SMAD2: Kruskal-Wallis p=0.010 Stage M.W.W. p = 0.56Normal CIS+AIS M.W.W. p ≤ 0.01 Primary 3000 M.W.W. p = 0.50M.W.W. p ≤ 0.01 2500 M.W.W. p = 0.44M.W. p = 0.88M.W.W. p = 0.101000 500 Higher Lower

Mutation Shared Proportion (Union)