Report

# contigs (>= 1000 bp)	scaffolds
_	31
# contigs (>= 5000 bp)	27
# contigs (>= 10000 bp)	26
# contigs (>= 25000 bp)	25
# contigs (>= 50000 bp)	22
Total length (>= 1000 bp)	4696185
Total length (>= 5000 bp)	4691496
Total length (>= 10000 bp)	4686030
Total length (>= 25000 bp)	4672863
Total length (>= 50000 bp)	4576116
# contigs	41
Largest contig	529644
Total length	4702926
Reference length	4641652
GC (%)	50.82
Reference GC (%)	50.78
N50	305443
NG50	305443
N75	176954
NG75	182223
L50	6
LG50	6
L75	12
LG75	11
# misassemblies	3
# misassembled contigs	2
Misassembled contigs length	578950
# local misassemblies	1
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	99.143
Duplication ratio	1.022
# N's per 100 kbp	0.47
# mismatches per 100 kbp	375.52
# indels per 100 kbp	1.00
Largest alignment	529644
I NATO	301763
NA50	301763
NGA50	
NGA50 NA75	128117
NGA50 NA75 NGA75	128117 128117
NGA50 NA75 NGA75 LA50	128117 7
NGA50 NA75 NGA75 LA50 LGA50	128117 7 7
NGA50 NA75 NGA75 LA50	128117 7

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

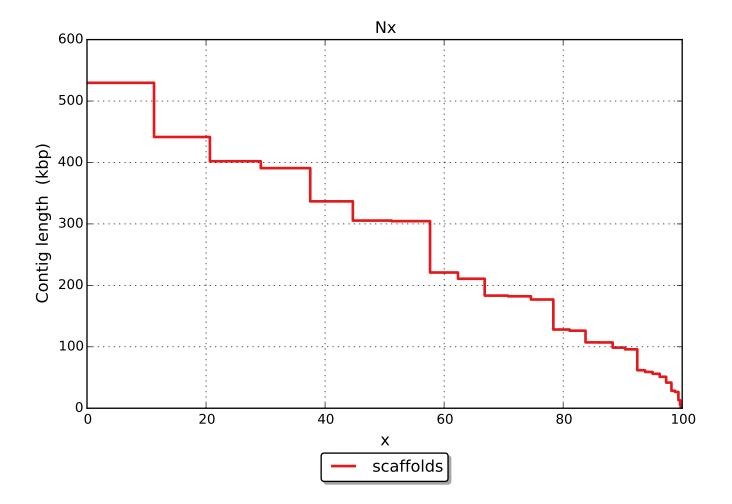
	scaffolds
# misassemblies	3
# relocations	3
# translocations	0
# inversions	0
# possibly misassembled contigs	1
# misassembled contigs	2
Misassembled contigs length	578950
# local misassemblies	1
# mismatches	17281
# indels	46
# short indels	45
# long indels	1
Indels length	67

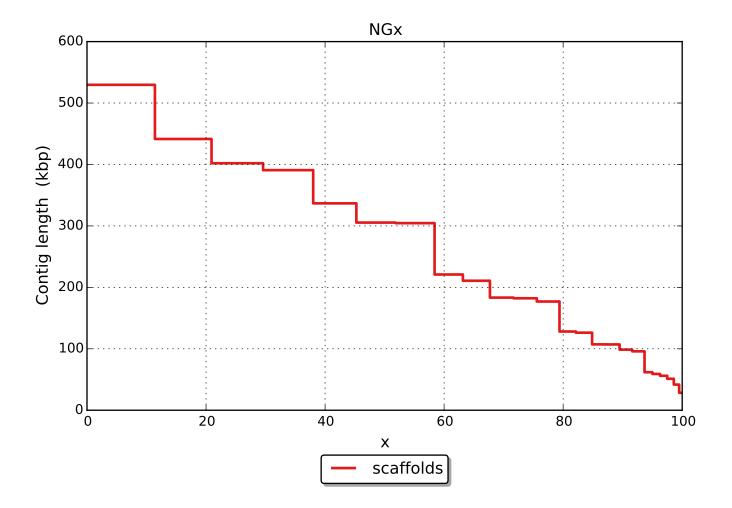
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

