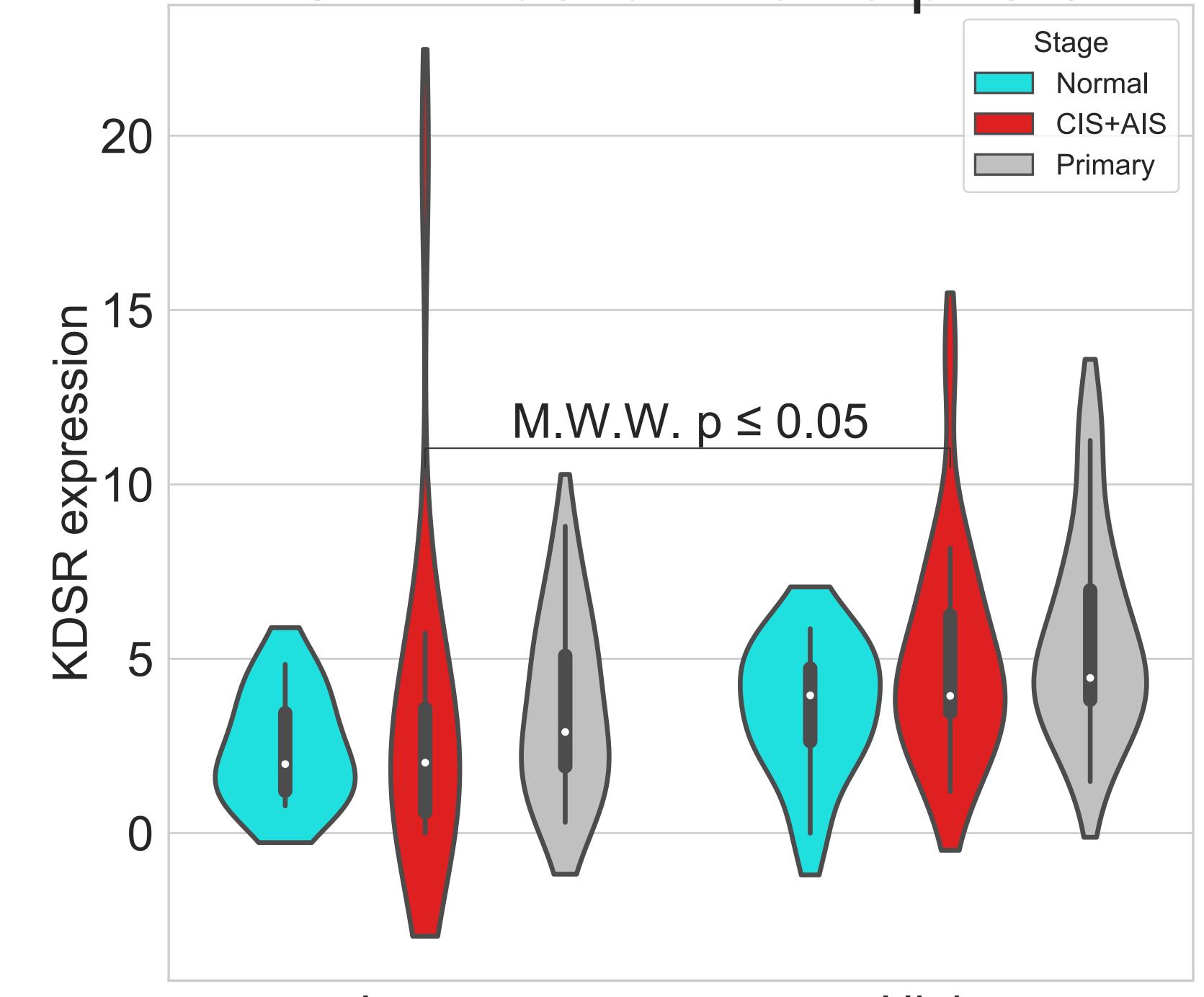
KDSR: Kruskal-Wallis p=0.012



Lower
Higher
Mutation Shared Proportion