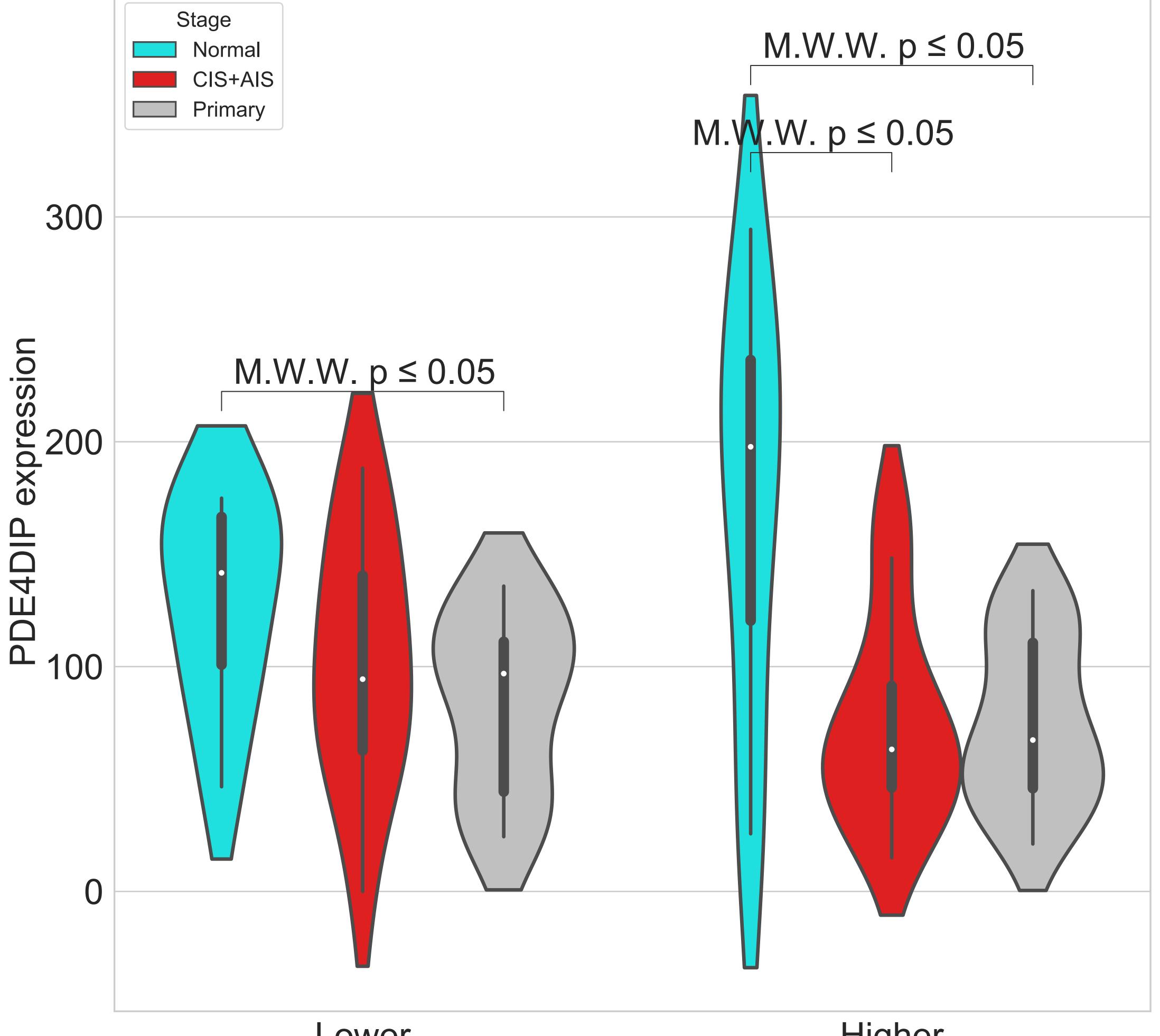
PDE4DIP: Kruskal-Wallis p=0.015

e mal 
S+AIS

M.W.W. p ≤ 0.05



Lower
Mutation Shared Count (SYN) per TMB