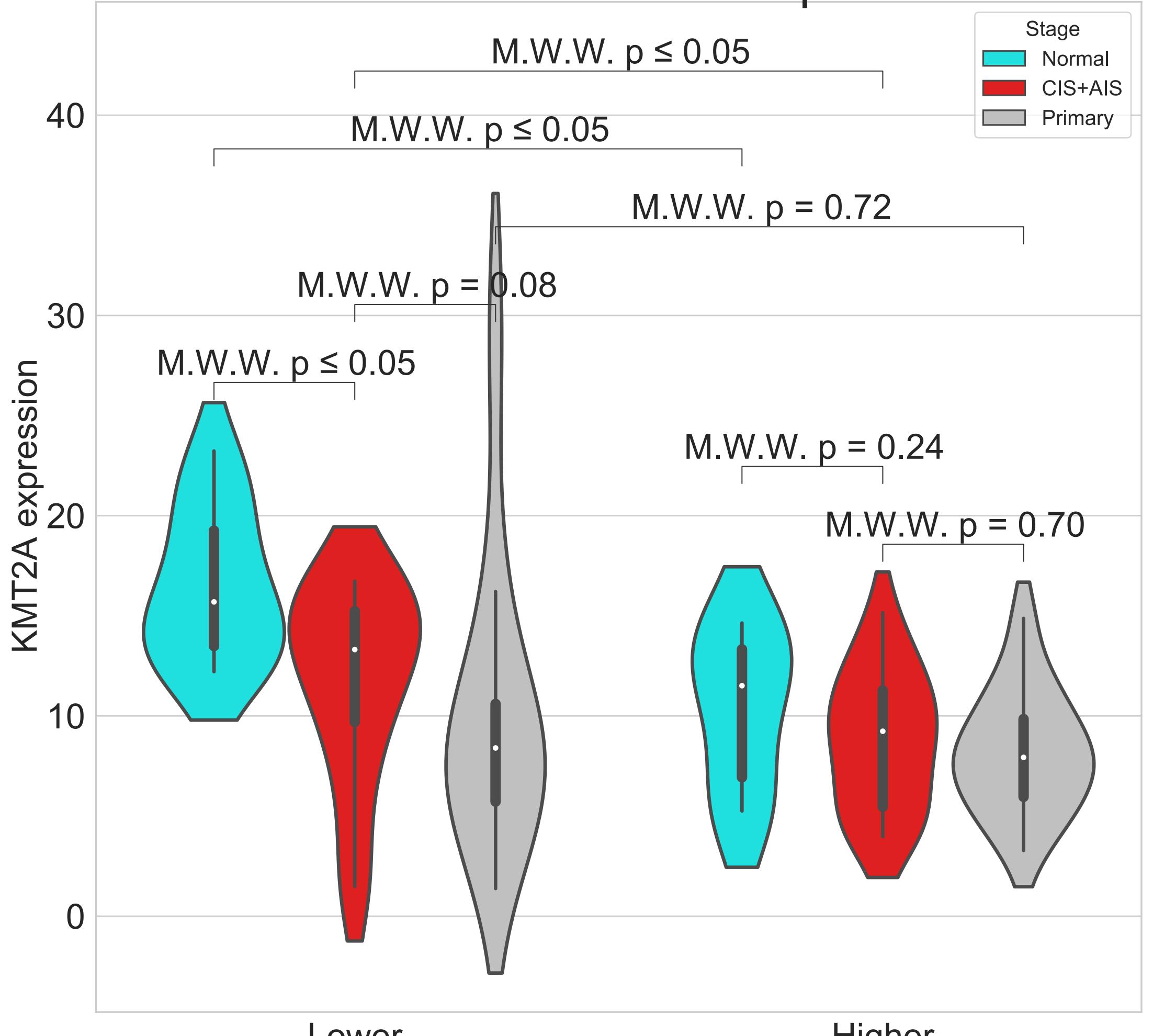
KMT2A: Kruskal-Wallis p=0.001

M.W.W. p ≤ 0.05



Lower
Mutation Shared Count per TMB