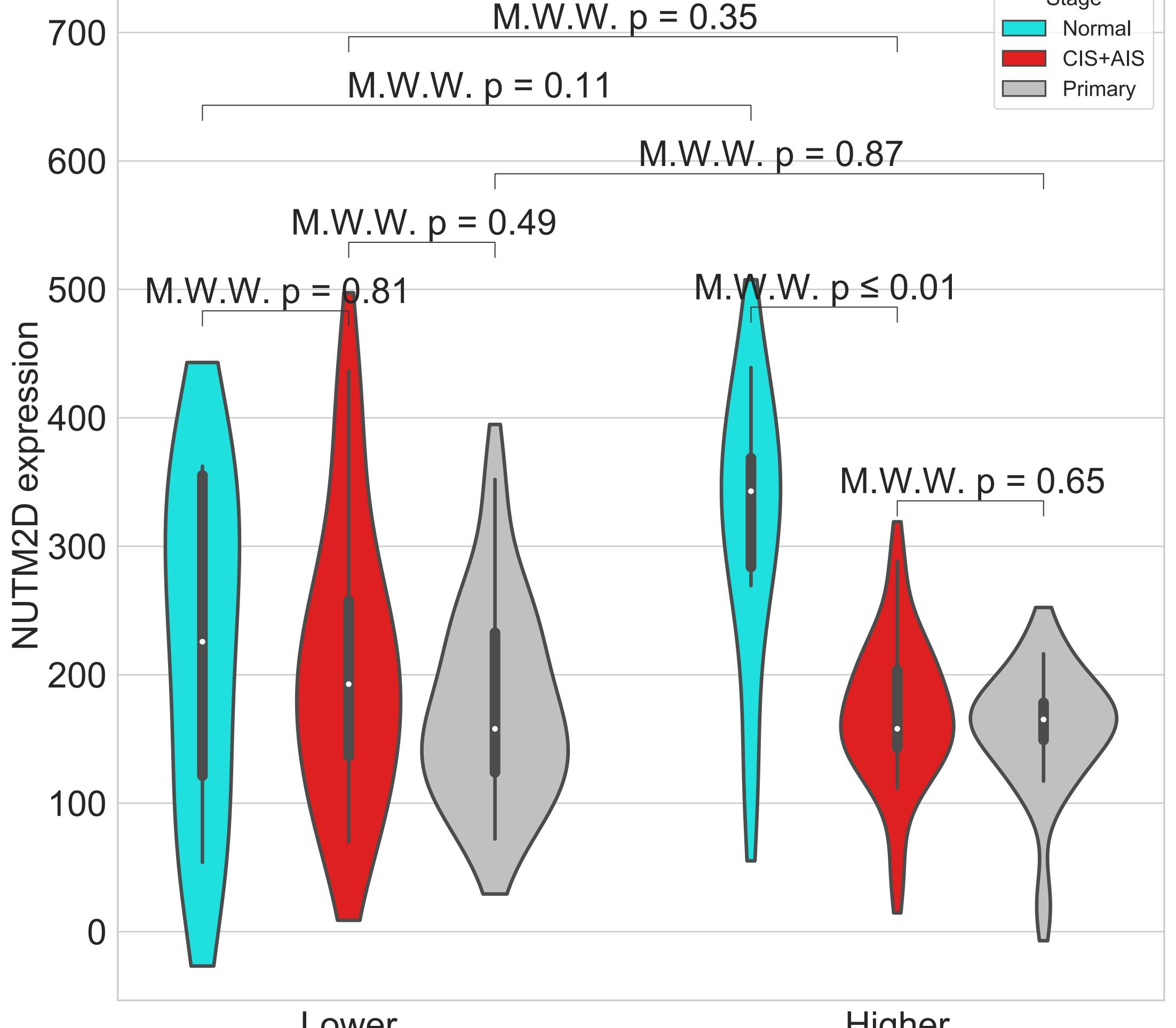
NUTM2D: Kruskal-Wallis p=0.039 Stage M.W.W. p = 0.35Normal CIS+AIS M.W. p = 0.11Primary M.W. p = 0.87M.W.W. p = 0.49M.W.W. p ≤ 0.01 M.W.W. p = 0.65



Higher Lower Mutation Shared Count (SYN)