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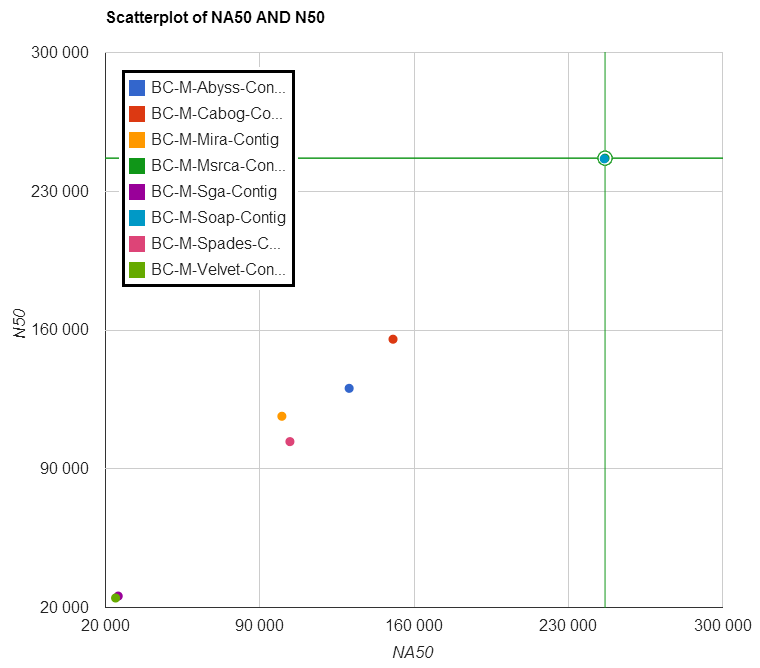


Figure S-A 1 Scatterplot of NA50 and N50 statistics on contigs  
from MiSeq assemblies on *Bacillus cereus*

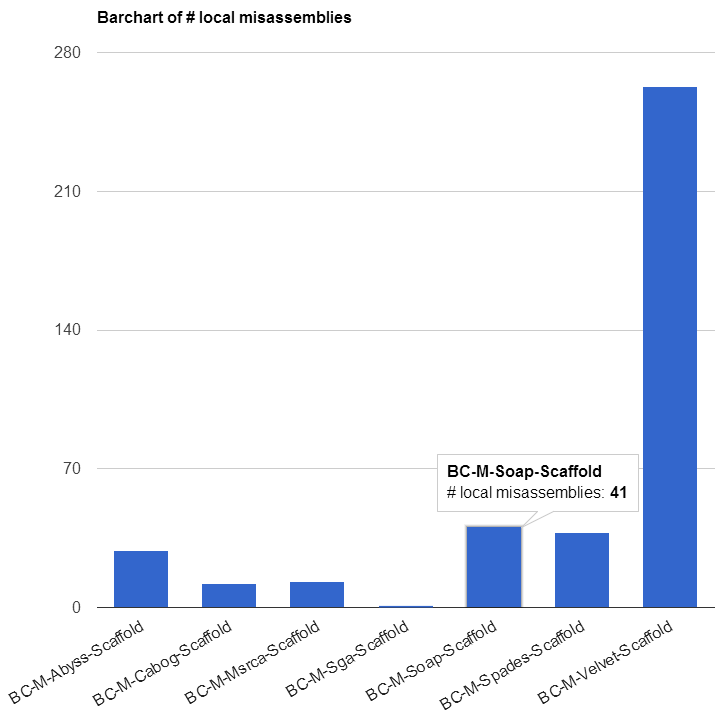


Figure S-A 2 The number of local misassemblies on scaffolds   
from MiSeq assemblies on *Bacillus cereus*

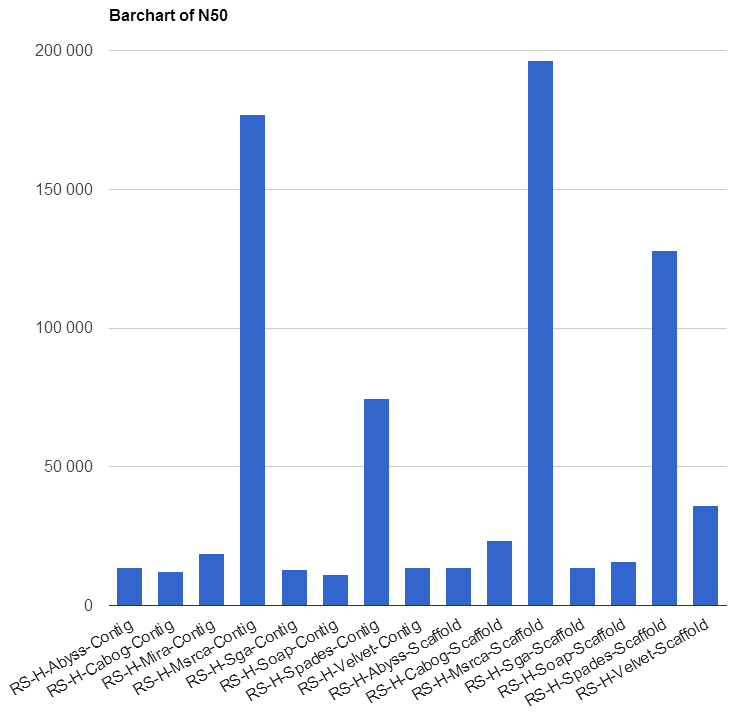


Figure S-A 3 N50 values for contigs and scaffolds from   
HiSeq assemblies on *Rhodobacter sphaeroides*

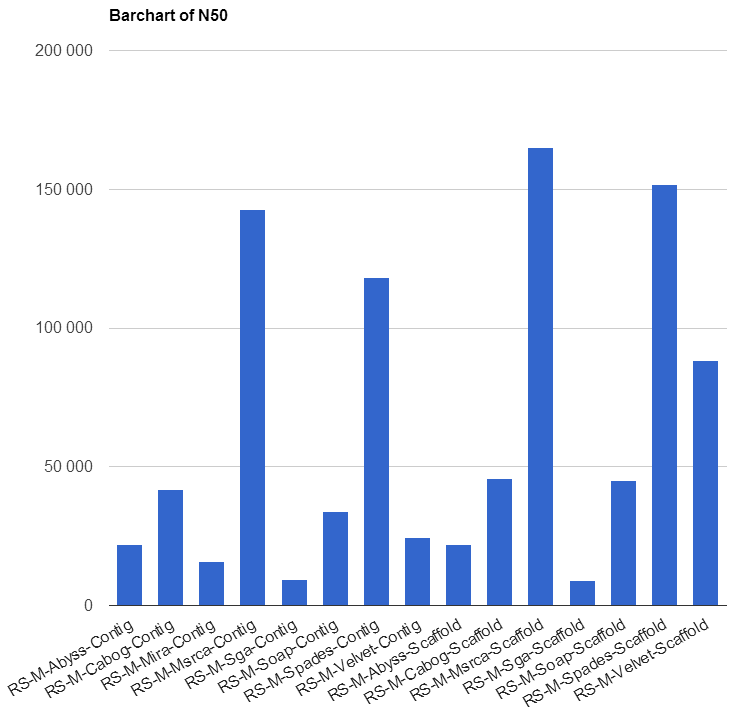


Figure S-A 4 N50 values for contigs and scaffolds from  
MiSeq assemblies on *Rhodobacter sphaeroides*

