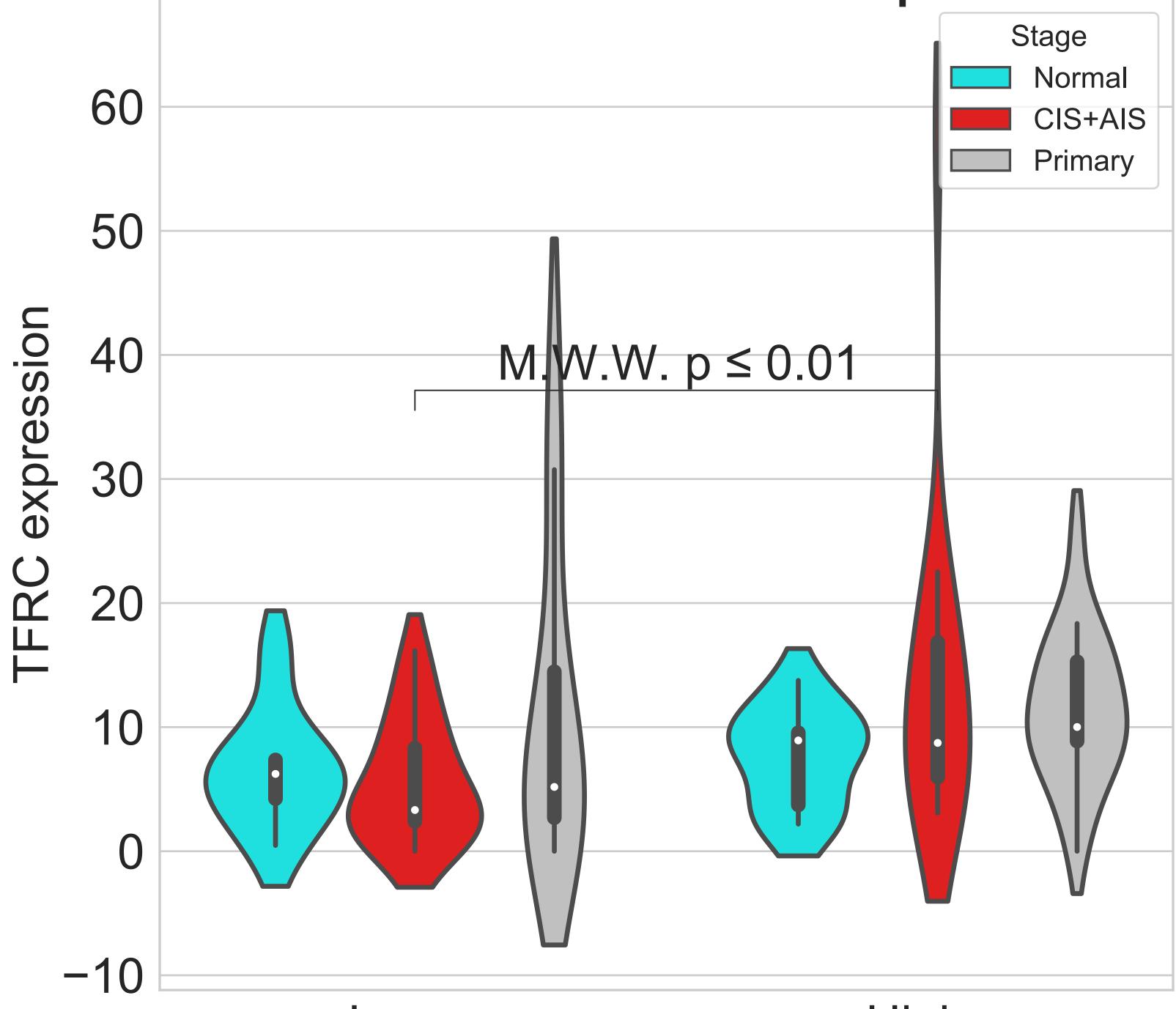
TFRC: Kruskal-Wallis p=0.041



Lower Higher Mutation Shared Count per TMB