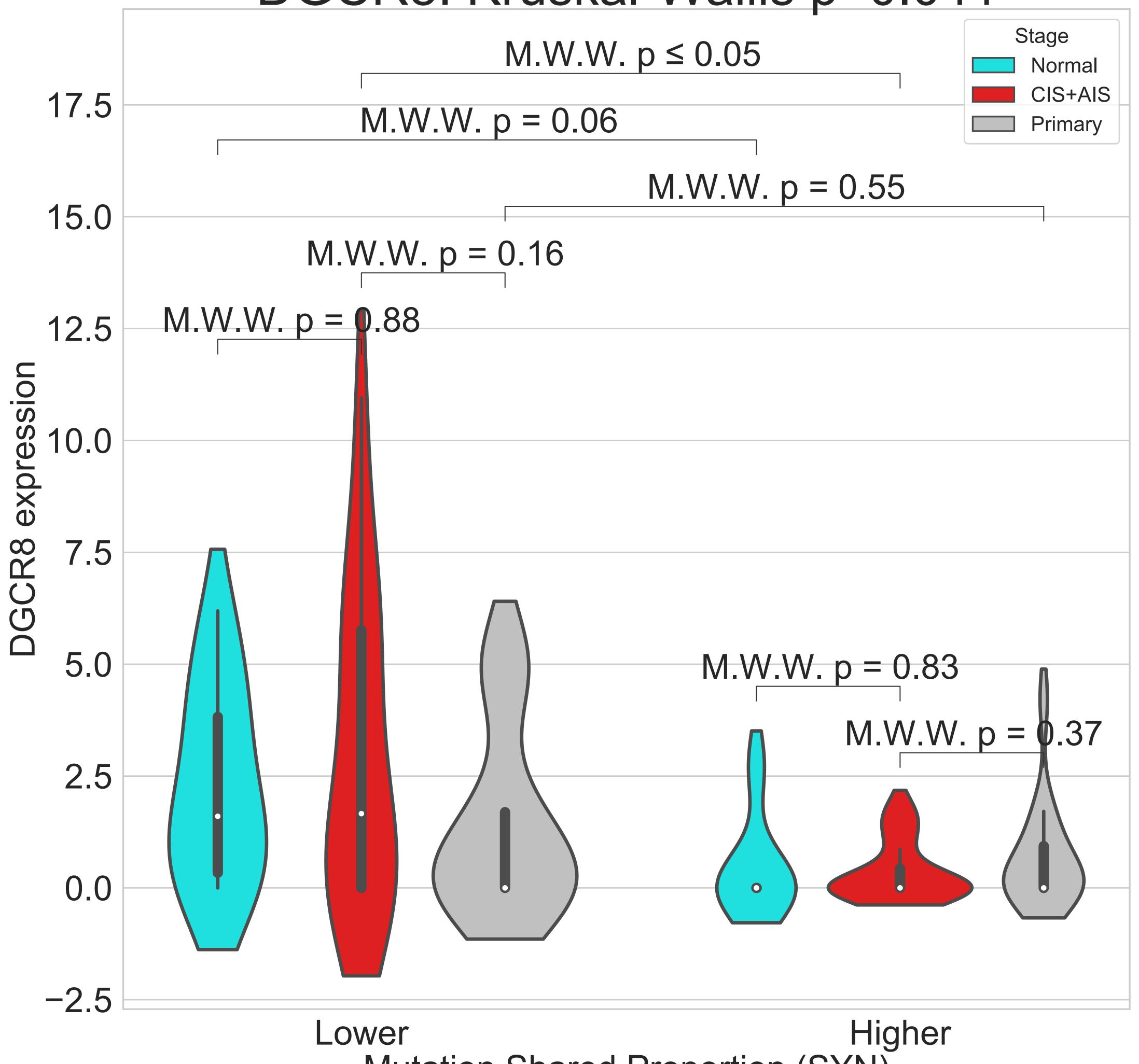
DGCR8: Kruskal-Wallis p=0.041



Mutation Shared Proportion (SYN)