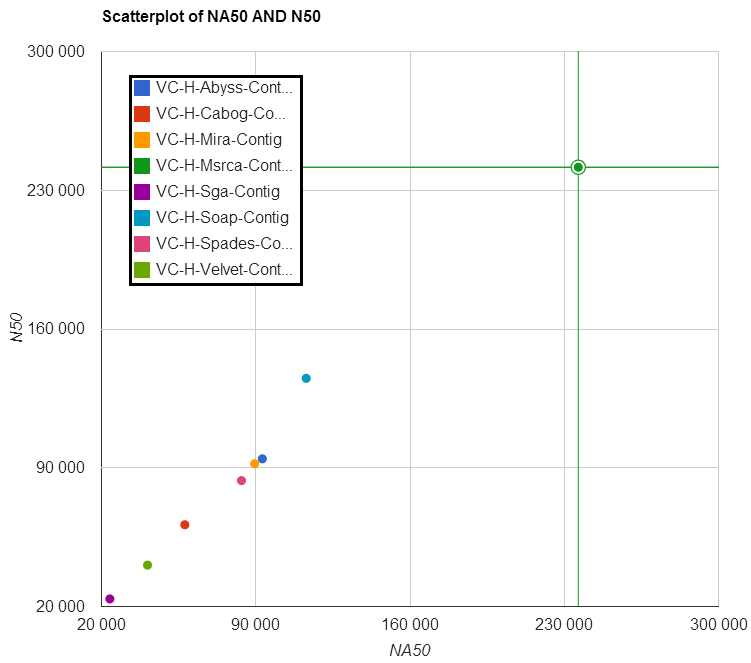


Figure S-A 5 Scatterplot on NA50 and N50 statistics on contigs  
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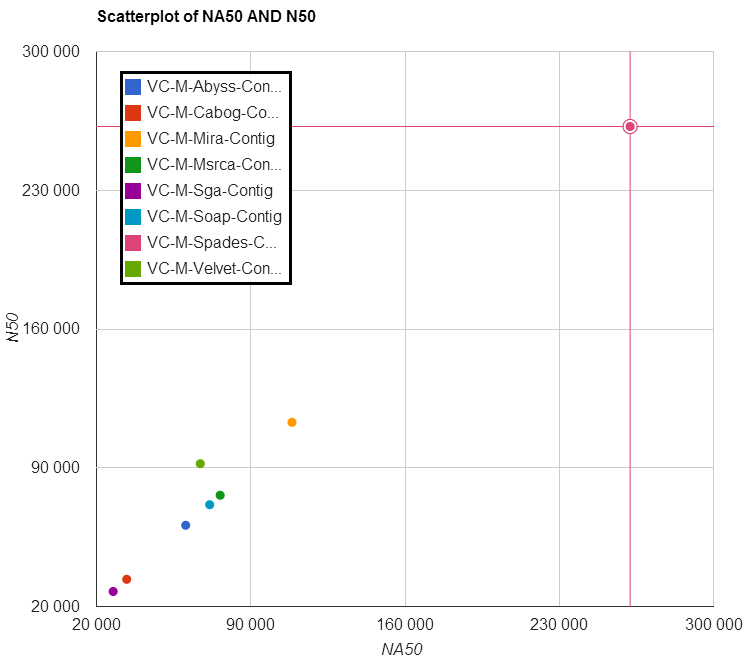


Figure S-A 6 Scatterplot on NA50 and N50 statistics on contigs   
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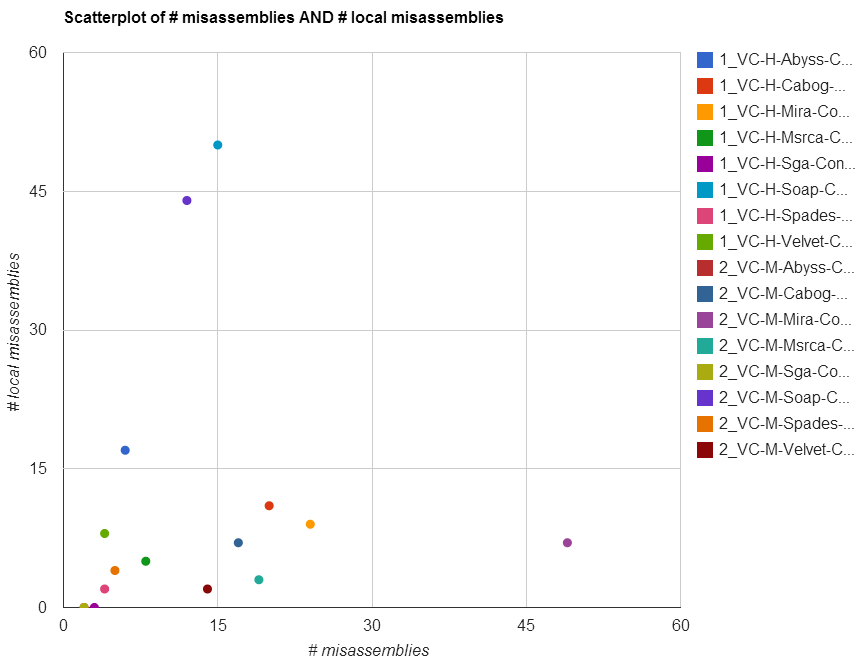


Figure S-A 7 Scatterplot of # misassemblies and # local misassemblies for contigs  
from HiSeq (1) and MiSeq (2) assemblies on *Vibrio cholerae*

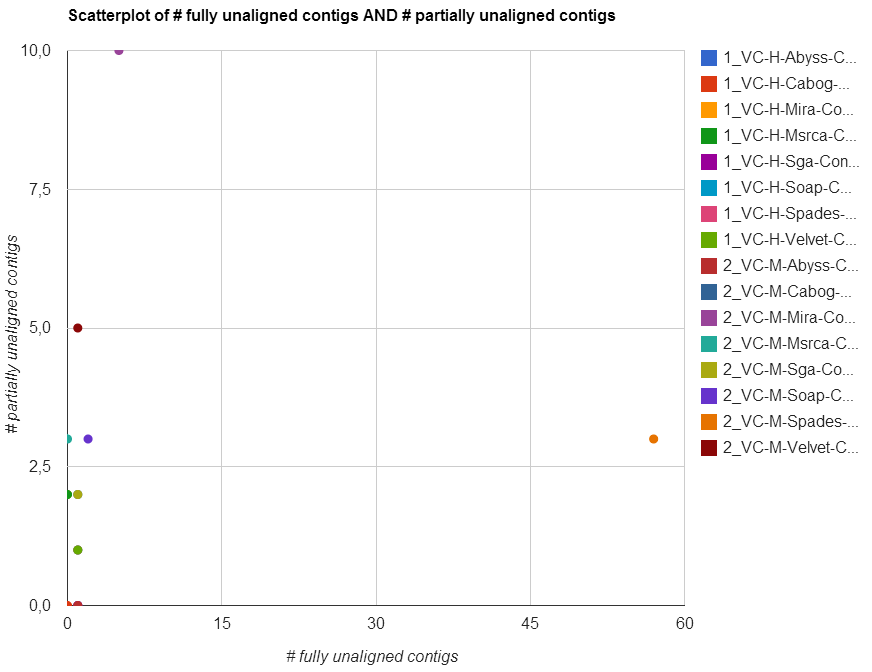


Figure S-A 8 Scatterplot of the number of unaligned contigs (Fully and partially) for contigs from HiSeq (1) and MiSeq (2) assemblies on *Vibrio cholerae*

