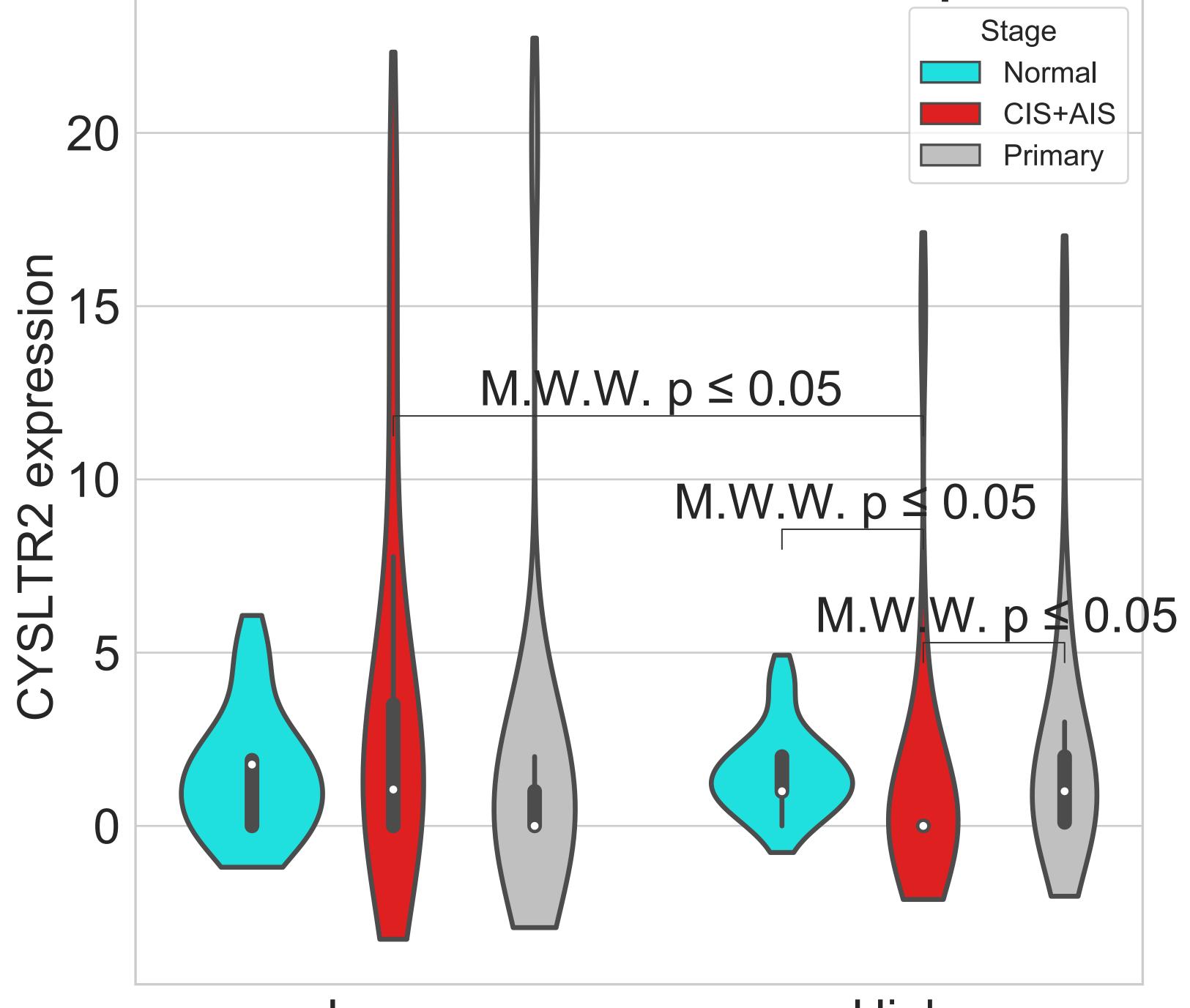
CYSLTR2: Kruskal-Wallis p=0.049



Lower Higher Mutation Shared Count