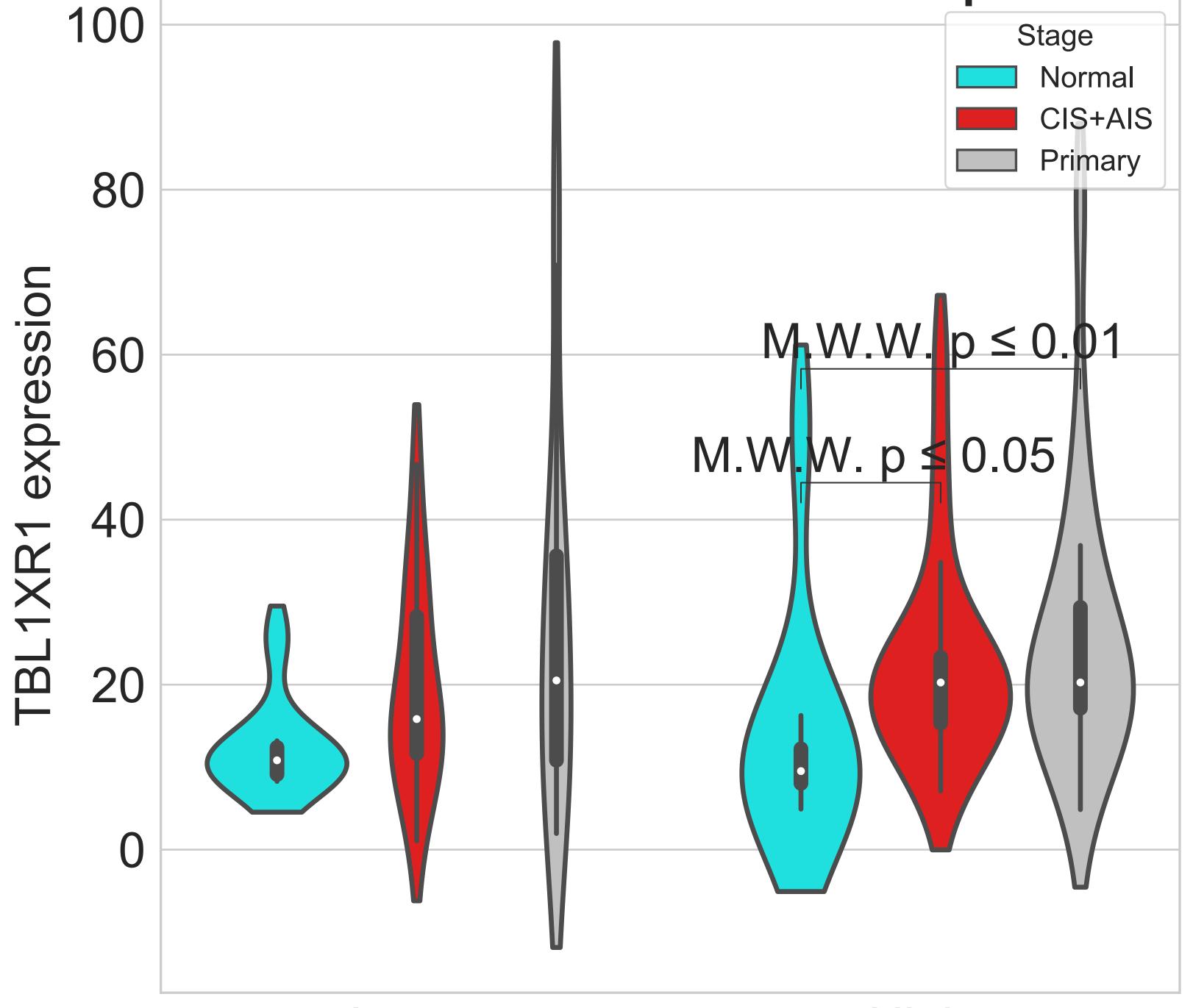
TBL1XR1: Kruskal-Wallis p=0.020



Lower Higher Mutation Shared Proportion (Union)