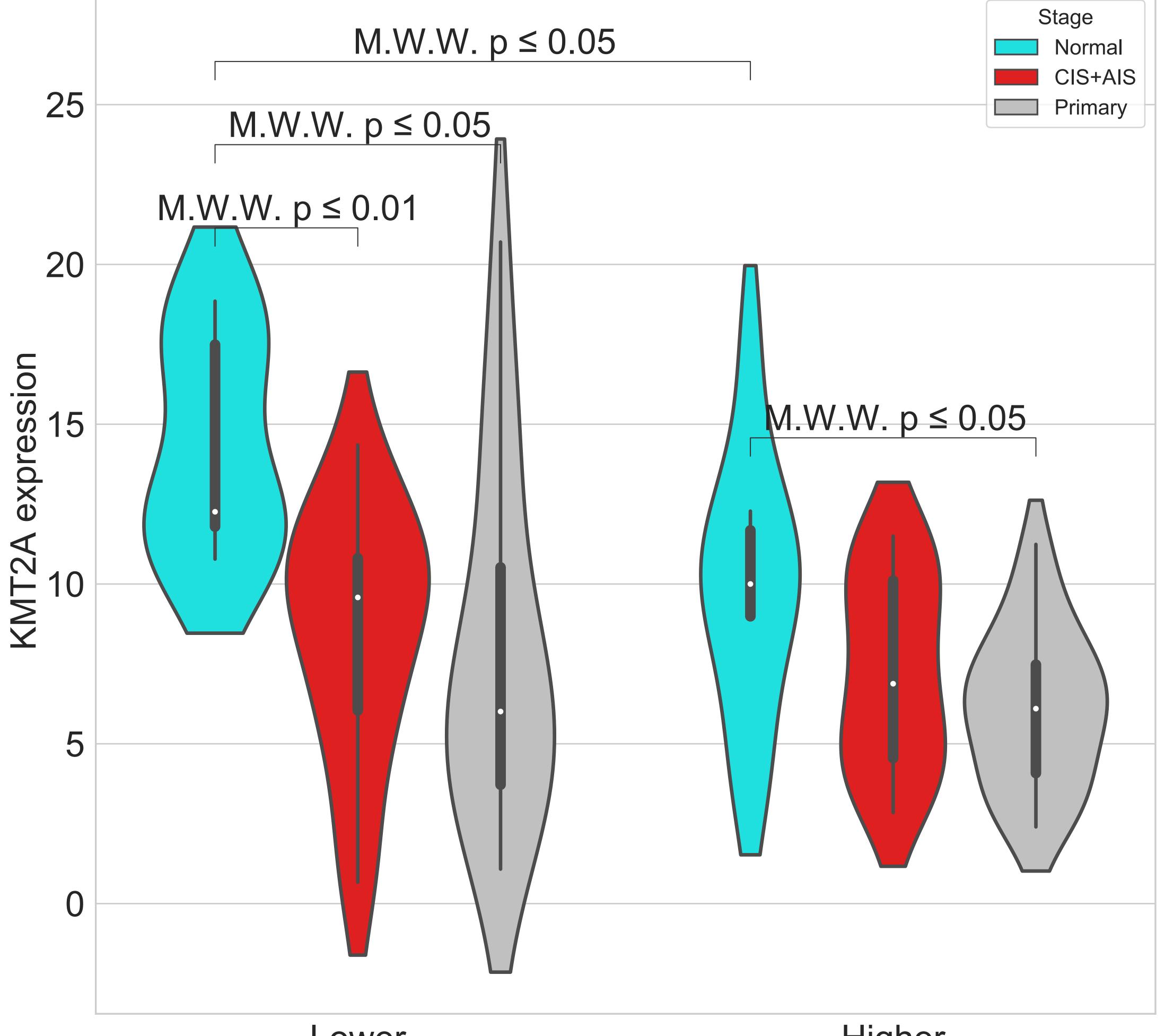
KMT2A: Kruskal-Wallis p=0.001 M.W.W. $p \le 0.05$



Higher Lower Mutation Shared Count per TMB