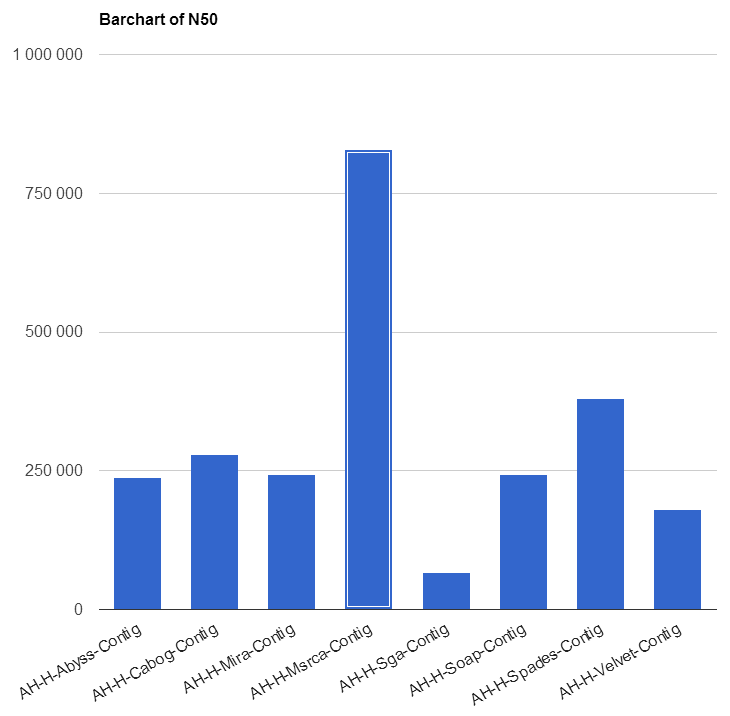


Figure S-A 9 Bar chart on N50 statistics on contigs from  
HiSeq assemblies on *Aeromonas hydrophila*

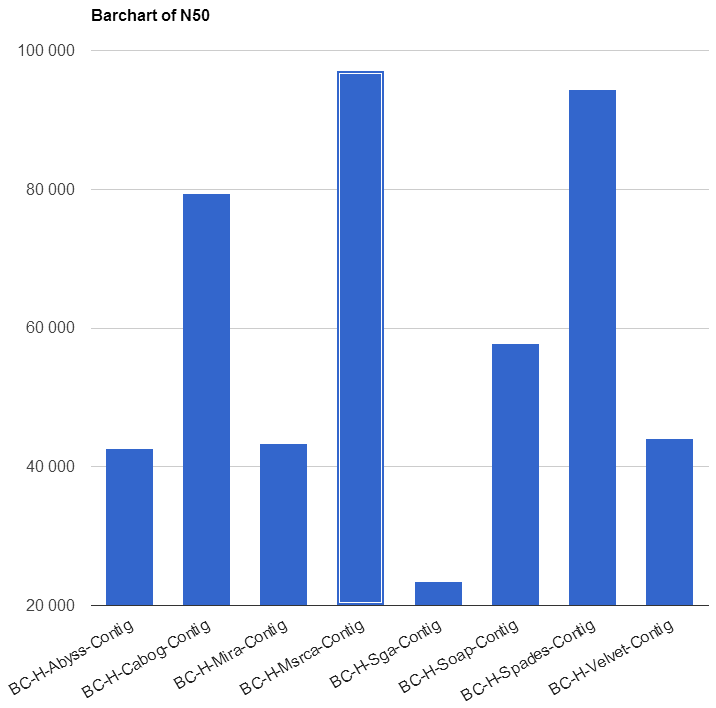


Figure S-A 10 Bar chart on N50 statistics on contigs   
from HiSeq assemblies on *Bacillus cereus*

