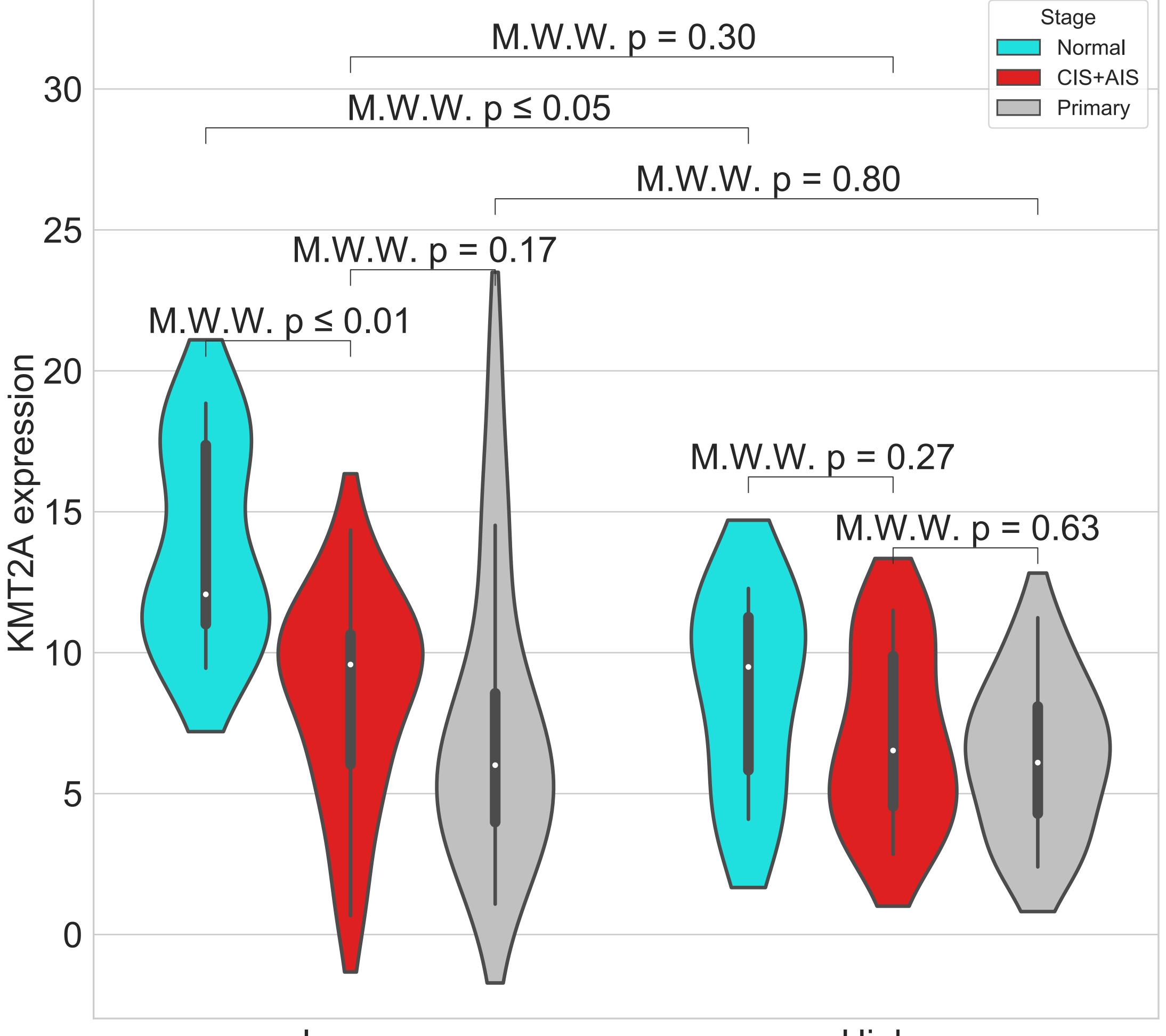
KMT2A: Kruskal-Wallis p=0.001

M.W.W. p = 0.30



Lower
Mutation Shared Count per TMB