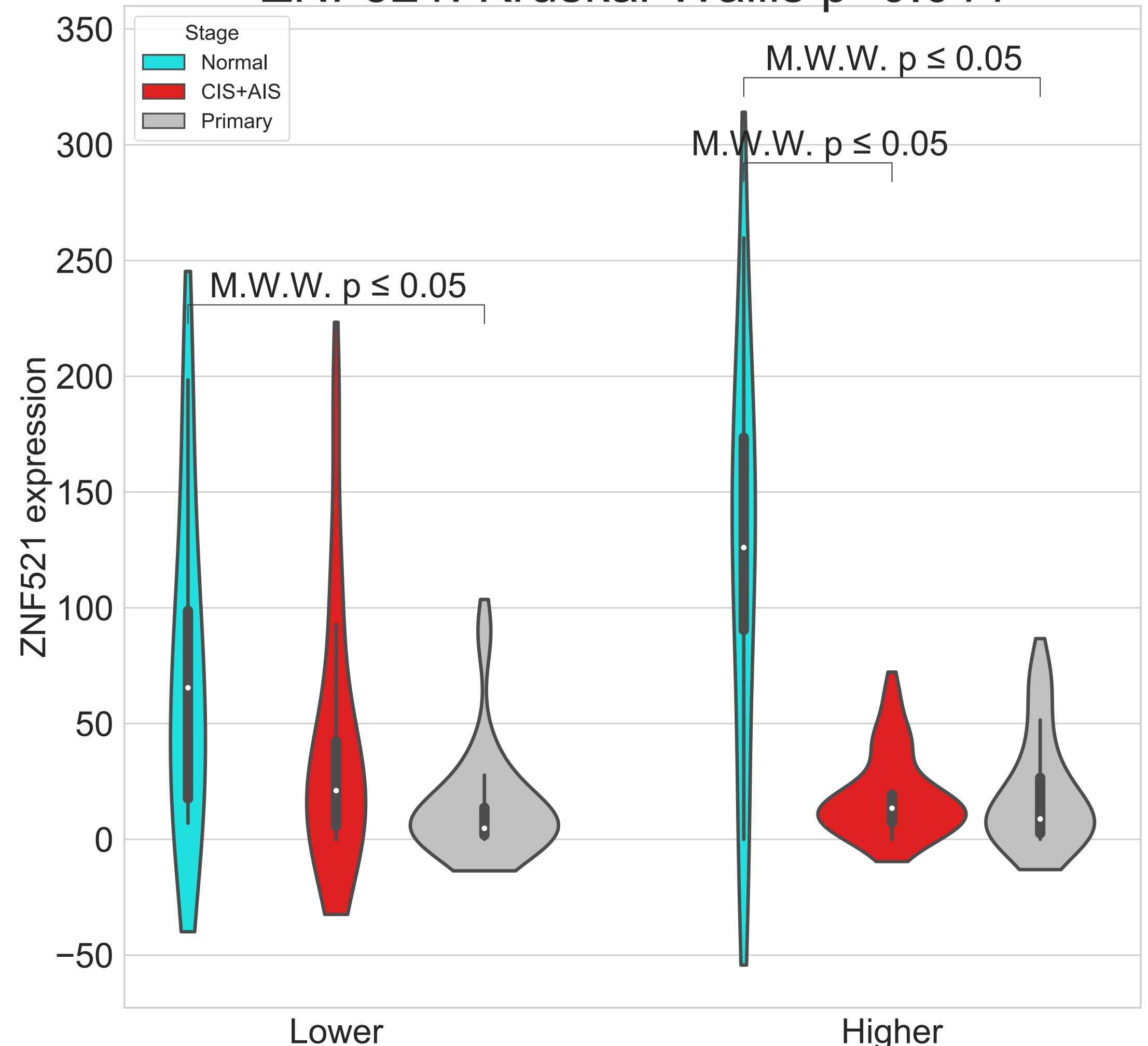
ZNF521: Kruskal-Wallis p=0.011



Higher Mutation Shared Proportion