HRAS: Kruskal-Wallis p=0.000 350 Stage M.W.W. p ≤ 0.05Normal CIS+AIS **Primary** 300 M.W.W. p ≤ 1e-4 250 M.W.W. $p \le 0.001$ 100 50 -50 Higher Lower **Mutation Shared Count**