CTNNB1: Kruskal-Wallis p=0.002 500 M.W.W. p ≤ 0.05 Stage Normal CIS+AIS M.W.W. p ≤ 0.05 Primary 400 M.W.W. p ≤ 0.05 M.W. p **≤**

Lower Higher Mutation Shared Count per TMB