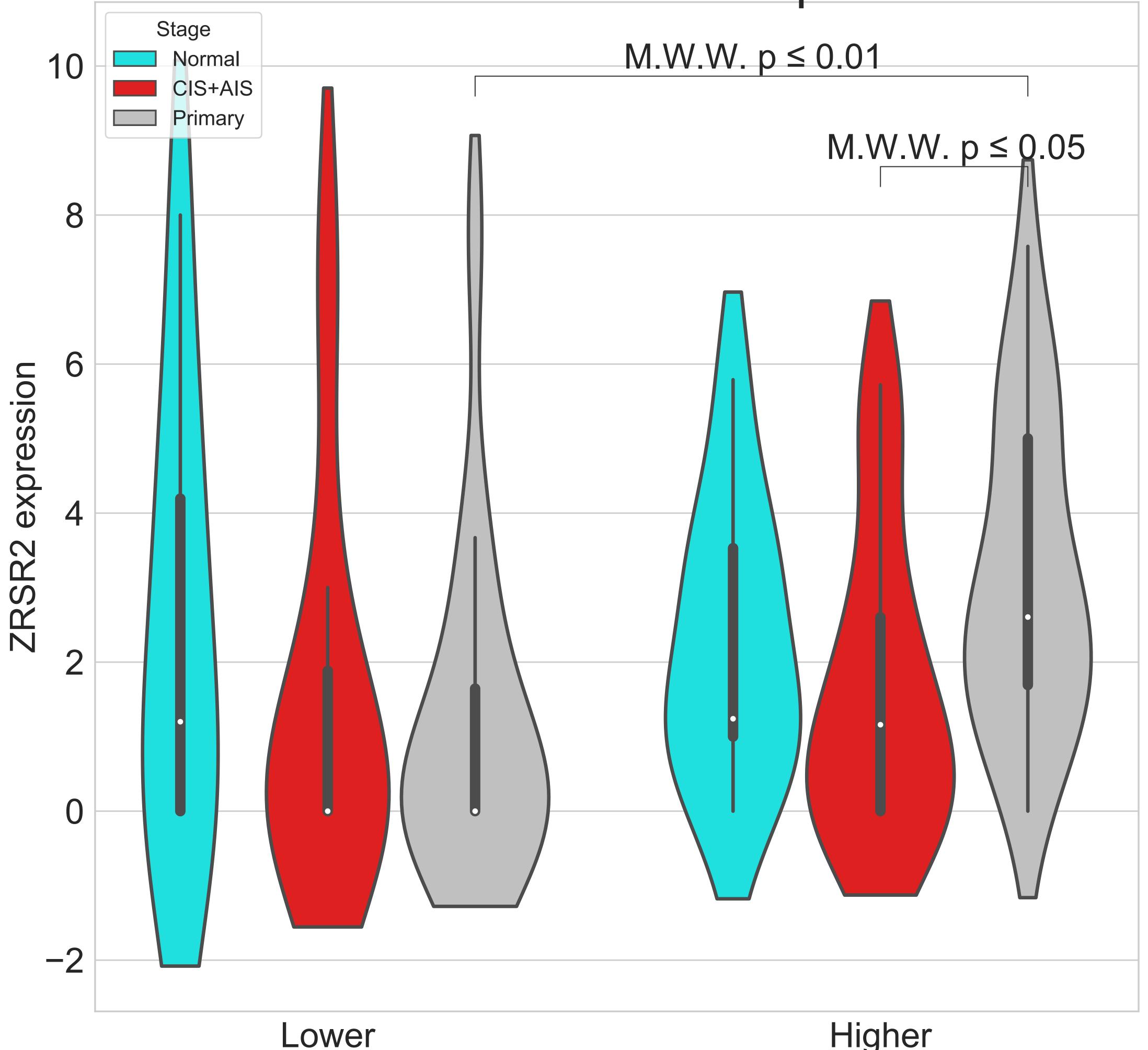
ZRSR2: Kruskal-Wallis p=0.024



wer
Mutation Shared Count per TMB