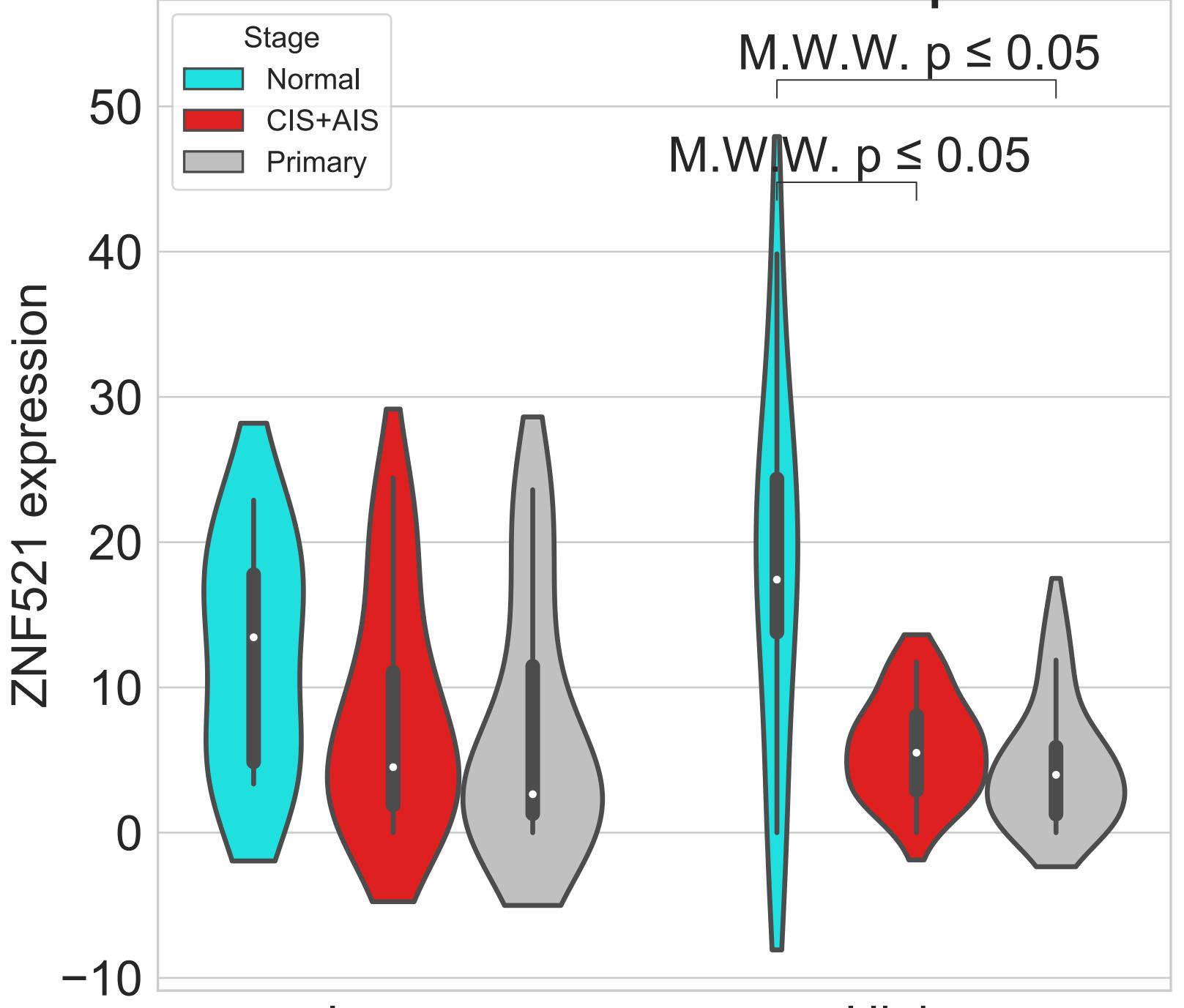
ZNF521: Kruskal-Wallis p=0.045



Lower Higher Mutation Shared Count (SYN) per TMB