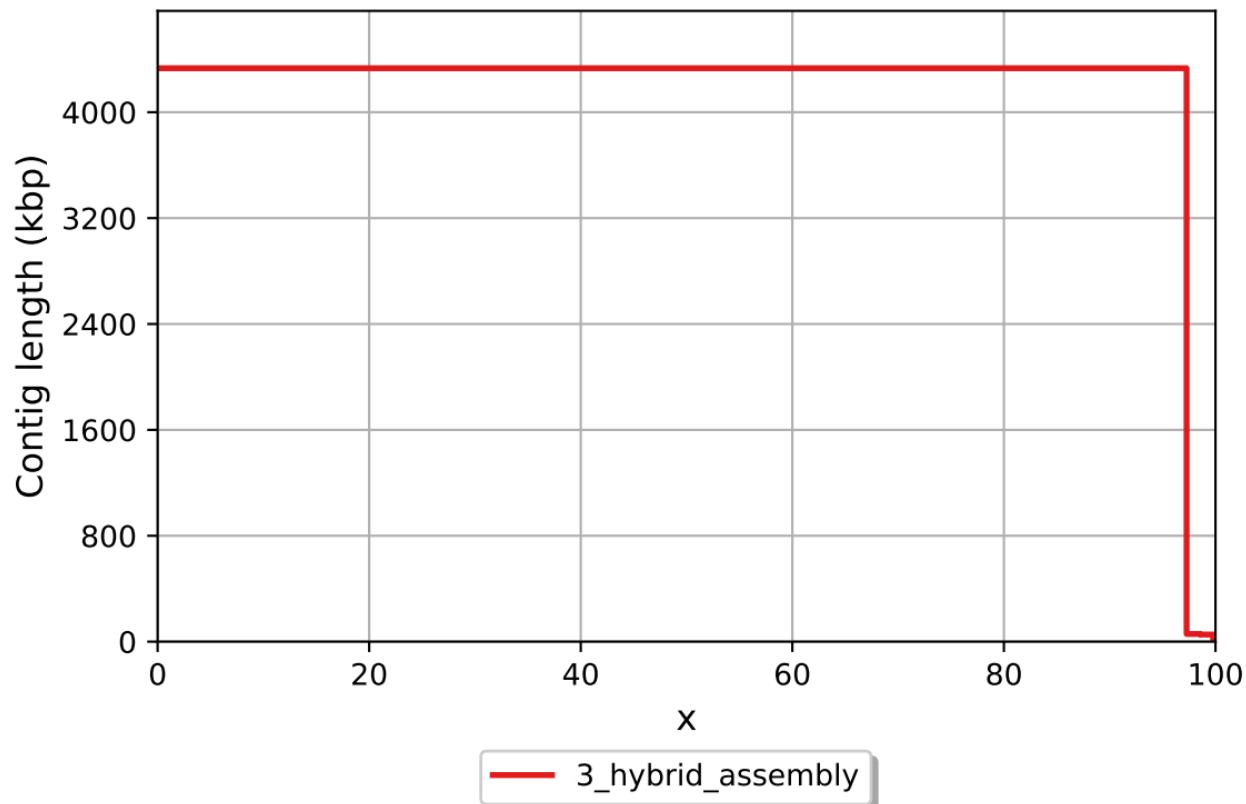


Report

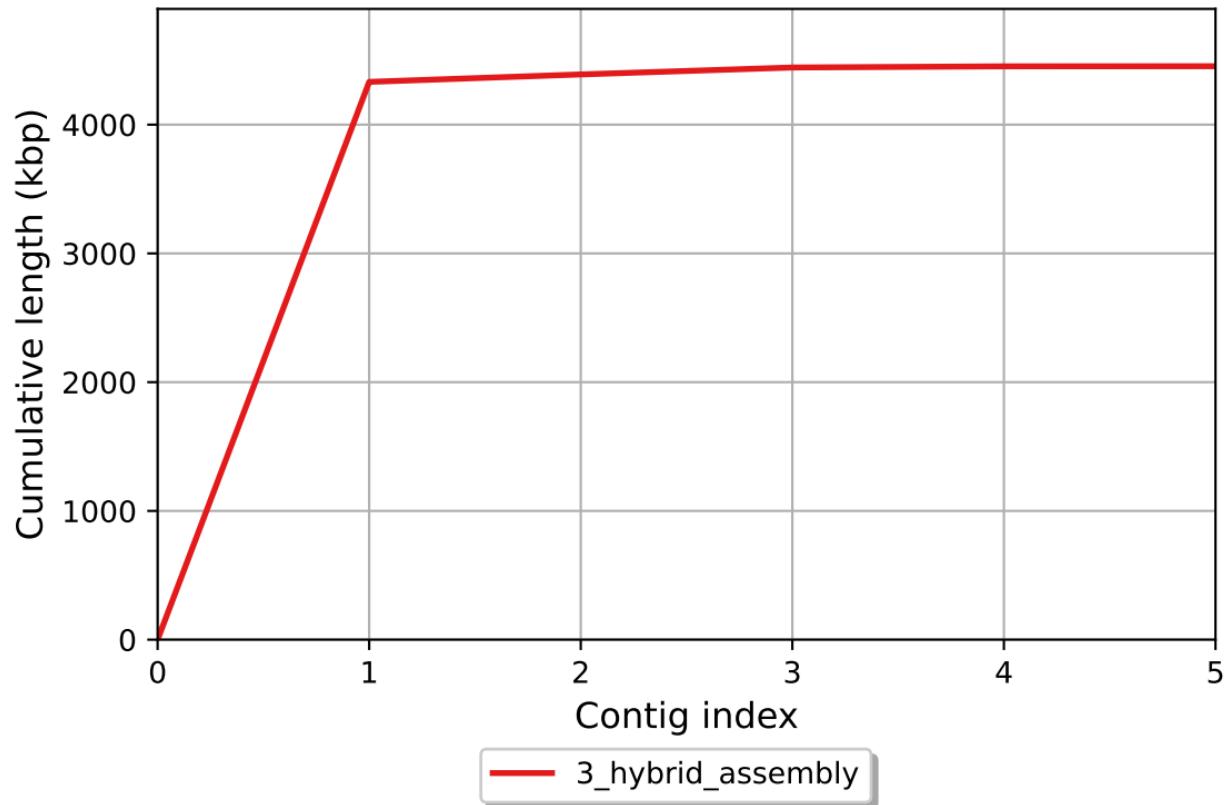
	3_hybrid_assembly
# contigs (>= 0 bp)	7
# contigs (>= 1000 bp)	4
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	4454332
Total length (>= 1000 bp)	4453091
Total length (>= 5000 bp)	4453091
Total length (>= 10000 bp)	4443965
Total length (>= 25000 bp)	4443965
Total length (>= 50000 bp)	4443965
# contigs	5
Largest contig	4332314
Total length	4453949
Reference length	-
Estimated reference length	-
GC (%)	45.71
Reference GC (%)	-
N50	4332314
NG50	-
N90	4332314
NG90	-
aun	4215421.0
aUNG	-
L50	1
LG50	-
L90	1
LG90	-
# total reads	-
# left	-
# right	-
Mapped (%)	-
Reference mapped (%)	-
Properly paired (%)	-
Reference properly paired (%)	-
Avg. coverage depth	-
Reference avg. coverage depth	-
Coverage >= 1x (%)	-
Reference coverage >= 1x (%)	-
# large blocks misassemblies	-
# misassemblies	-
# misassembled contigs	-
Misassembled contigs length	-
# local misassemblies	-
# scaffold gap ext. mis.	-
# scaffold gap loc. mis.	-
# structural variations	-
# possible TEs	-
# unaligned mis. contigs	-
# unaligned contigs	-
Unaligned length	-
Genome fraction (%)	-
Duplication ratio	-
Avg contig read support	-
# N's per 100 kbp	0.00
# mismatches per 100 kbp	-
# indels per 100 kbp	-
# genomic features	-
# operons	-
Complete BUSCO (%)	-
Partial BUSCO (%)	-
# predicted genes (unique)	2495
# predicted genes (>= 0 bp)	2520 + 0 part
# predicted genes (>= 300 bp)	2445 + 0 part
# predicted genes (>= 1500 bp)	816 + 0 part
# predicted genes (>= 3000 bp)	216 + 0 part
# predicted rRNA genes	15 + 0 part
Largest alignment	-
Total aligned length	-
NA50	-
NGA50	-
NA90	-
NGA90	-
auna	-
aUNGA	-
LA50	-
LGA50	-
LA90	-
LGA90	-
K-mer-based compl. (%)	-
K-mer-based cor. length (%)	-
K-mer-based mis. length (%)	-

