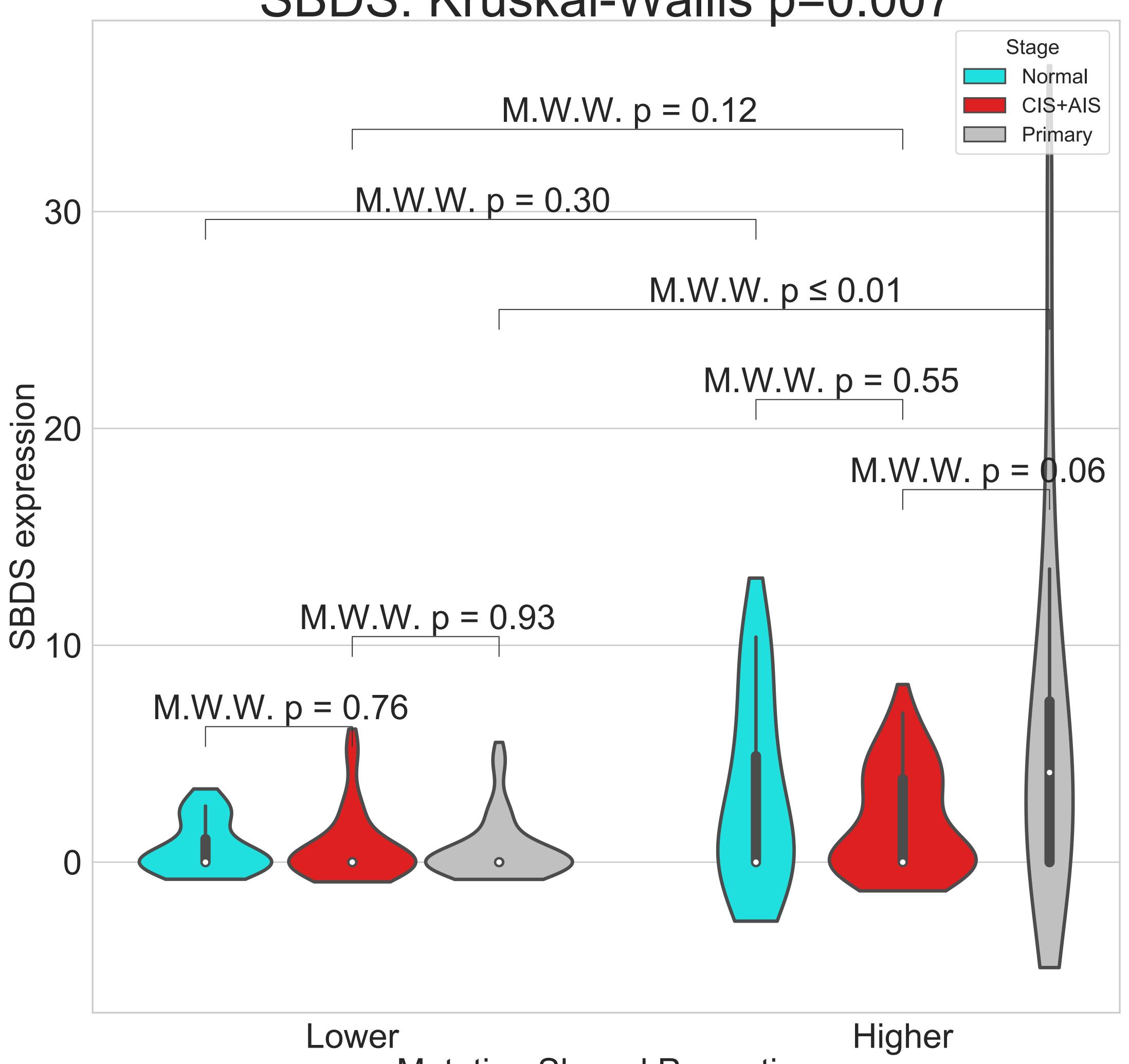
SBDS: Kruskal-Wallis p=0.007



Mutation Shared Proportion