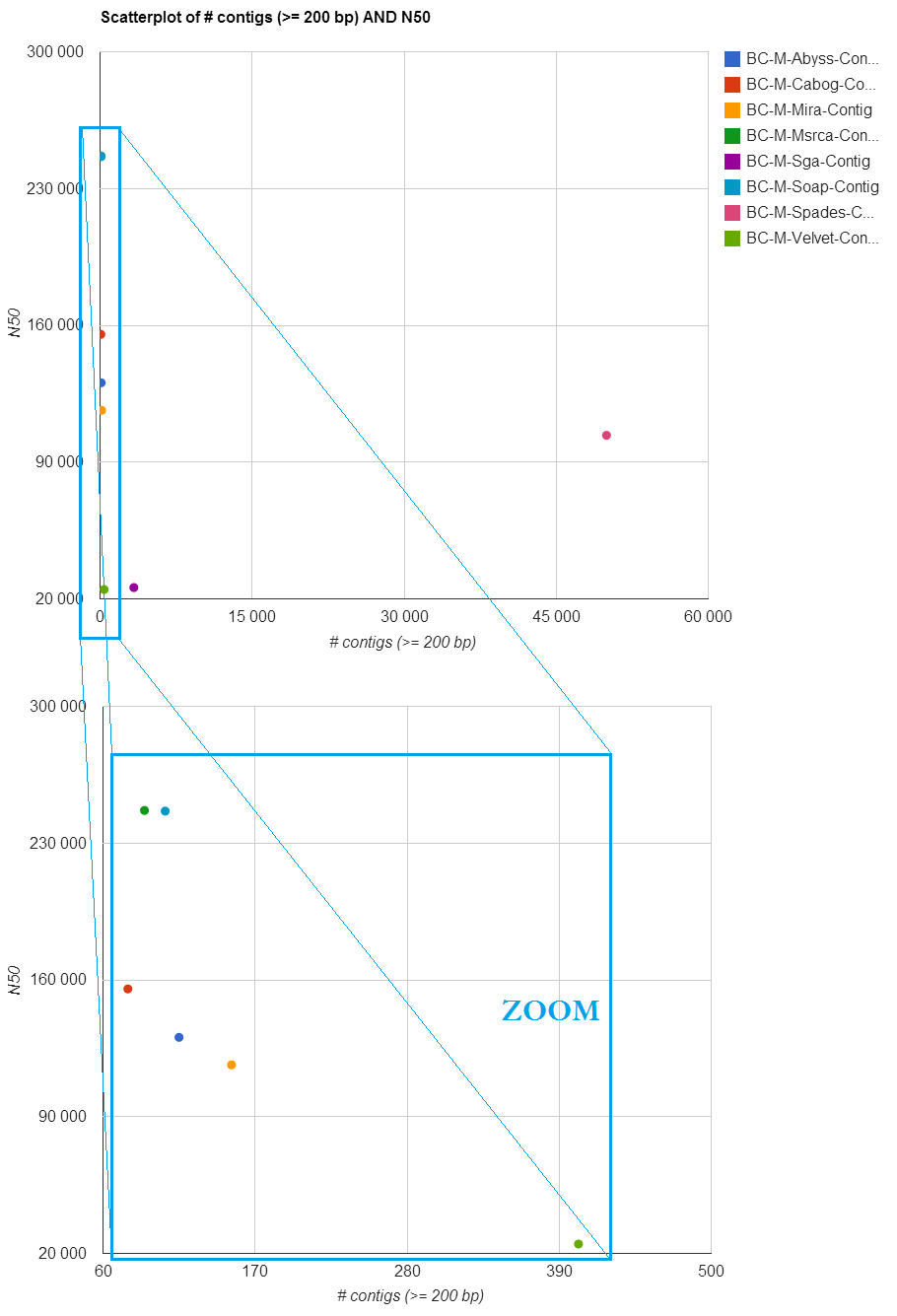


Figure S-A 11 Scatterplot of the number of contigs and N50 for contigs from MiSeq assemblies on *Bacillus cereus ATCC 10987*

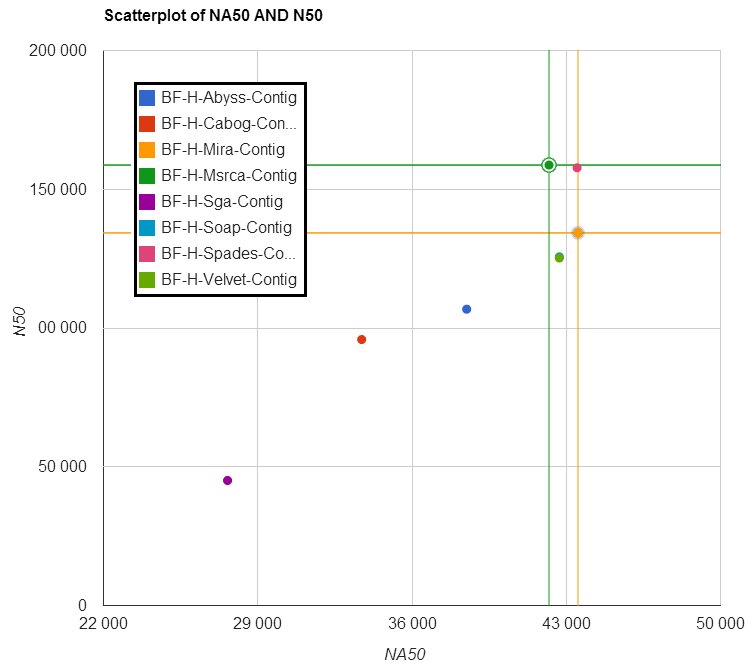


Figure S-A Scatterplot of NA50 and N50 statistics on contigs  
from HiSeq assemblies on *Bacteroides fragilis*

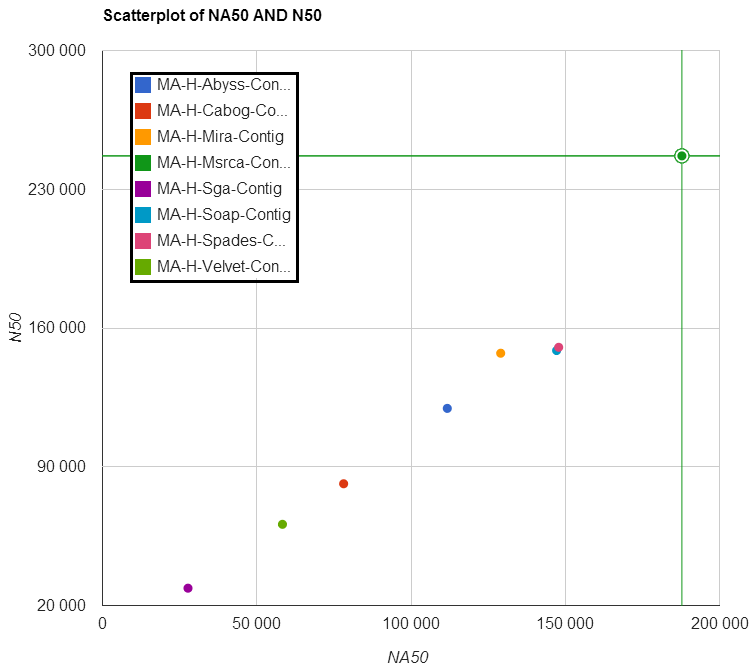


Figure S-A Scatterplot of NA50 and N50 statistics on contigs  
from HiSeq assemblies on *Mycobacterium abscessus*

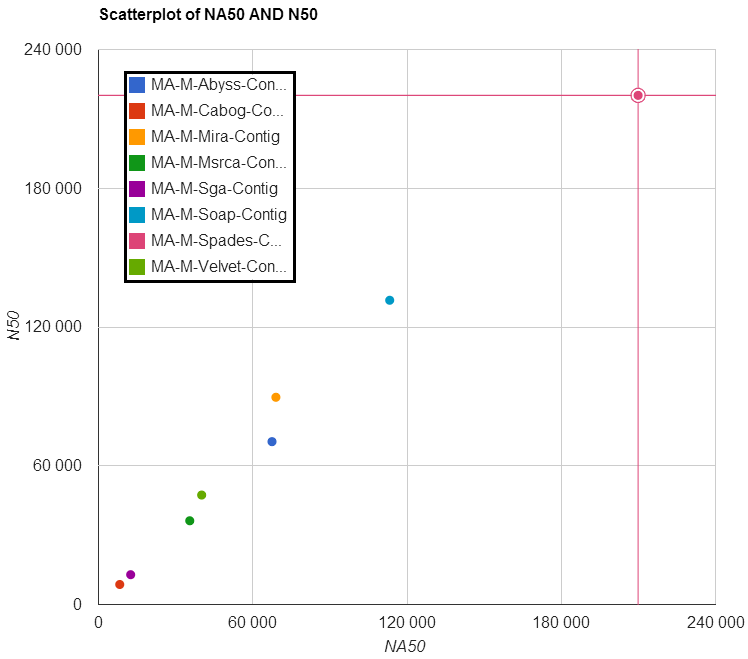


Figure S-A Scatterplot on NA50 and N50 statistics on contigs   
from MiSeq assemblies on *Mycobacterium abscessus*

