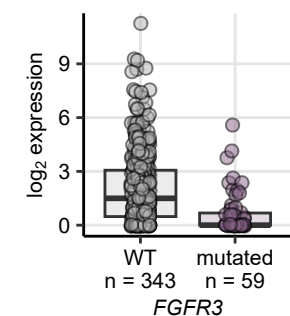
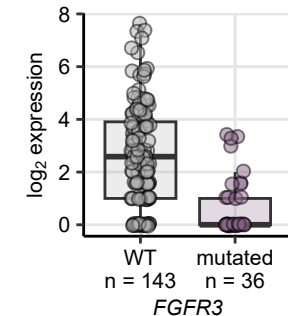
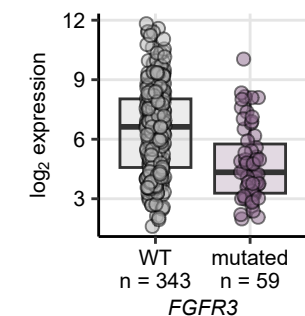


A***FGFR3* mutations, TCGA BLCA**

WT: n = 343, mutated: n = 59

***FGFR3* mutations, IMvigor**

WT: n = 143, mutated: n = 36

**B*****FGFR1*, TCGA BLCA**d = -0.97, $p < 0.001$ ***FGFR1*, IMvigor**d = -0.98, $p < 0.001$ ***FGFR3*, TCGA BLCA**d = 1.1, $p < 0.001$ ***FGFR3*, IMvigor**d = 2.1, $p < 0.001$ ***FGF2*, TCGA BLCA**d = -0.69, $p < 0.001$ ***FGF2*, IMvigor**d = -0.82, $p < 0.001$ ***FGF5*, TCGA BLCA**d = -0.86, $p < 0.001$ ***FGF5*, IMvigor**d = -1.2, $p < 0.001$ ***FGF7*, TCGA BLCA**d = -0.84, $p < 0.001$ ***FGF7*, IMvigor**d = -1.2, $p < 0.001$ 