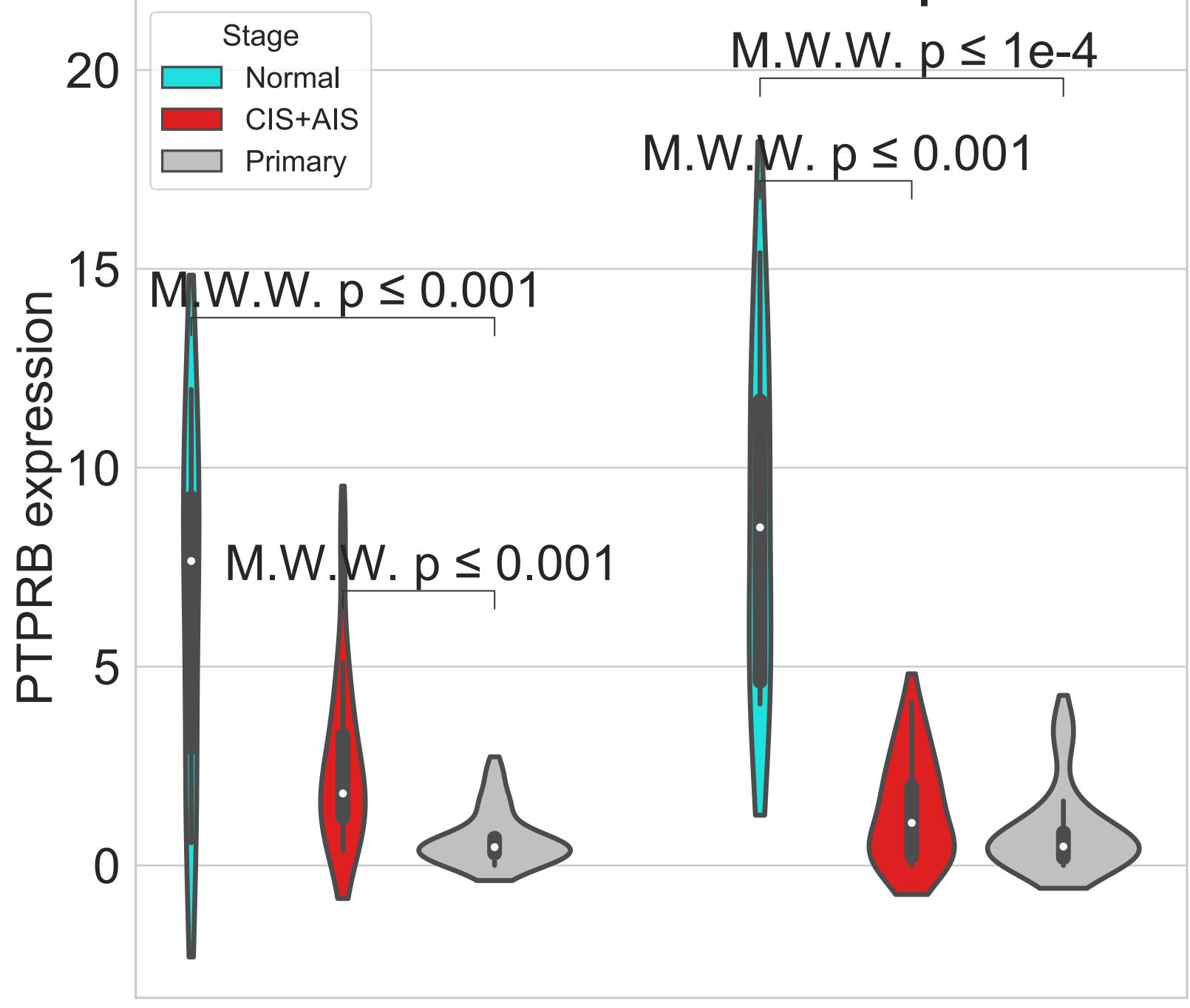
PTPRB: Kruskal-Wallis p=0.000



Lower

Higher

Mutation Shared Count (SYN)