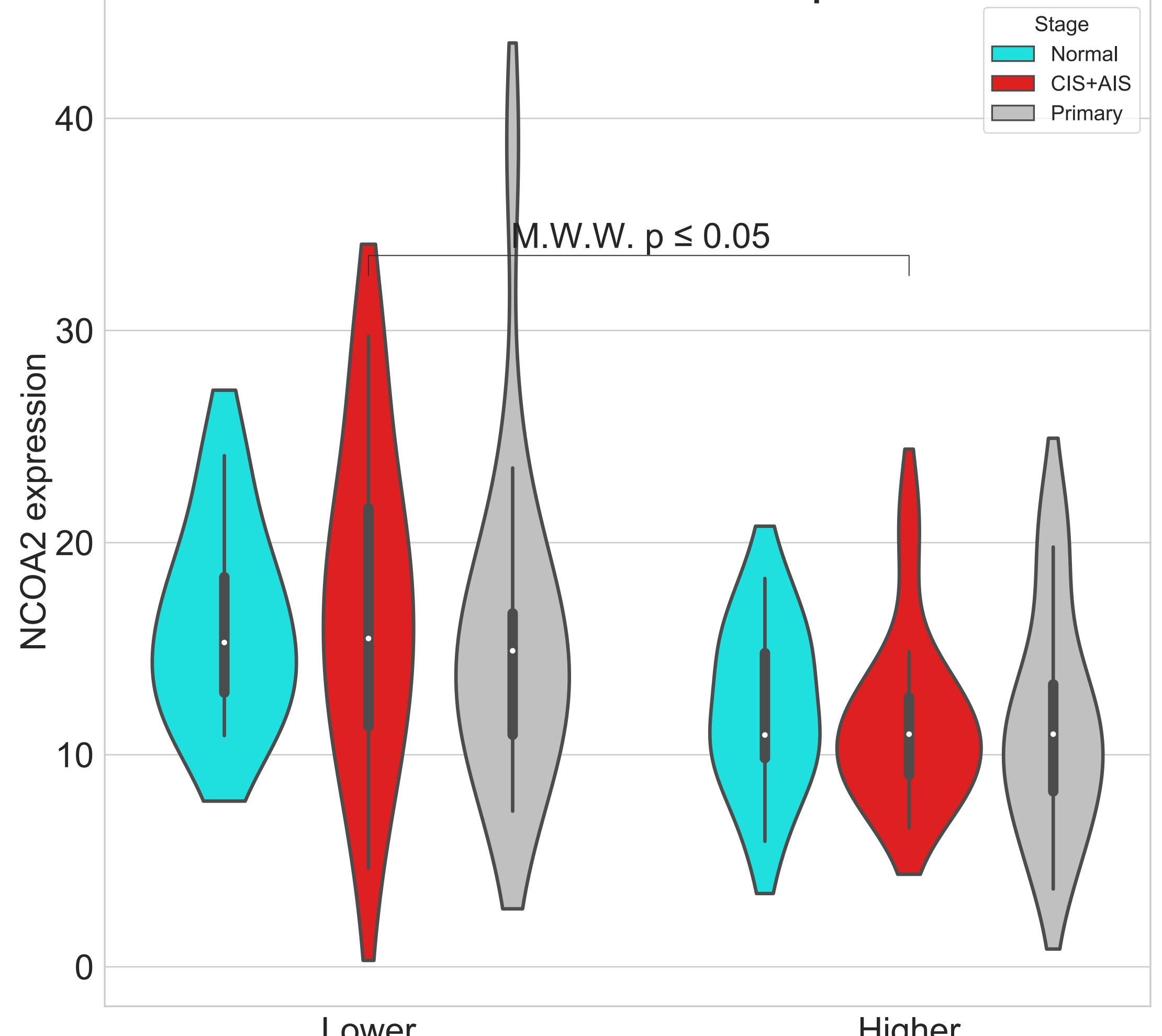
NCOA2: Kruskal-Wallis p=0.018



Higher Lower Mutation Shared Proportion (SYN)