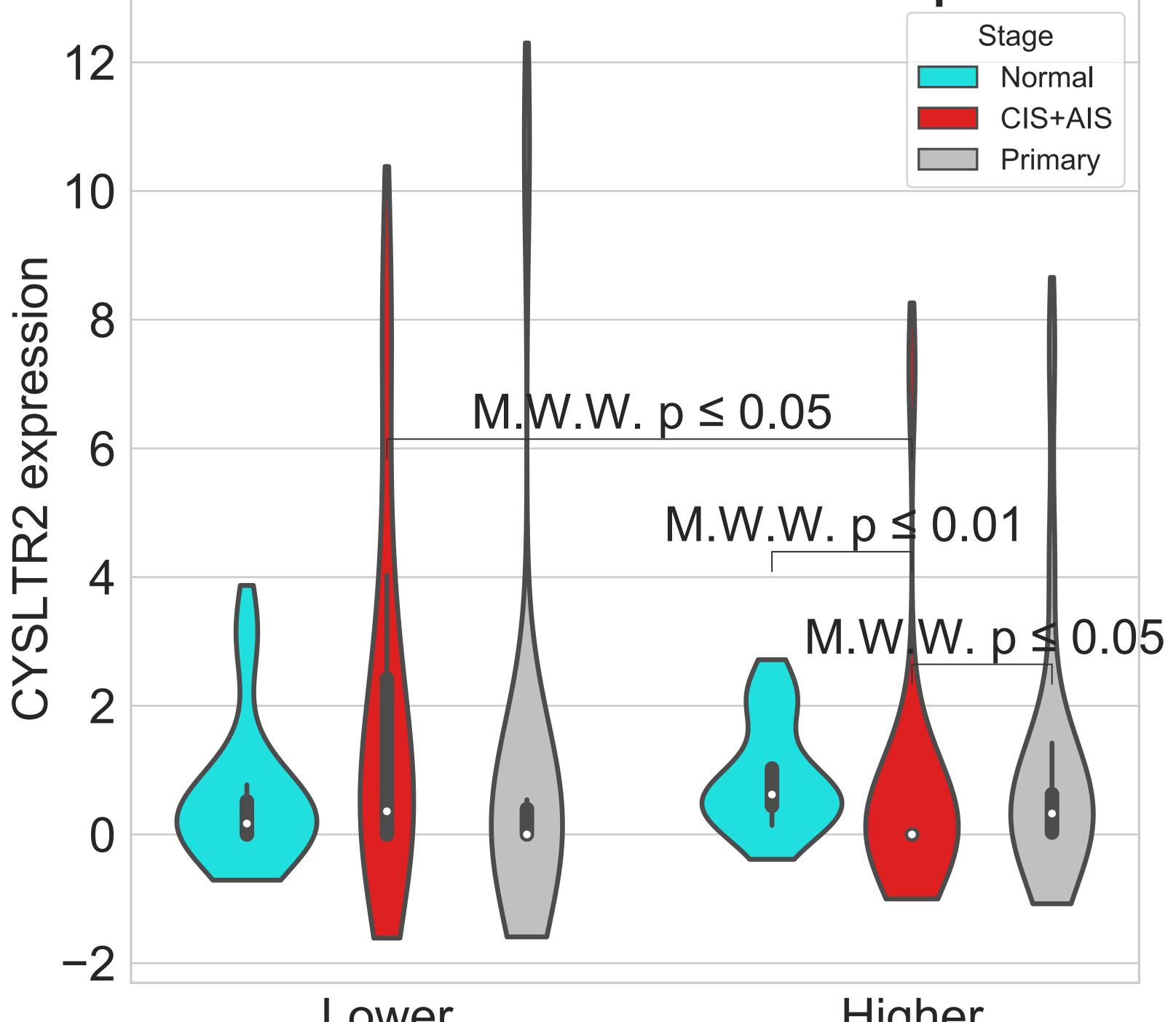
CYSLTR2: Kruskal-Wallis p=0.025



Lower
Mutation Shared Count (SYN)