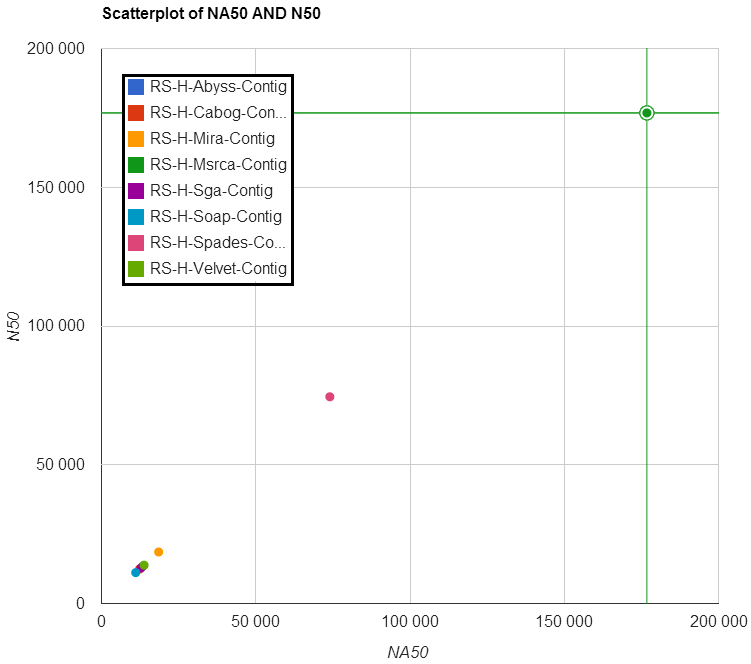


Figure S-A Scatterplot on NA50 and N50 statistics on contigs  
from HiSeq assemblies on *Rhodobacter sphaeroides*

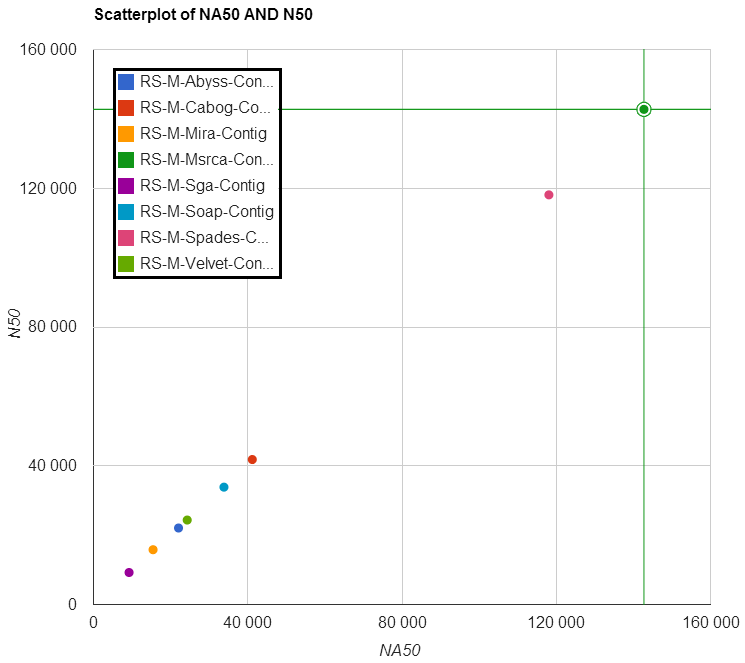


Figure S-A Scatterplot on NA50 and N50 statistics on contigs  
from MiSeq assemblies on Rhodobacter sphaeroides

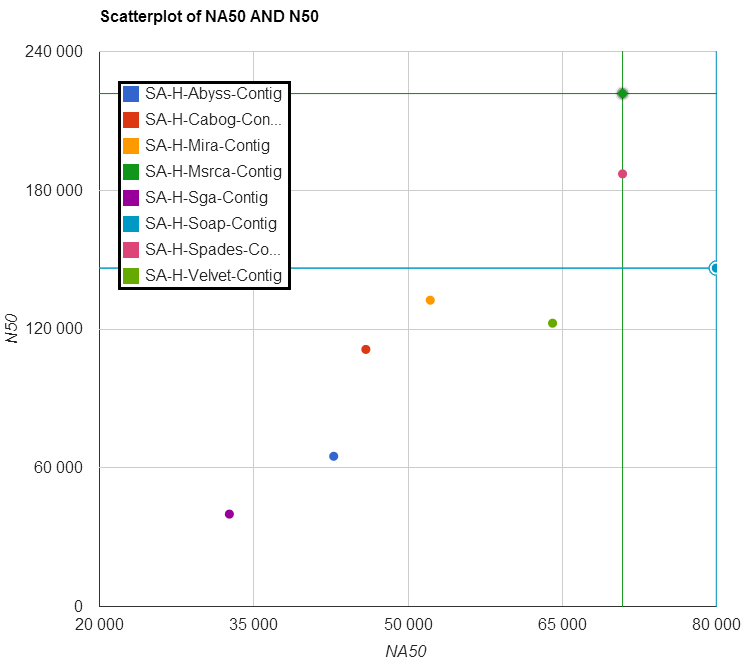


Figure S-A Scatterplot on NA50 and N50 statistics on contigs  
from HiSeq assemblies on *Staphylococcus aureus*

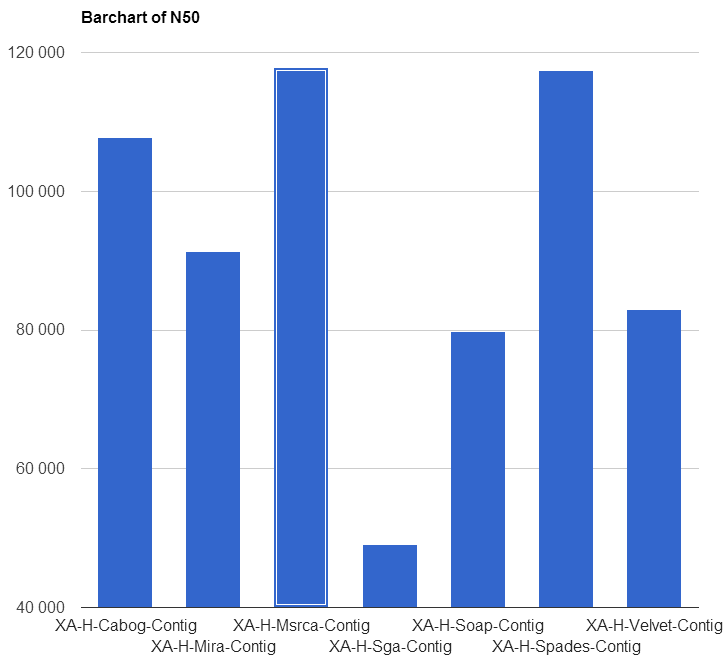


Figure S-A Bar chart on N50 statistics on contigs  
from HiSeq assemblies on *Xanthomonas axonopodis*