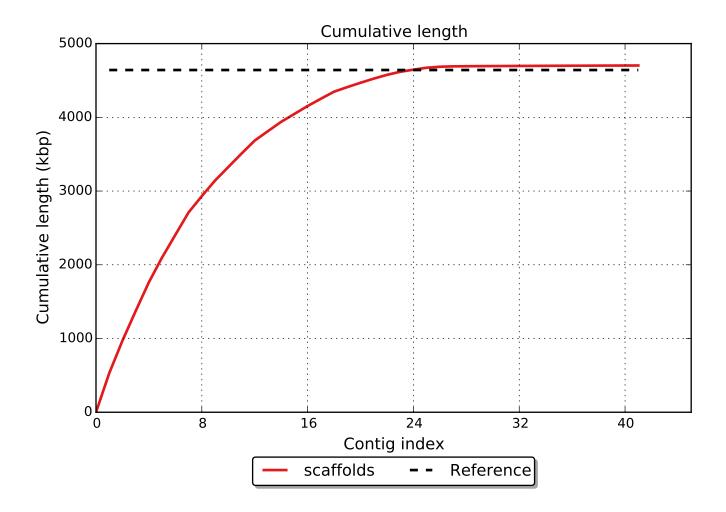
Unaligned report

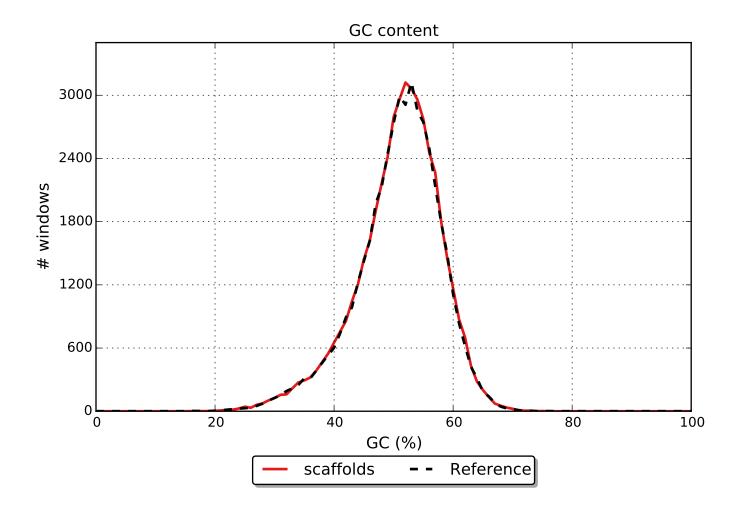
	scaffolds
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	22

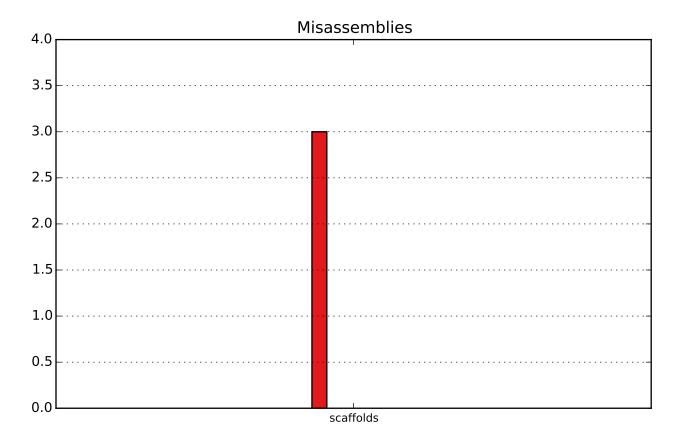
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).











relocations

