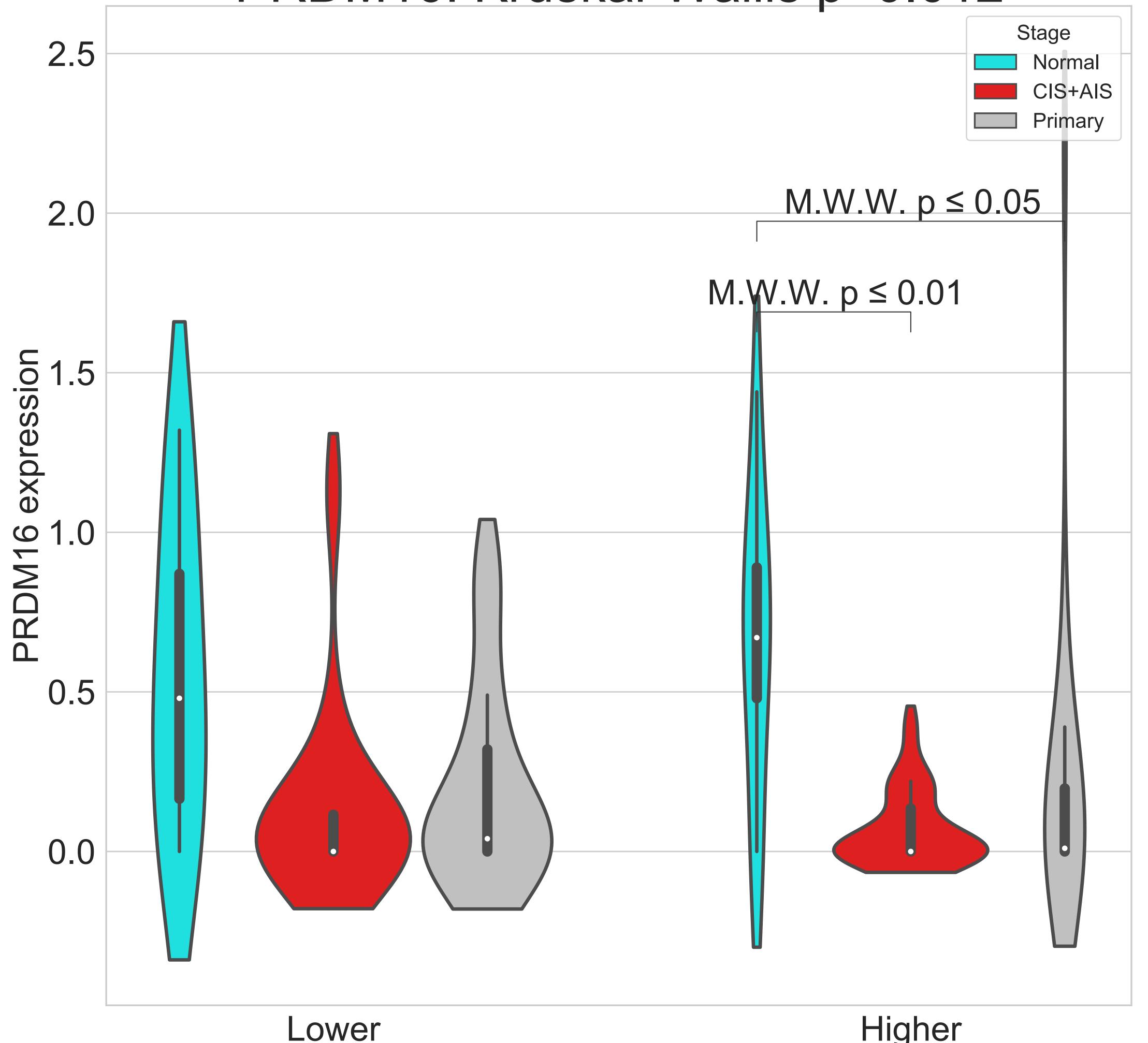
PRDM16: Kruskal-Wallis p=0.012



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Mutation Shared Count per TMB