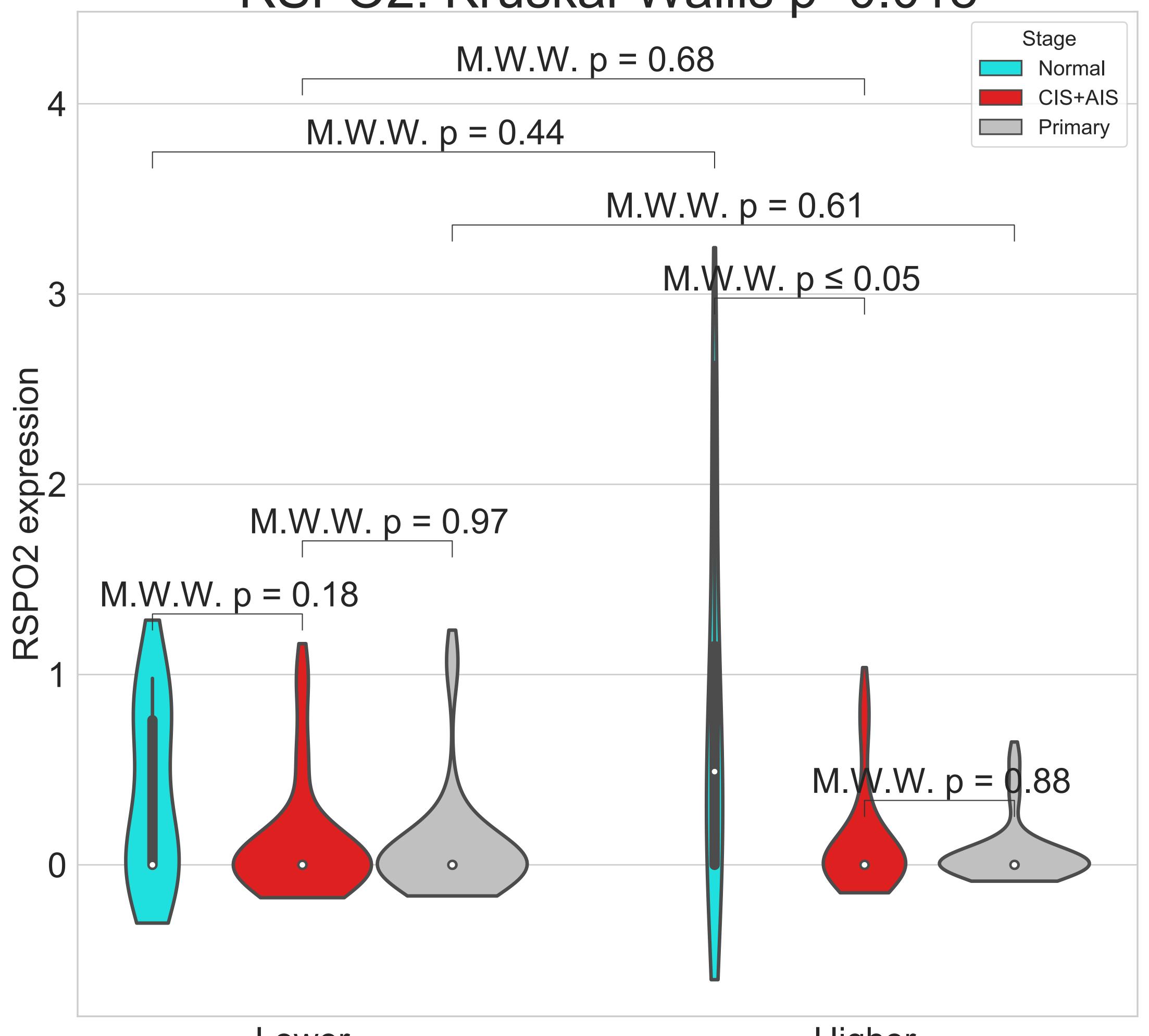
RSP02: Kruskal-Wallis p=0.018



Lower

Mutation Shared Count per TMB