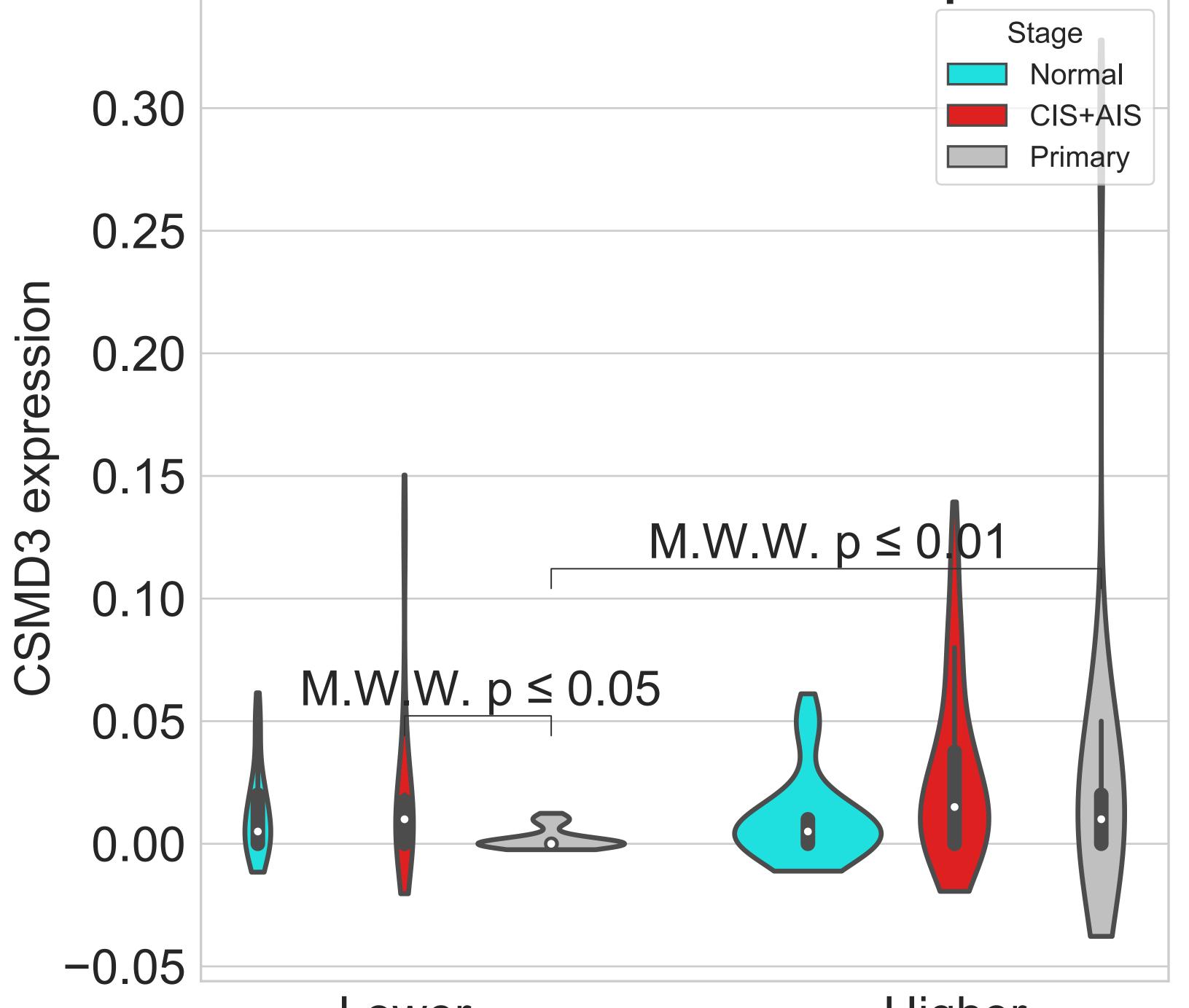
CSMD3: Kruskal-Wallis p=0.039



Lower
Mutation Shared Proportion (Union & SYN)