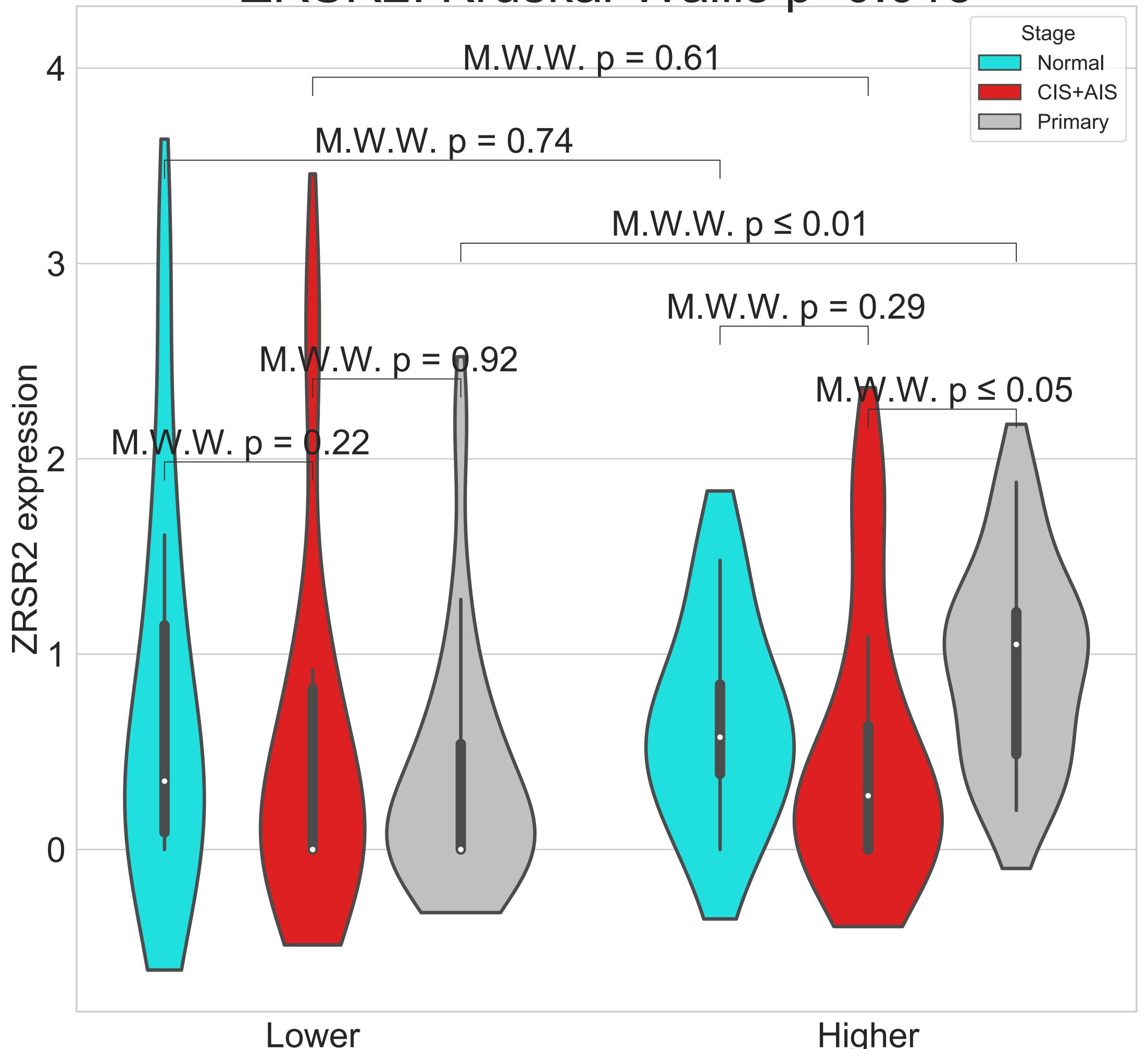
ZRSR2: Kruskal-Wallis p=0.015



ver
Mutation Shared Count per TMB