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 = 0.502500 M.W.W. p ≤ 0.01 M.W.W. p = 0.44M.W. p = 0.88M.W.W. p = 0.10SMAI 1000 500 Higher Lower

Mutation Shared Proportion (Union & SYN)