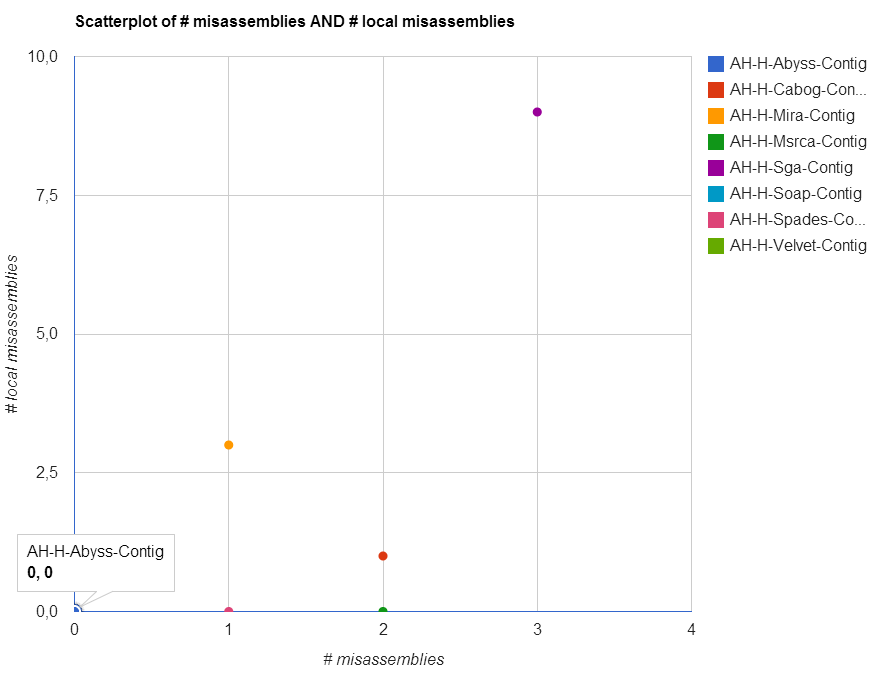


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Aeromonas hydrophila*

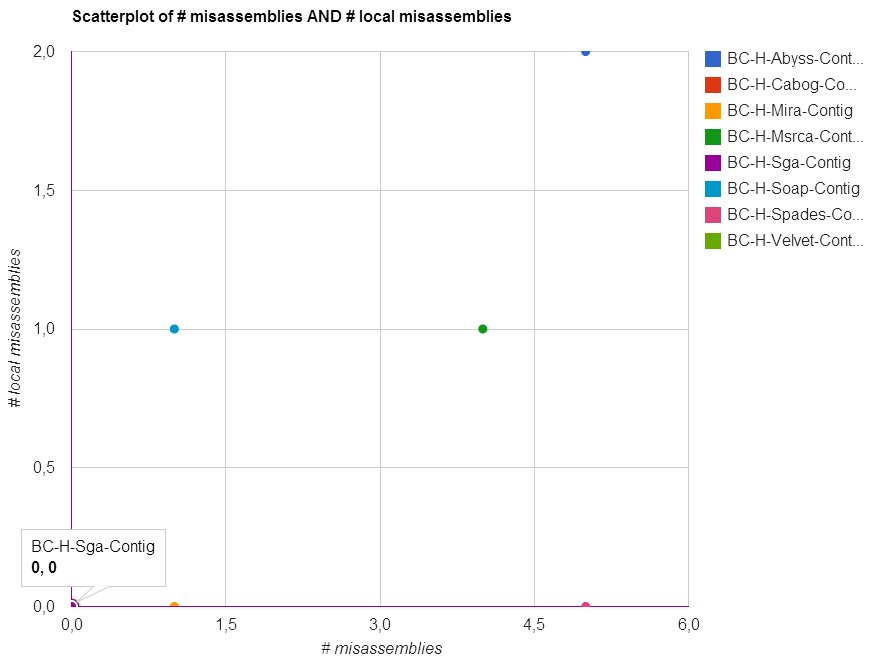


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Bacillus cereus*

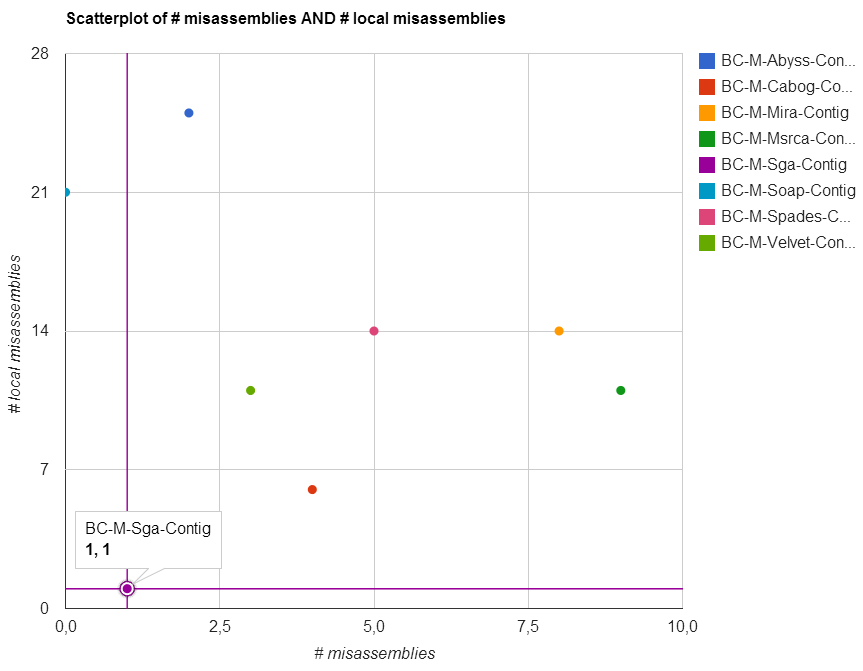


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from MiSeq assemblies on *Bacillus cereus*

