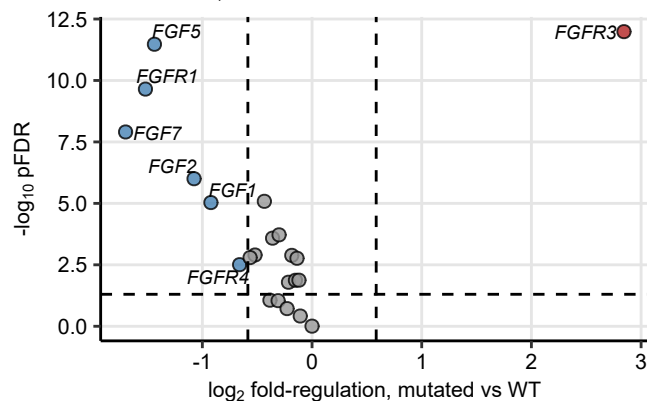
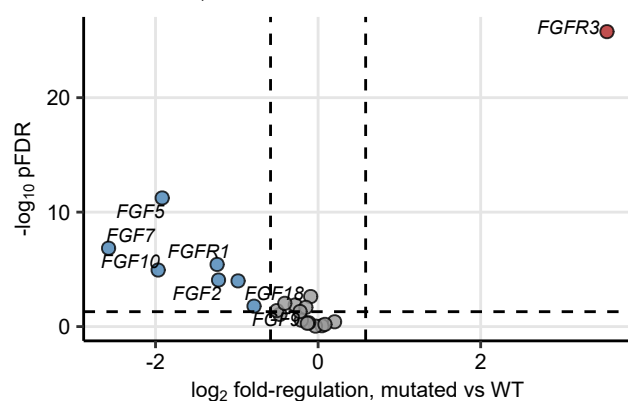


**A****FGFR3 mutations, TCGA BLCA**

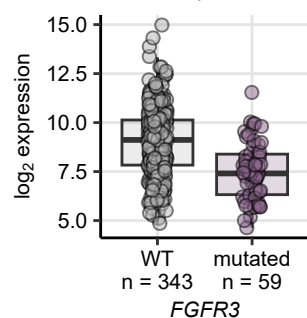
WT: n = 343, mutated: n = 59

**FGFR3 mutations, IMvigor**

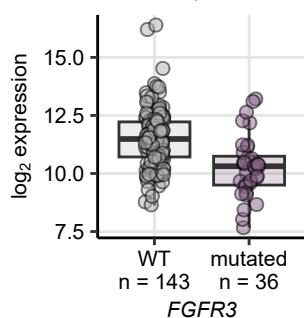
WT: n = 143, mutated: n = 36

**B****FGFR1, TCGA BLCA**

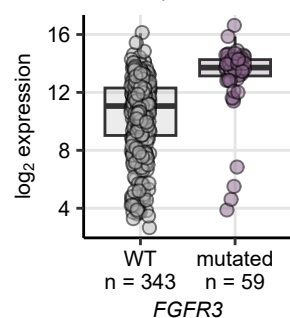
d = -0.97, p &lt; 0.001

**FGFR1, IMvigor**

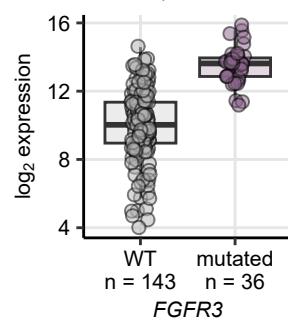
d = -0.98, p &lt; 0.001

**FGFR3, TCGA BLCA**

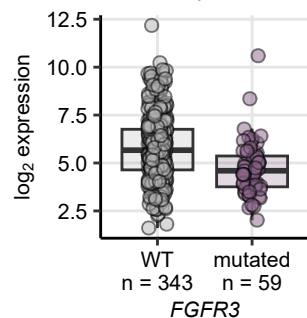
d = 1.1, p &lt; 0.001

**FGFR3, IMvigor**

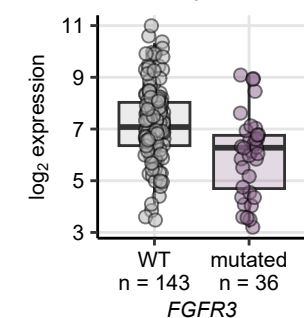
d = 2.1, p &lt; 0.001

**FGF2, TCGA BLCA**

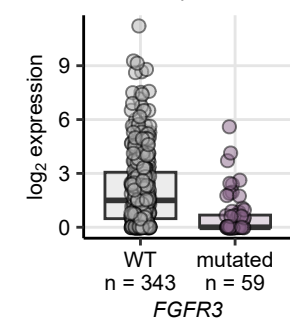
d = -0.69, p &lt; 0.001

**FGF2, IMvigor**

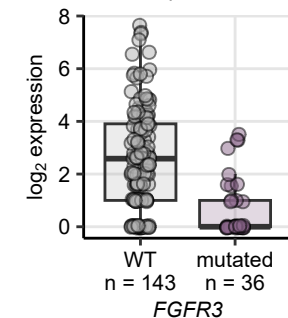
d = -0.82, p &lt; 0.001

**FGF5, TCGA BLCA**

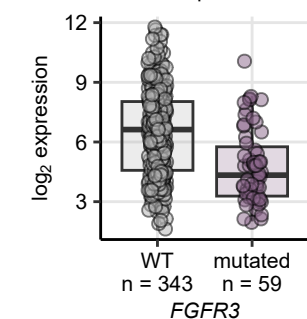
d = -0.86, p &lt; 0.001

**FGF5, IMvigor**

d = -1.2, p &lt; 0.001

**FGF7, TCGA BLCA**

d = -0.84, p &lt; 0.001

**FGF7, IMvigor**

d = -1.2, p &lt; 0.001

