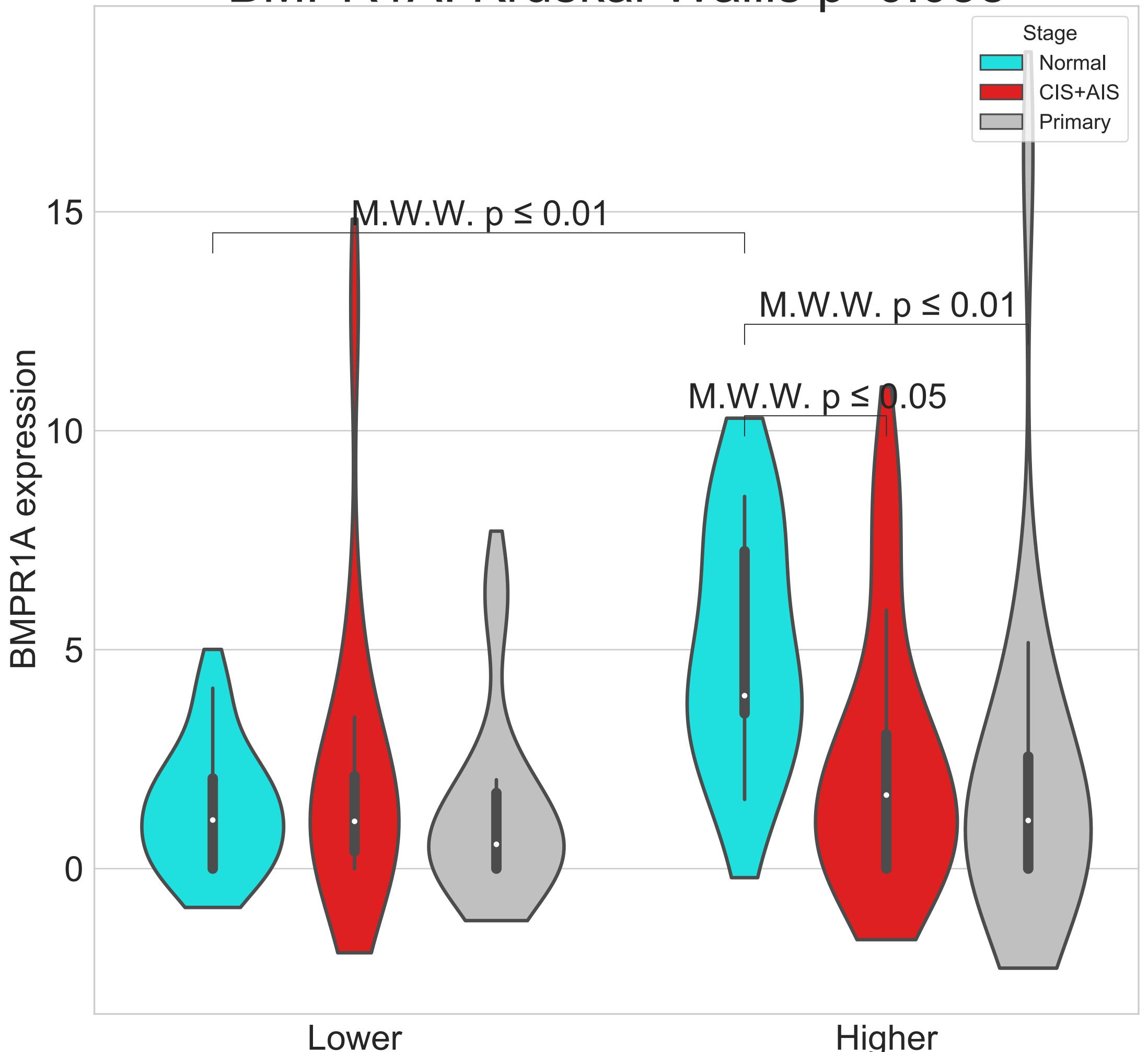
BMPR1A: Kruskal-Wallis p=0.038



ver
Mutation Shared Count (SYN)