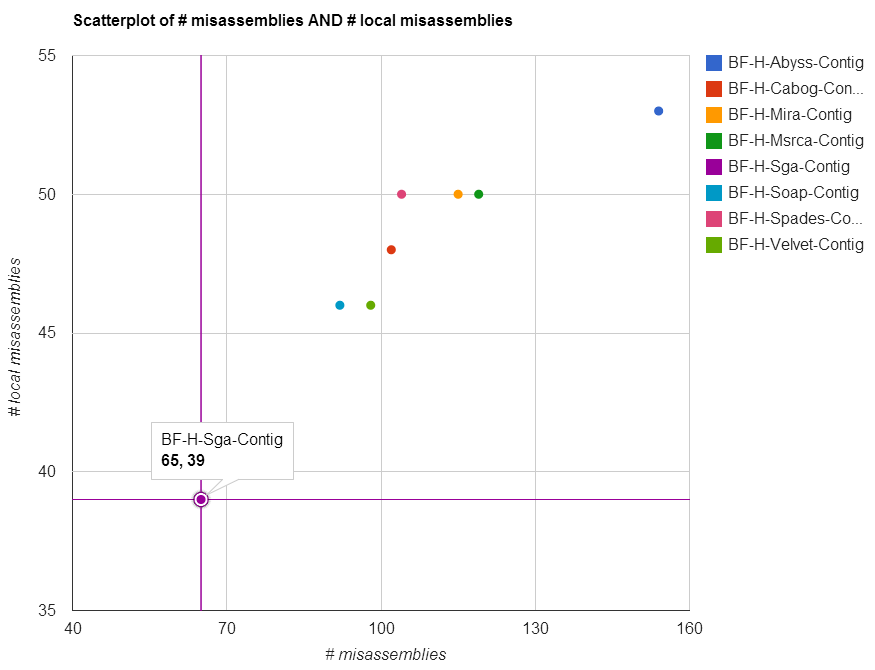


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Bacteroides fragilis*

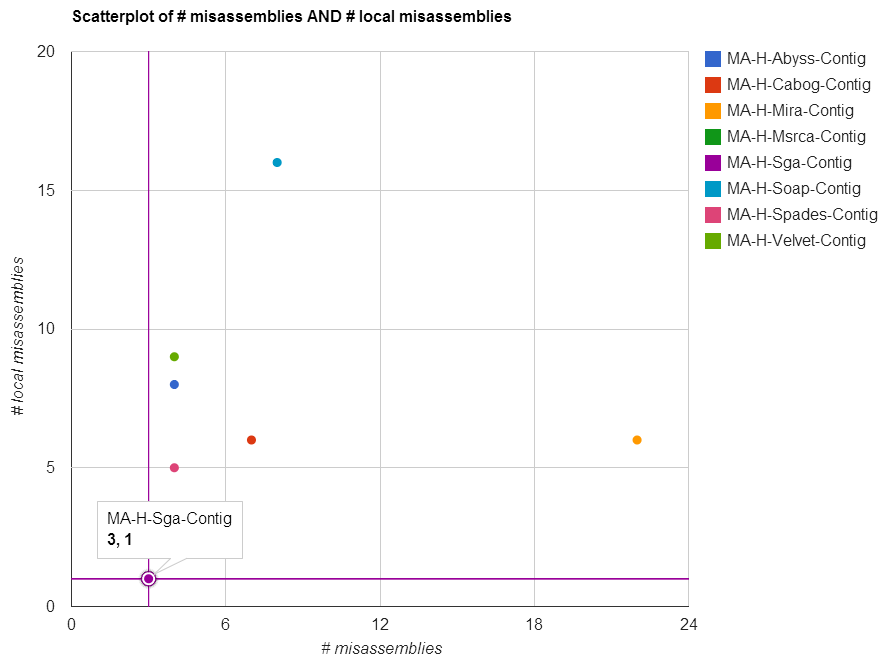


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Mycobacterium abscessus*

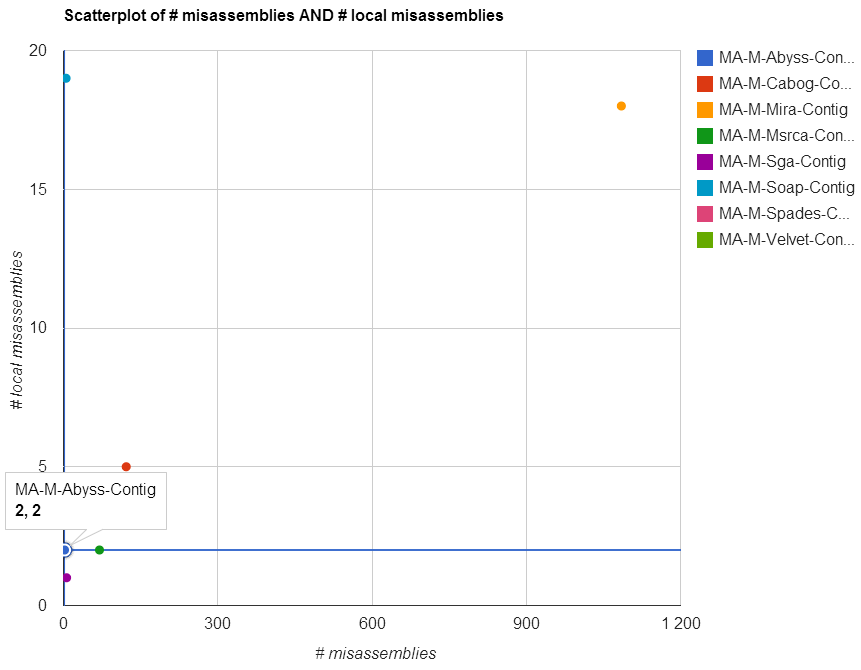


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from MiSeq assemblies on *Mycobacterium abscessus*

