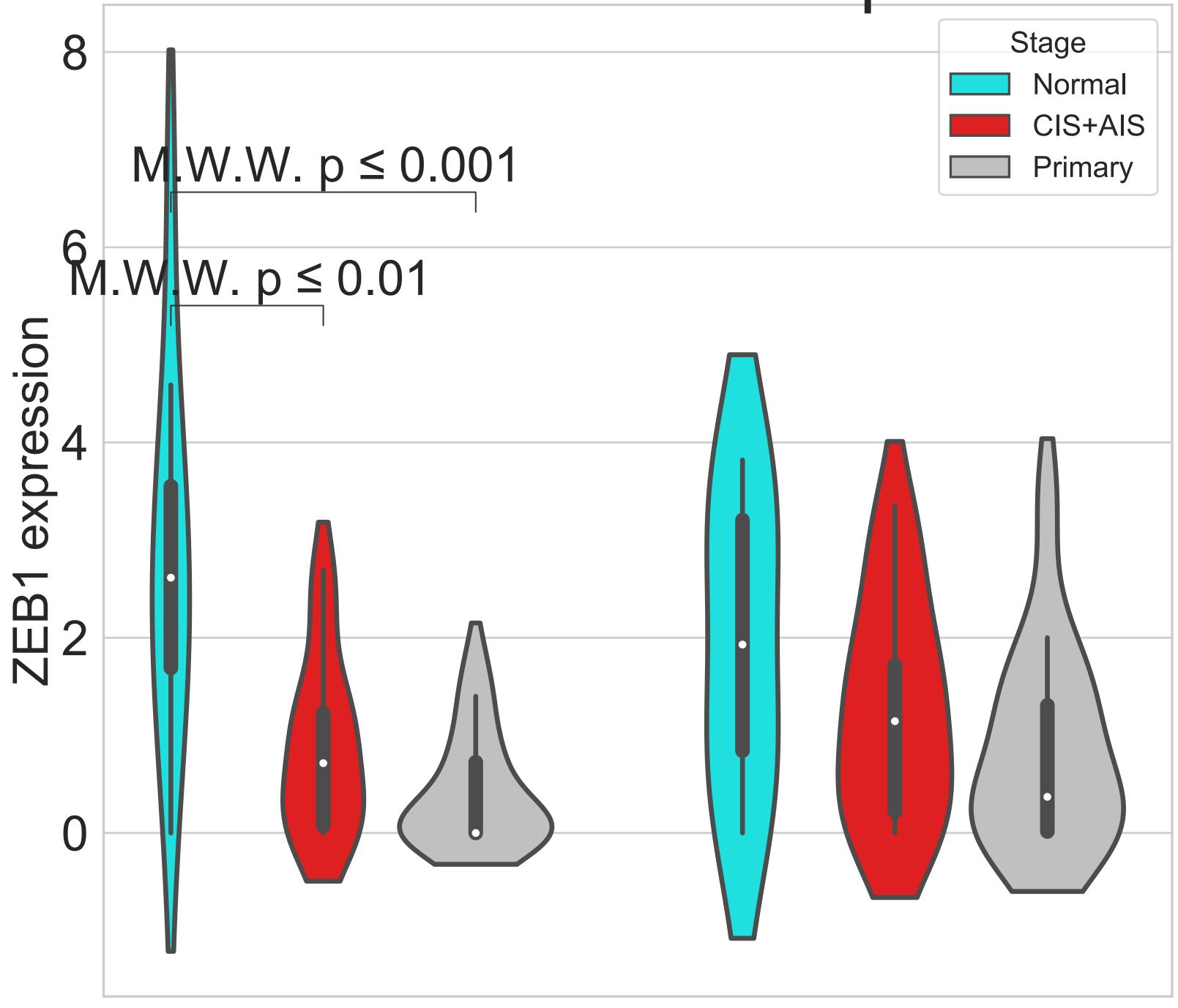
ZEB1: Kruskal-Wallis p=0.001



Lower
Higher
Mutation Shared Count per TMB