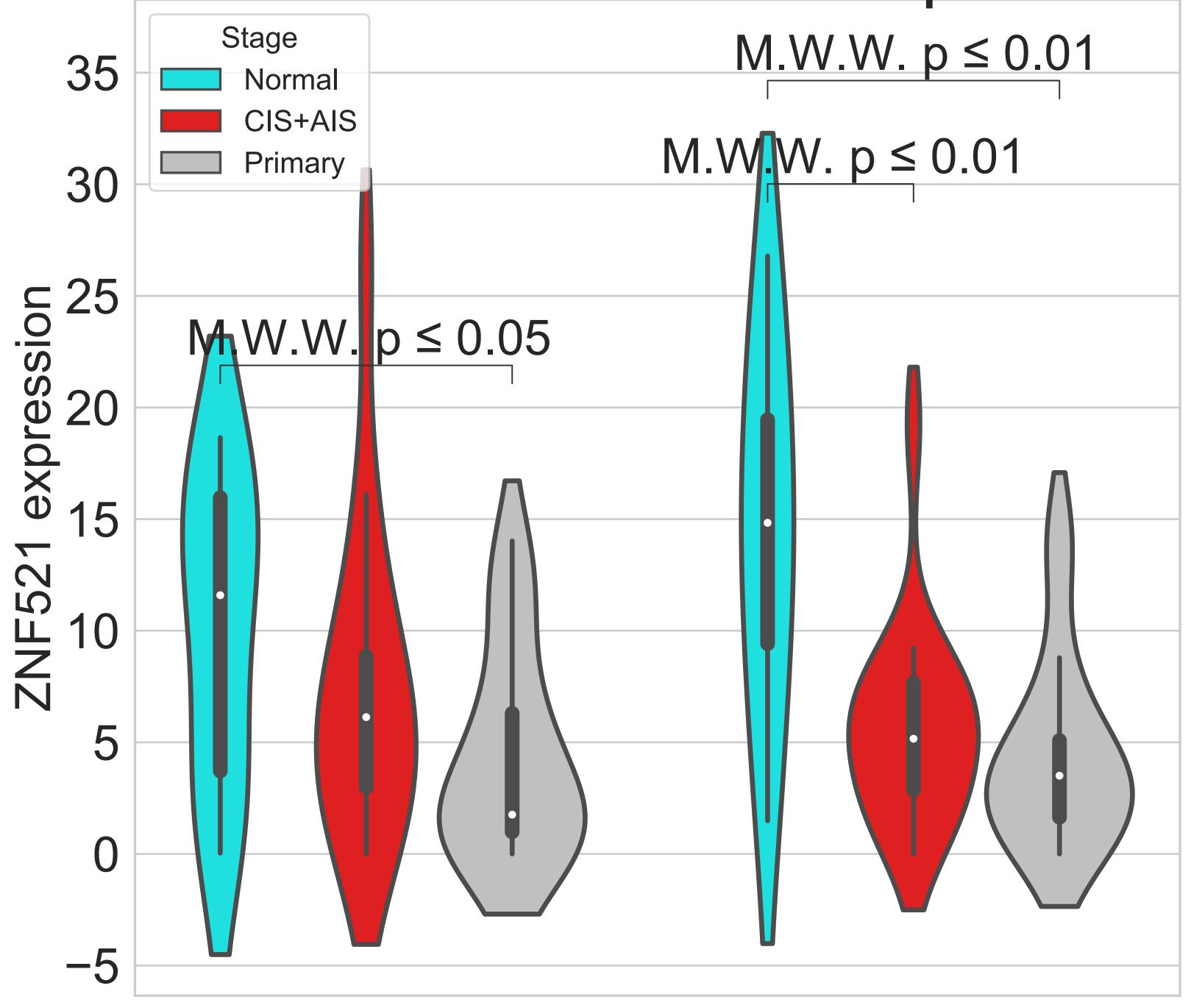
ZNF521: Kruskal-Wallis p=0.013



Lower
Higher
Mutation Shared Count (SYN)