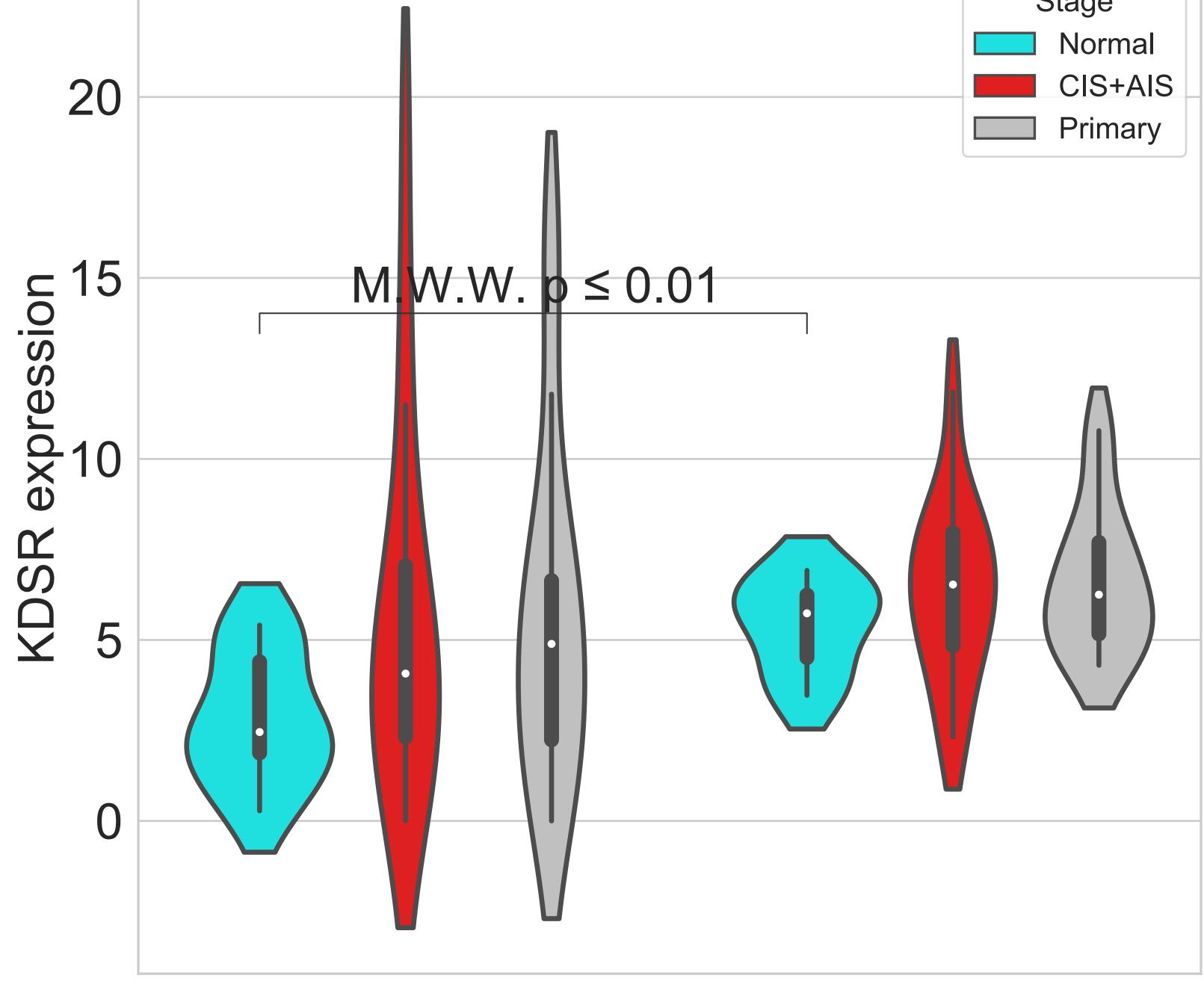
KDSR: Kruskal-Wallis p=0.012

Stage
Normal
CIS+AIS
Primary



Lower Higher Mutation Shared Count (SYN) per TMB