SMAD2: Kruskal-Wallis p=0.024 Stage M.W. p = 0.76Normal CIS+AIS M.W.W. p ≤ 0.05 Primary M.W.W. p = 0.84M.W.W. p ≤ 0.01 M.W.W. p = 0.54M.W.W. p = 0.75M.W.W. p = 0.10

3000

2500

1000

500

Lower
Mutation Shared Proportion (SYN)