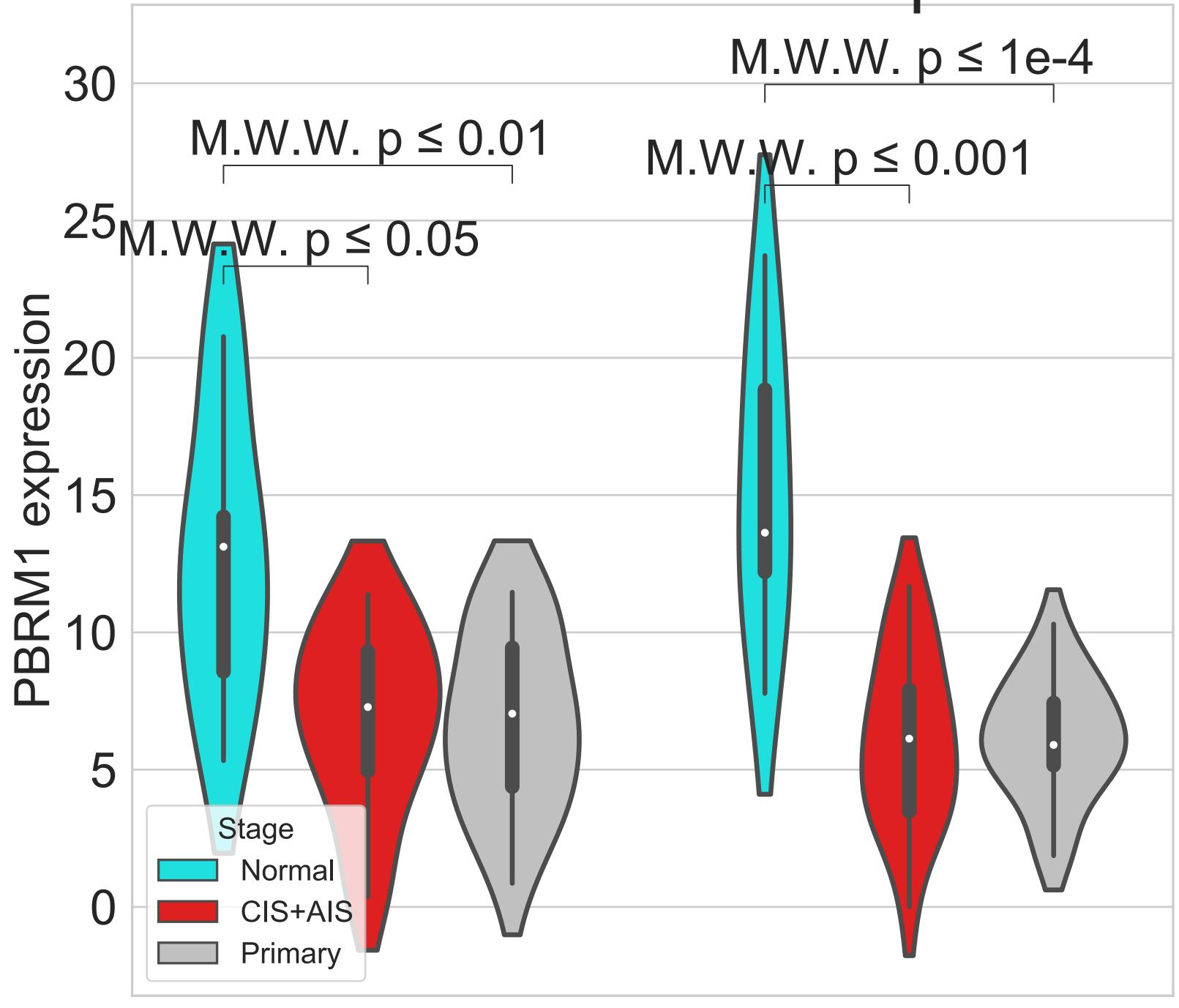
PBRM1: Kruskal-Wallis p=0.000



Lower
Higher
Mutation Shared Count (SYN)