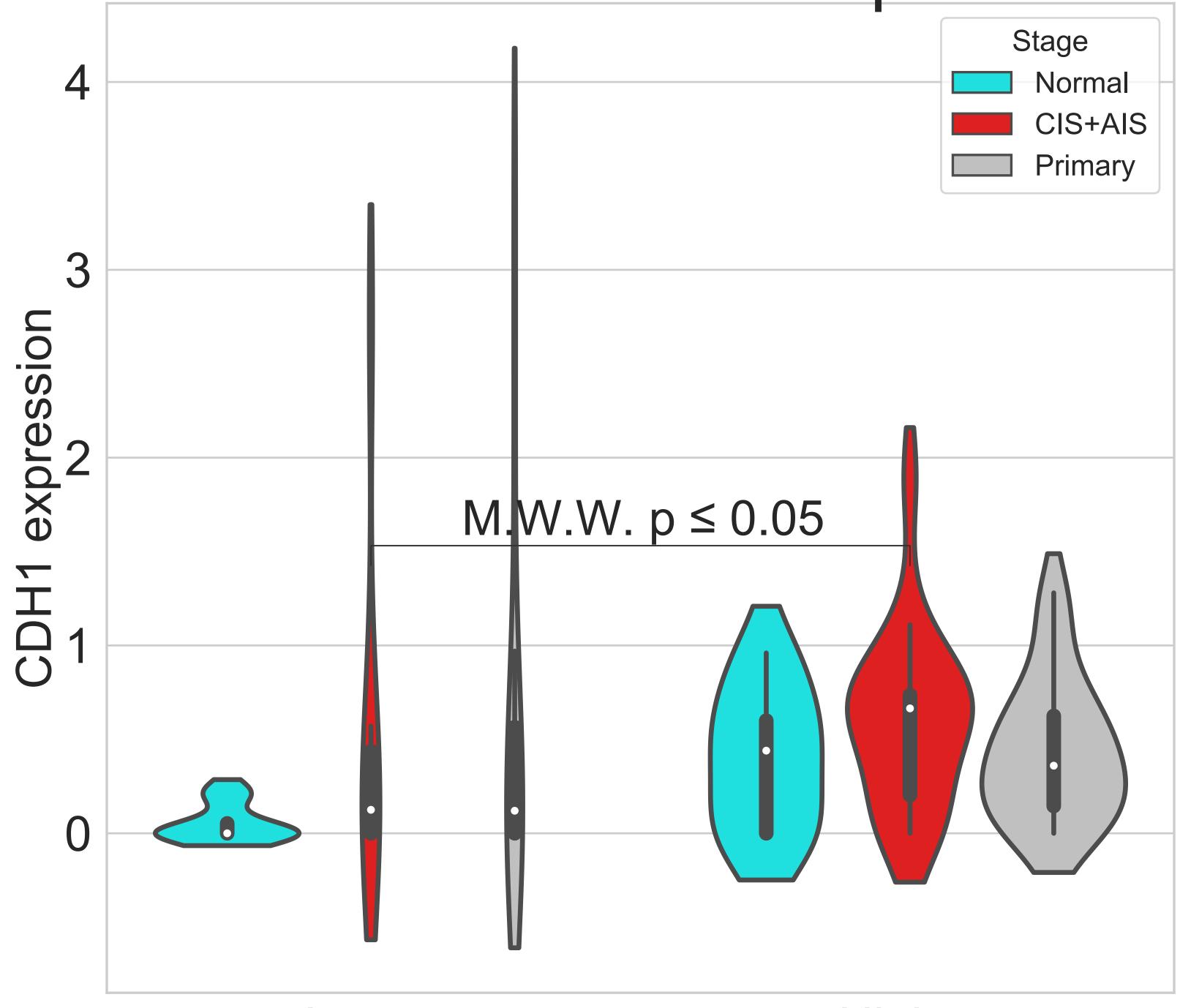
CDH1: Kruskal-Wallis p=0.018



Lower Higher Mutation Shared Proportion (Union & SYN)