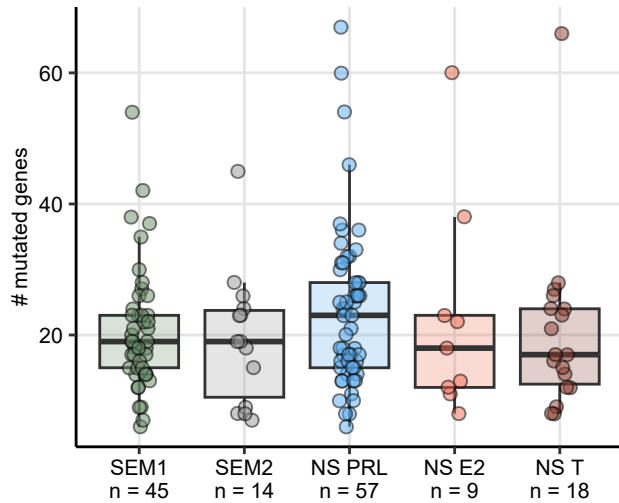


**A****Total mutation burden, TCGA** $\eta^2 = -0.0017$ , ns ( $p = 0.44$ )**B****Copy number alterations, TCGA** $\eta^2 = 0.02$ , ns ( $p = 0.14$ )