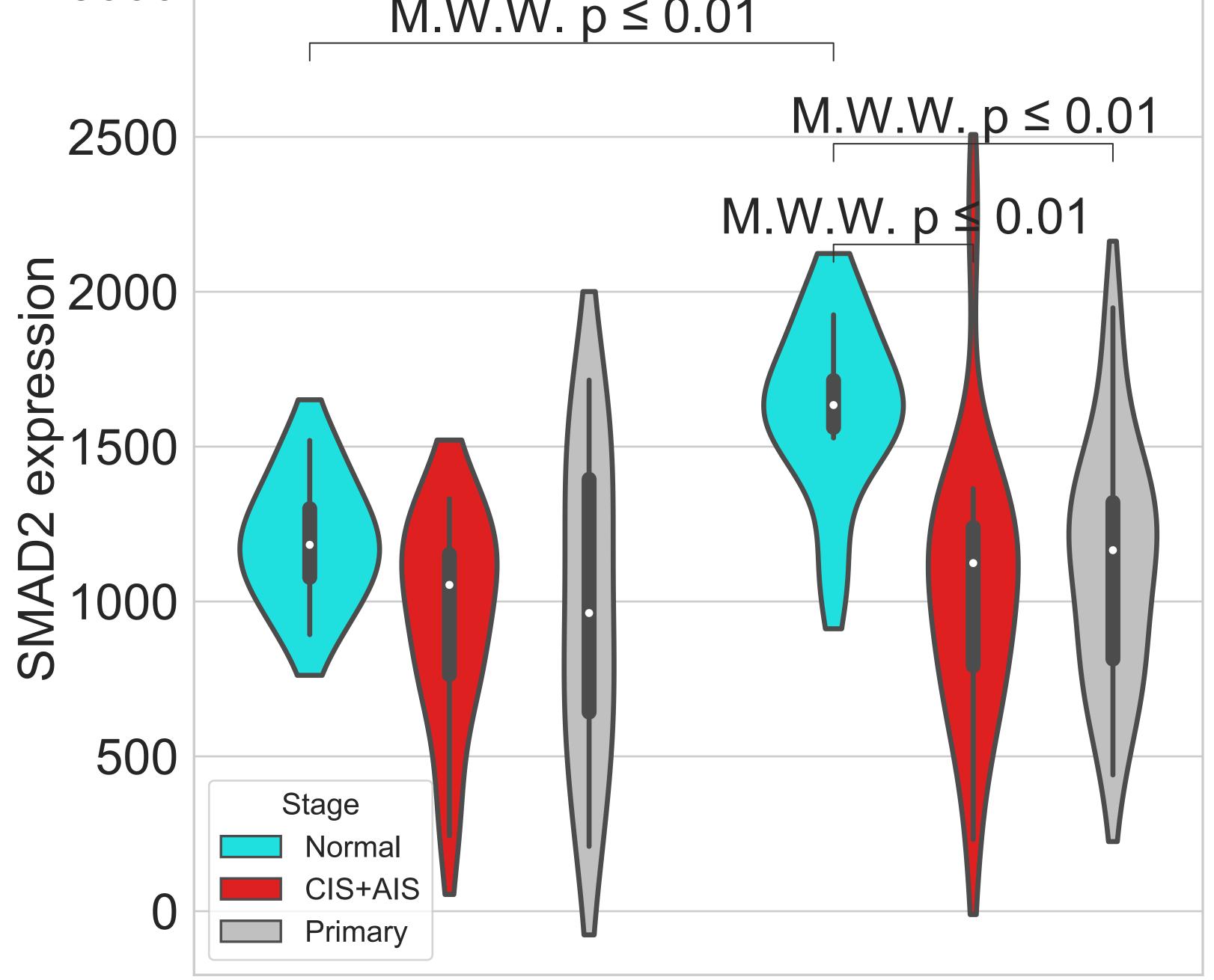
SMAD2: Kruskal-Wallis p=0.010

M.W.W. p ≤ 0.01

M.W.W. p ≤ 0.01



Lower Higher Mutation Shared Proportion (Union & SYN)