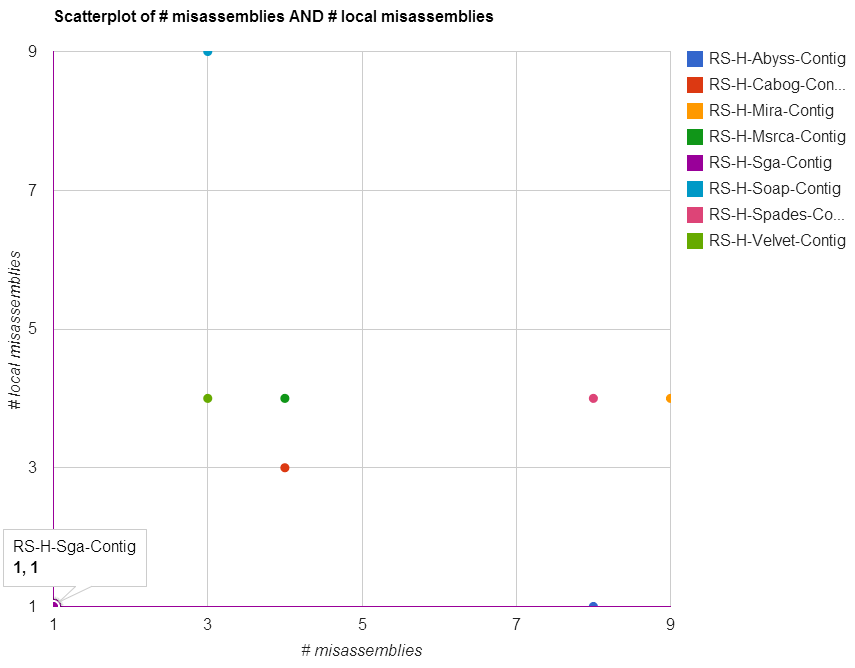


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Rhodobacter sphaeroides*

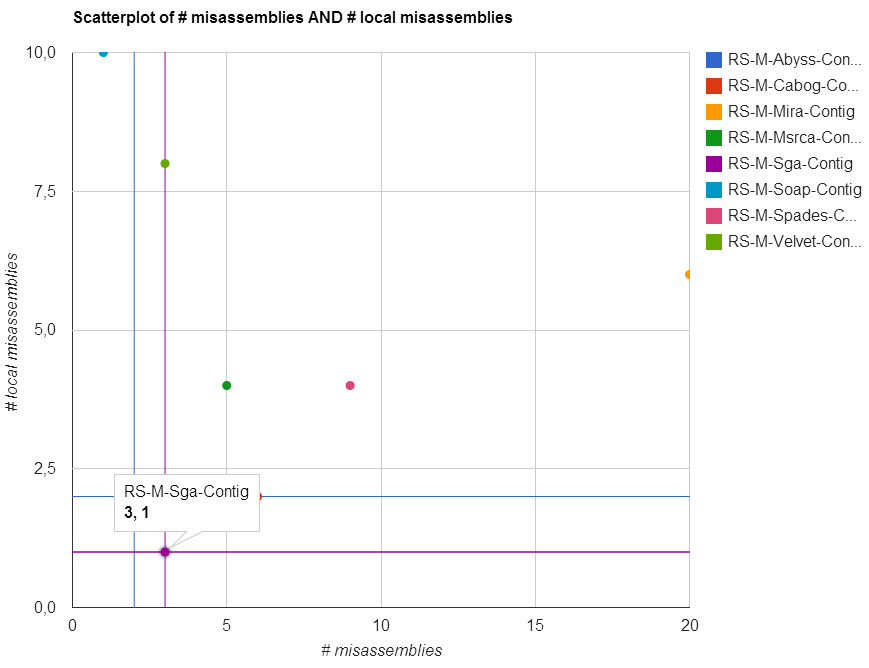


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from MiSeq assemblies on *Rhodobacter sphaeroides*

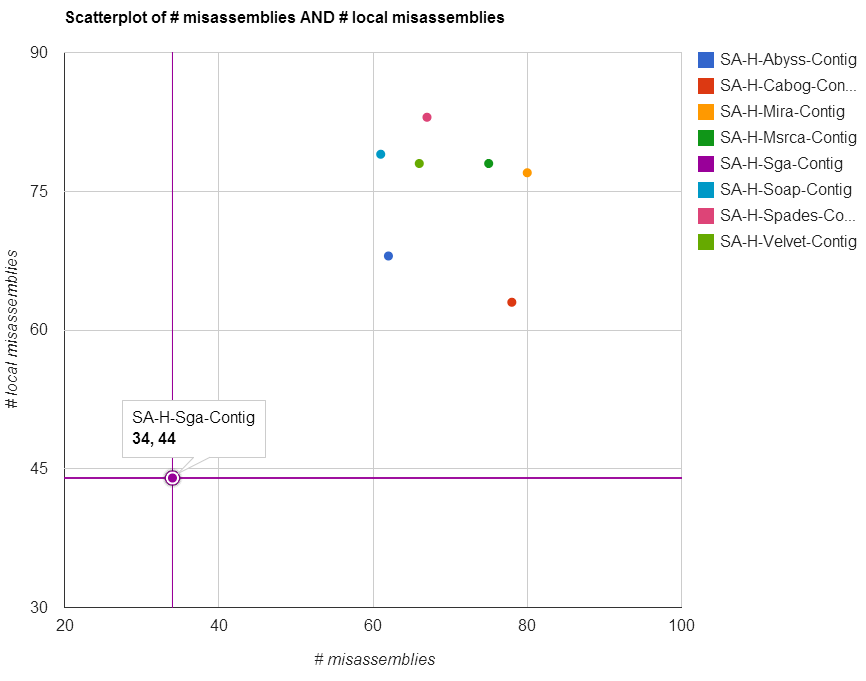


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Staphylococcus aureus*

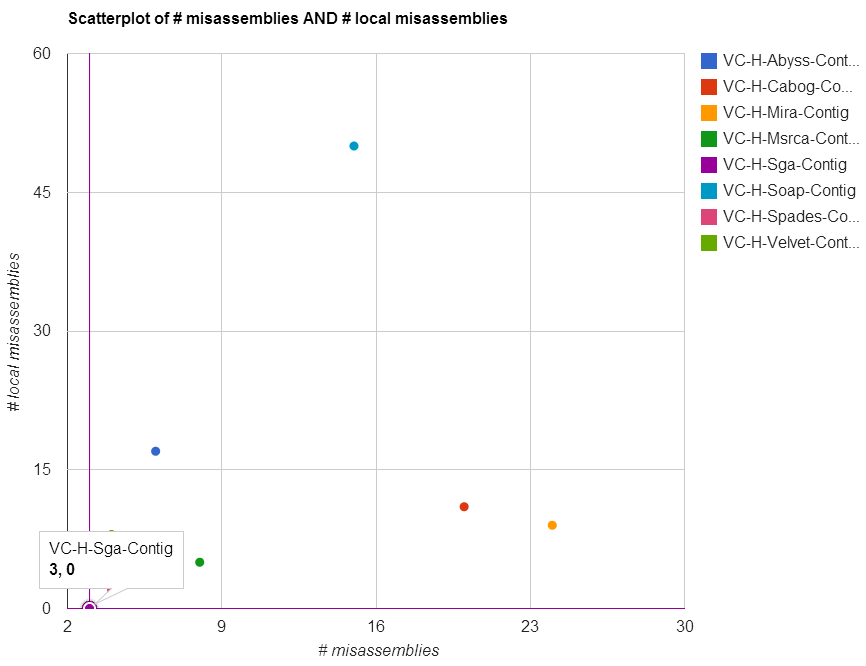


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from HiSeq assemblies on *Vibrio cholerae*

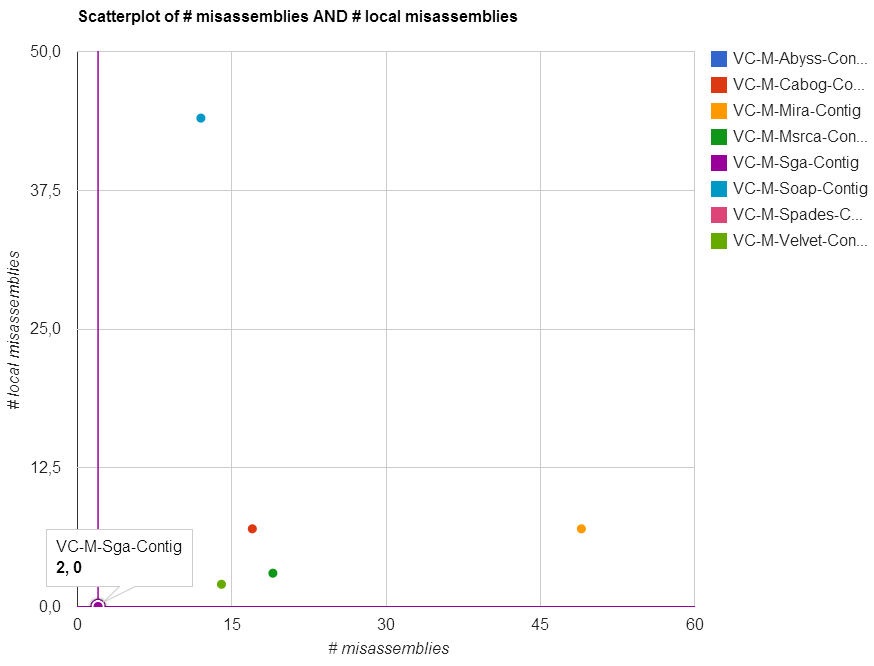


Figure S-A Scatterplot on #misassemblies and # local misassemblies on contigs from MiSeq assemblies on *Vibrio cholerae*