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).

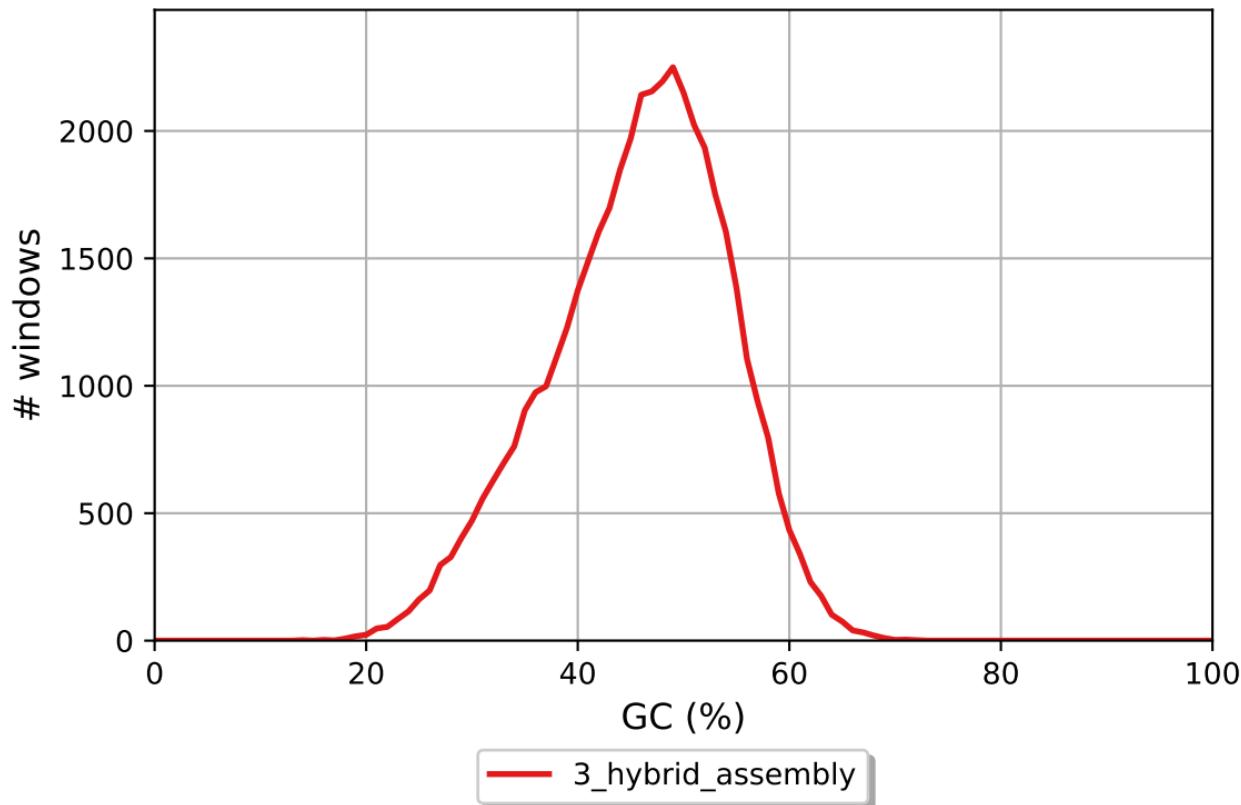
Nx



Cumulative length



GC content



3_hybrid_assembly GC content

