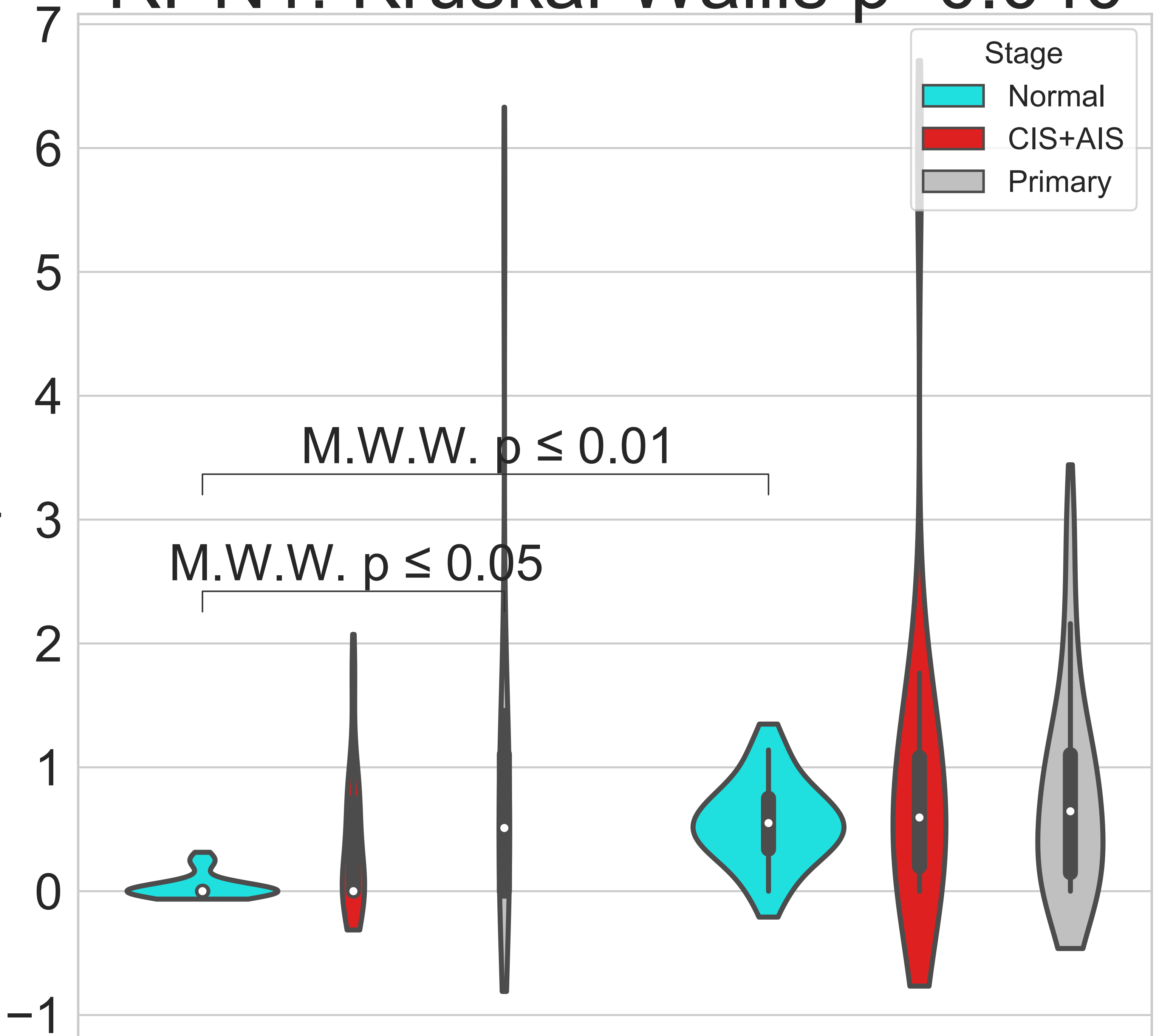


RPN1: Kruskal-Wallis p=0.040

RPN1 expression



Lower Higher
Mutation Shared Count per TMB