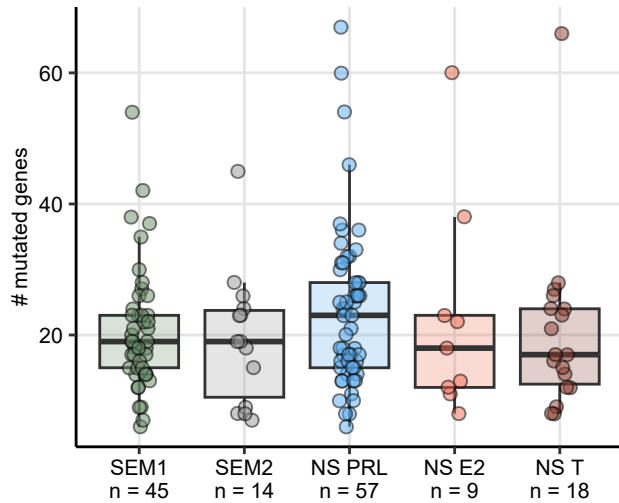


A**Total mutation burden, TCGA** $\eta^2 = -0.0017$, ns ($p = 0.44$)**B****Copy number alterations, TCGA** $\eta^2 = 0.02$, ns ($p = 0.14$)