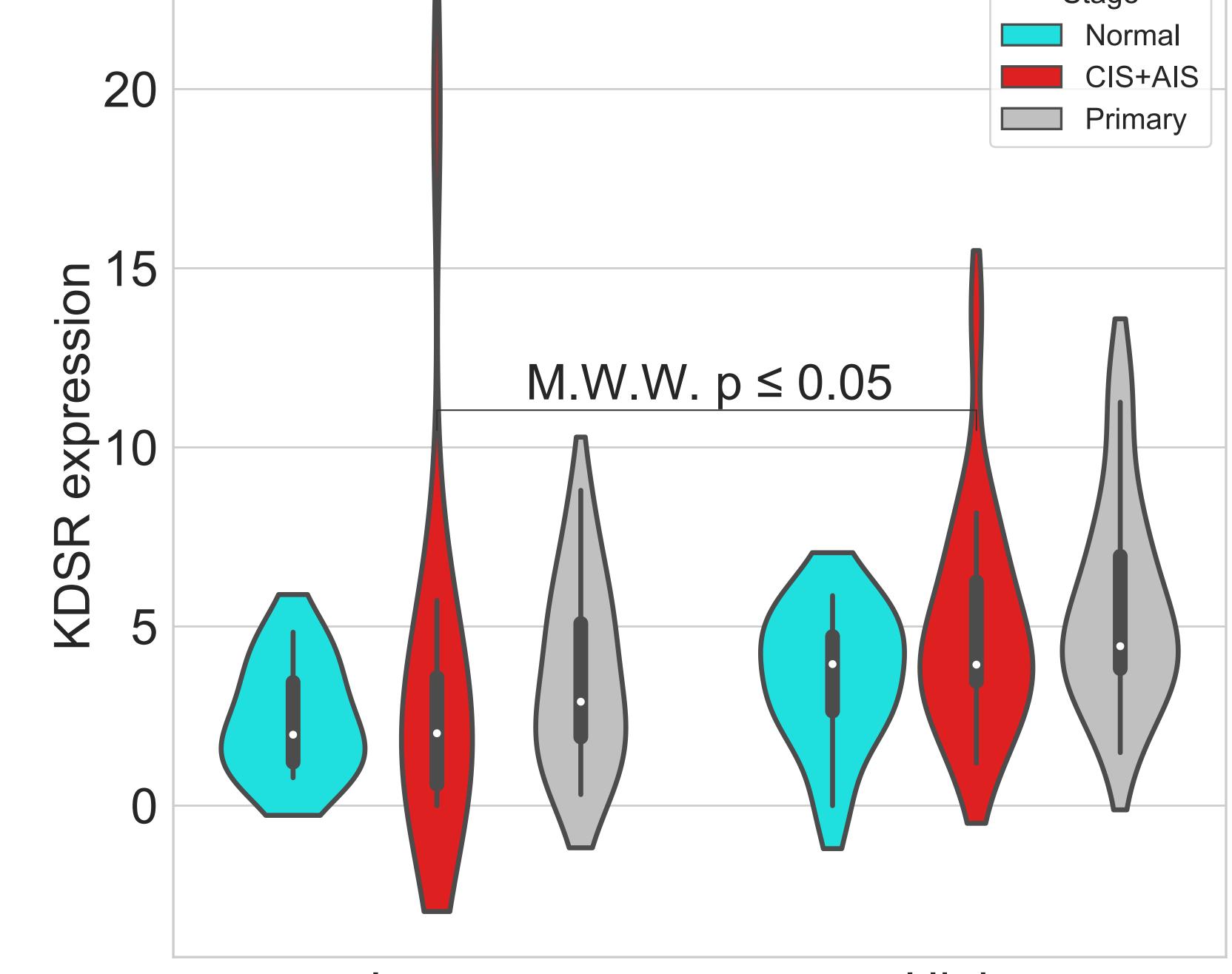
KDSR: Kruskal-Wallis p=0.012 Stage Normal CIS+AIS Primary



Higher Lower Mutation Shared Count per TMB