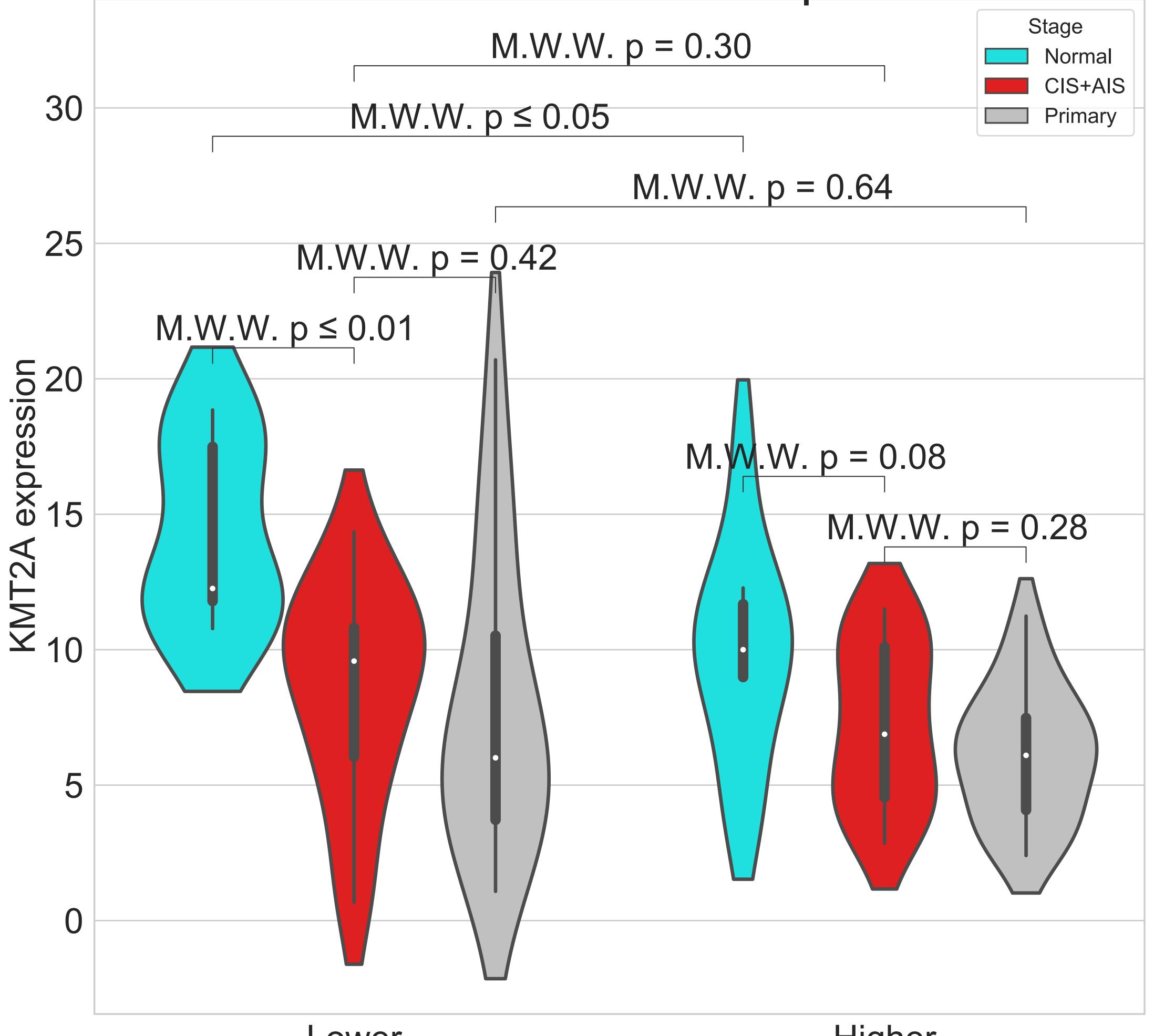
KMT2A: Kruskal-Wallis p=0.001 M.W.W. p = 0.30



Higher Lower Mutation Shared Count per TMB