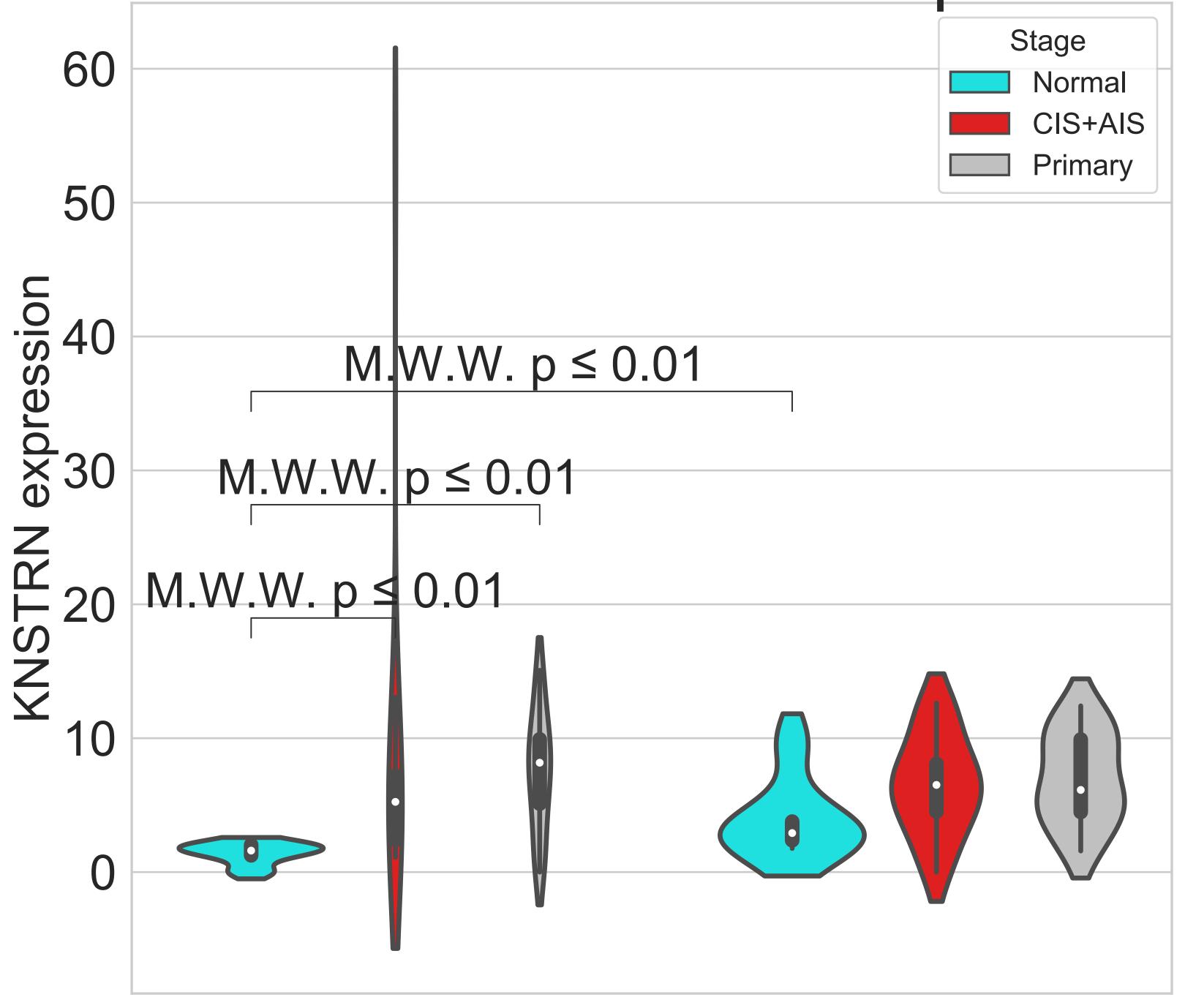
KNSTRN: Kruskal-Wallis p=0.002



Lower

Mutation Shared Proportion