Report

	conting
# conting (> 1000 hm)	contigs 54
# contigs (>= 1000 bp)	
# contigs (>= 5000 bp)	45
# contigs (>= 10000 bp)	42
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	26
Total length (>= 1000 bp)	4708714
Total length (>= 5000 bp)	4691082
Total length (>= 10000 bp)	4668137
Total length (>= 25000 bp)	4620809
Total length (>= 50000 bp)	4177287
# contigs	64
Largest contig	349059
Total length	4716262
Reference length	4641652
GC (%)	50.72
Reference GC (%)	50.79
N50	177303
NG50	177303
N75	99302
NG75	99302
L50	9
LG50	9
L75	18
LG75	18
# misassemblies	4
# misassembled contigs	3
Misassembled contigs length	711322
# local misassemblies	1
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	97.664
Duplication ratio	1.040
# N's per 100 kbp	0.00
# mismatches per 100 kbp	246.71
# indels per 100 kbp	0.44
Largest alignment	333124
NA50	173841
NGA50	173841
NA75	87711
NGA75	94467
LA50	10
LGA50	10
LA75	20
LGA75	19
L 25A/3	19

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Misassemblies report

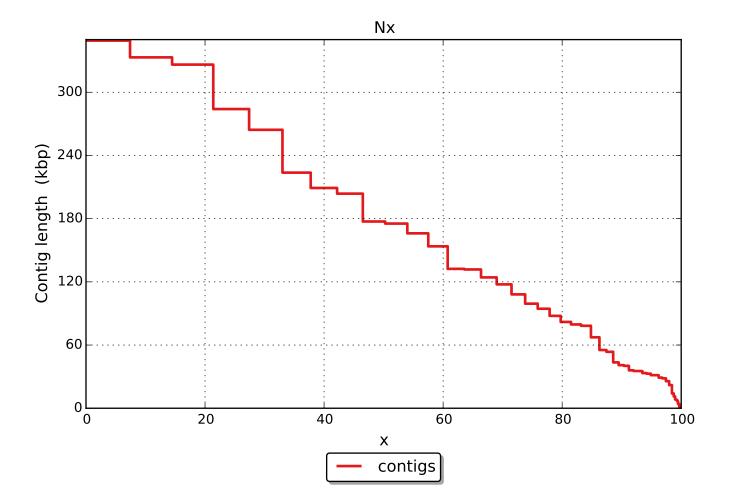
	contigs
# misassemblies	4
# relocations	4
# translocations	0
# inversions	0
# possibly misassembled contigs	1
# misassembled contigs	3
Misassembled contigs length	711322
# local misassemblies	1
# mismatches	11184
# indels	20
# short indels	20
# long indels	0
Indels length	20

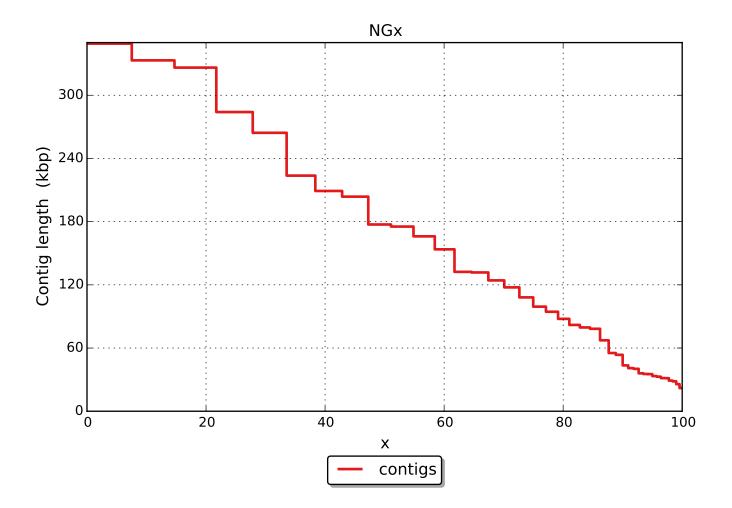
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

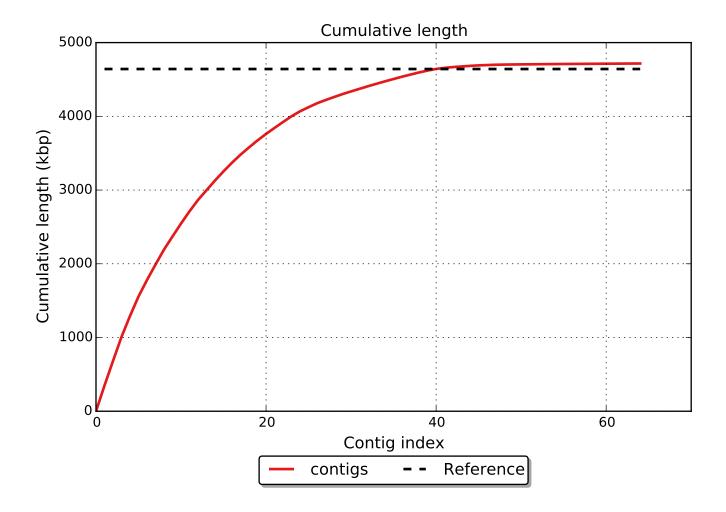
Unaligned report

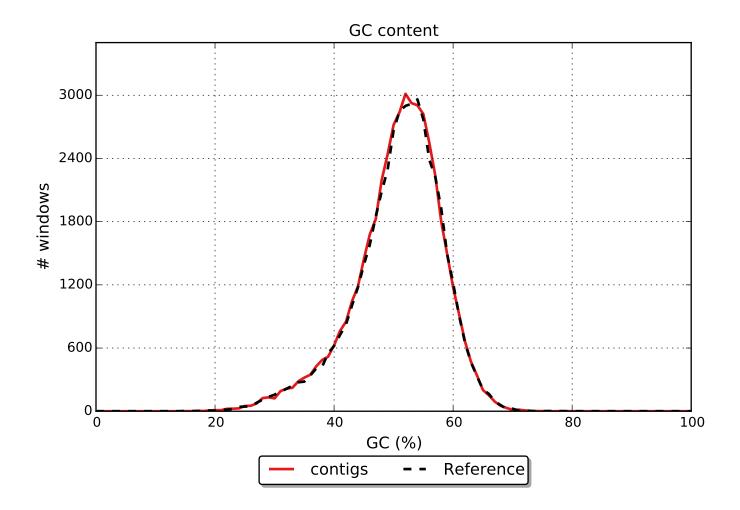
	contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

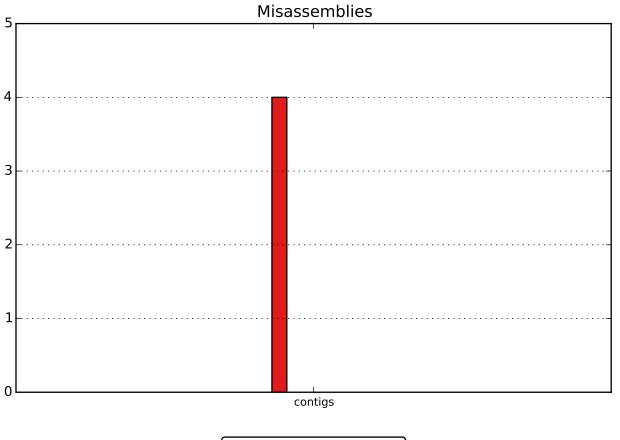
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).











relocations

