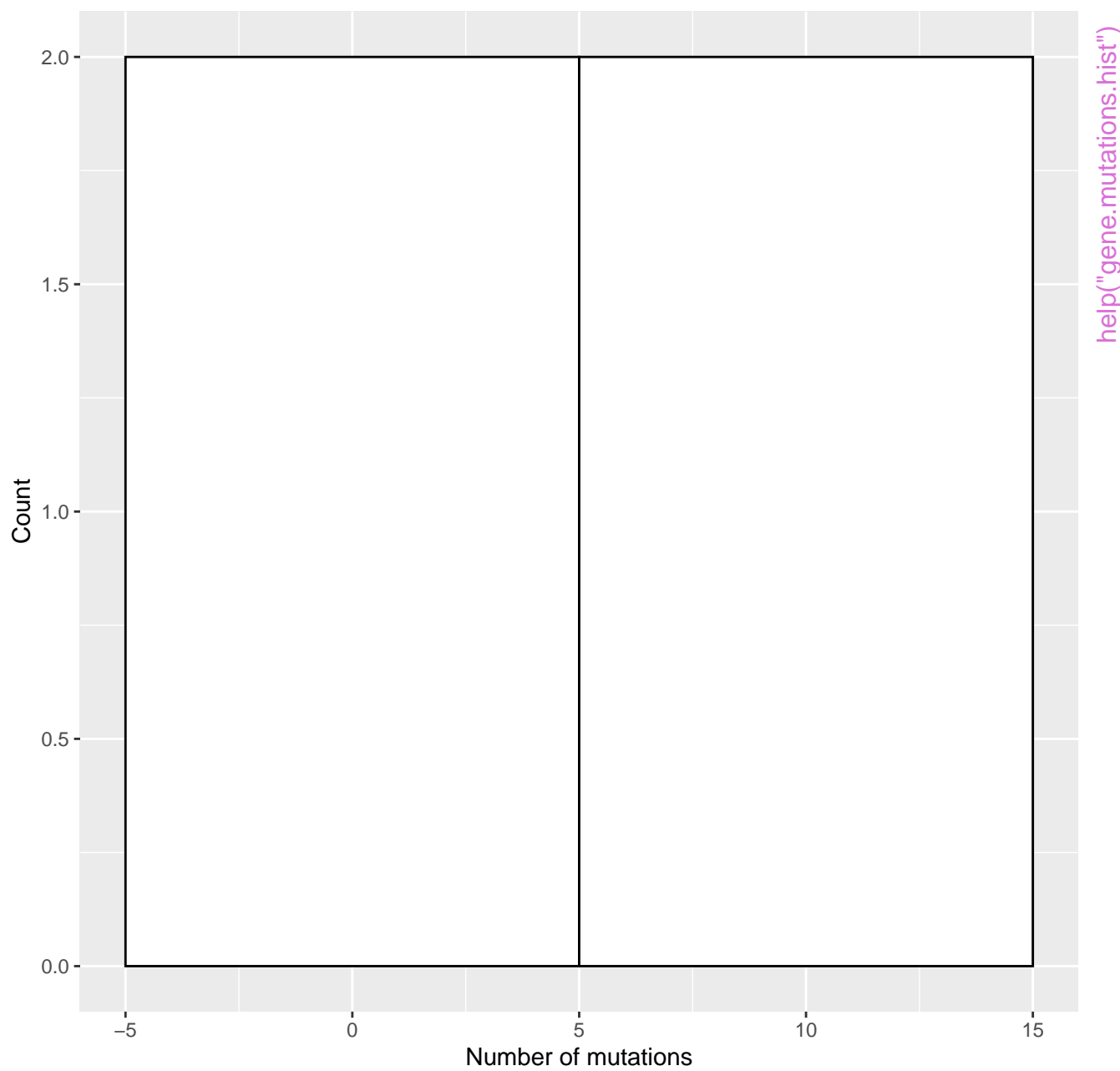


help("draw.ggraph")

Number of mutations per gene (4 genes)



Number of mutations per sample (8 samples)

