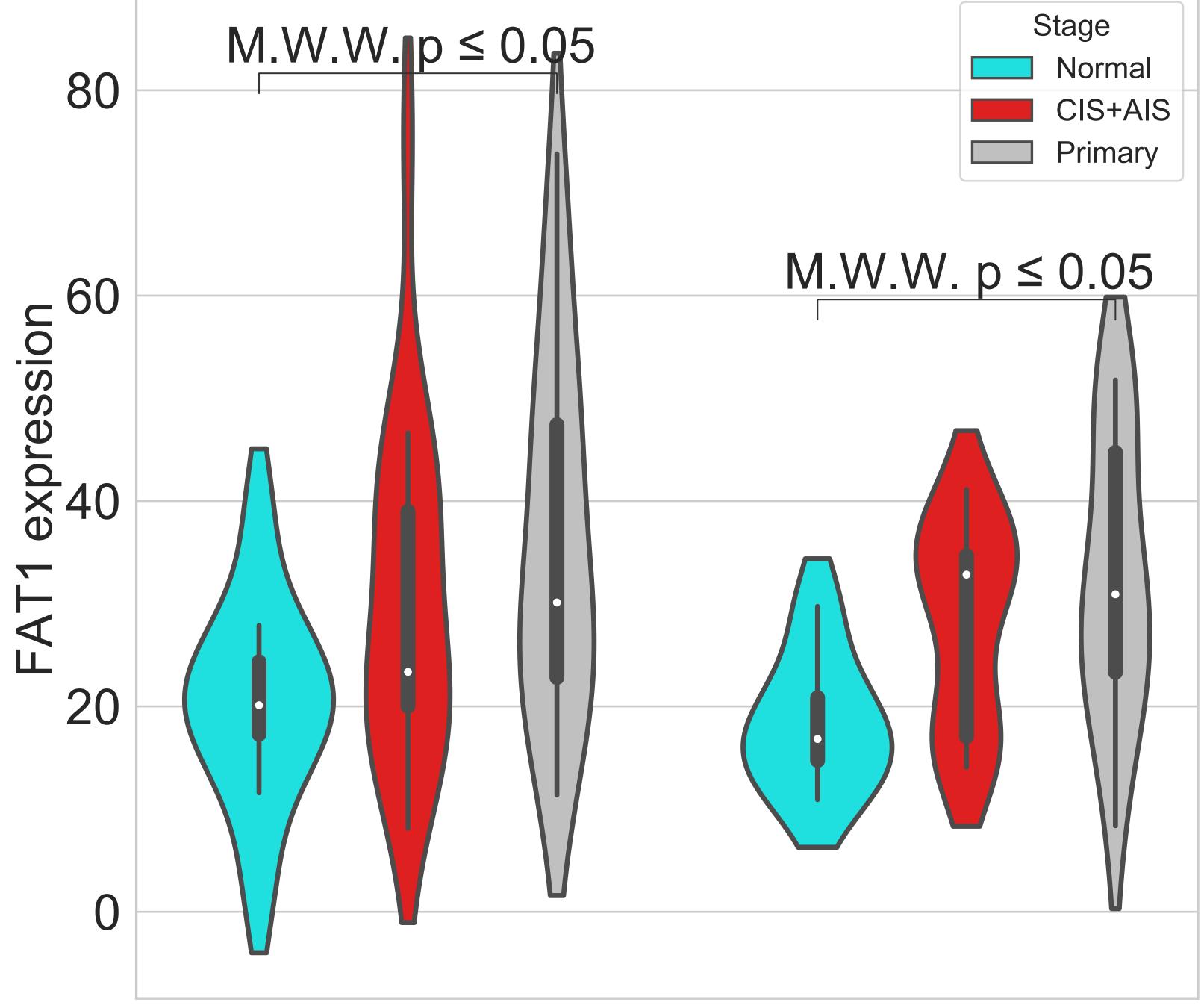
FAT1: Kruskal-Wallis p=0.033

M.W.W. p ≤ 0.05

Stage
Normal



Lower
Mutation Shared Count per TMB