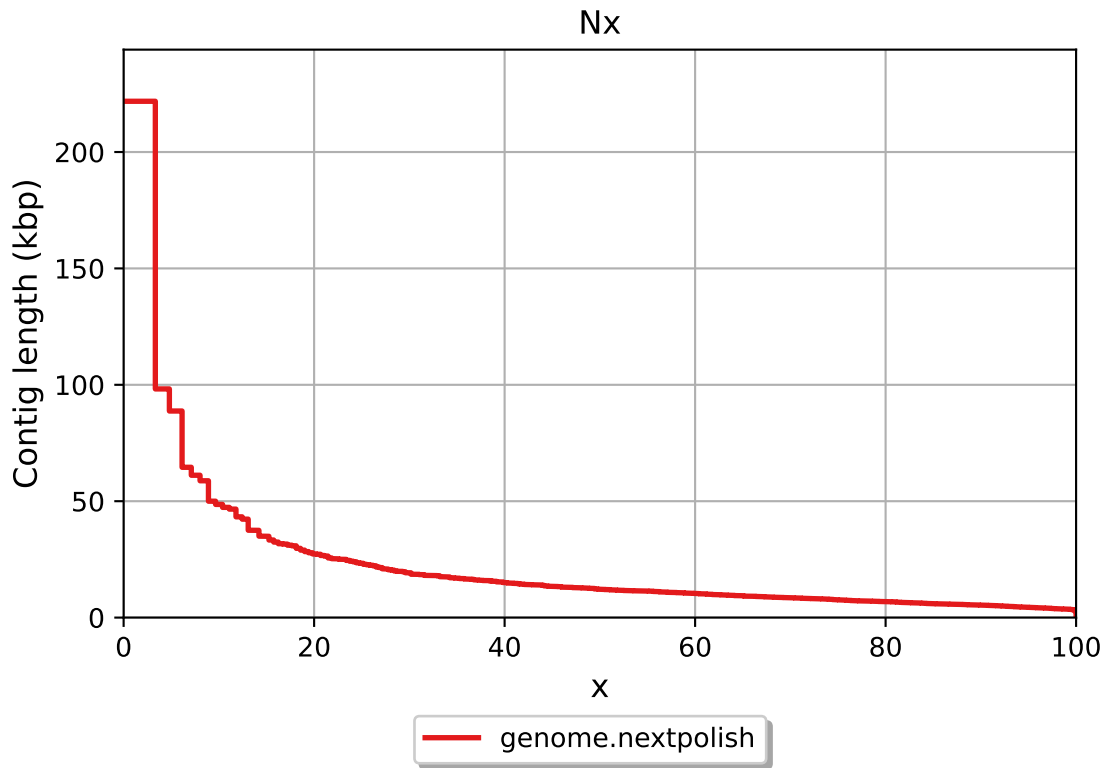


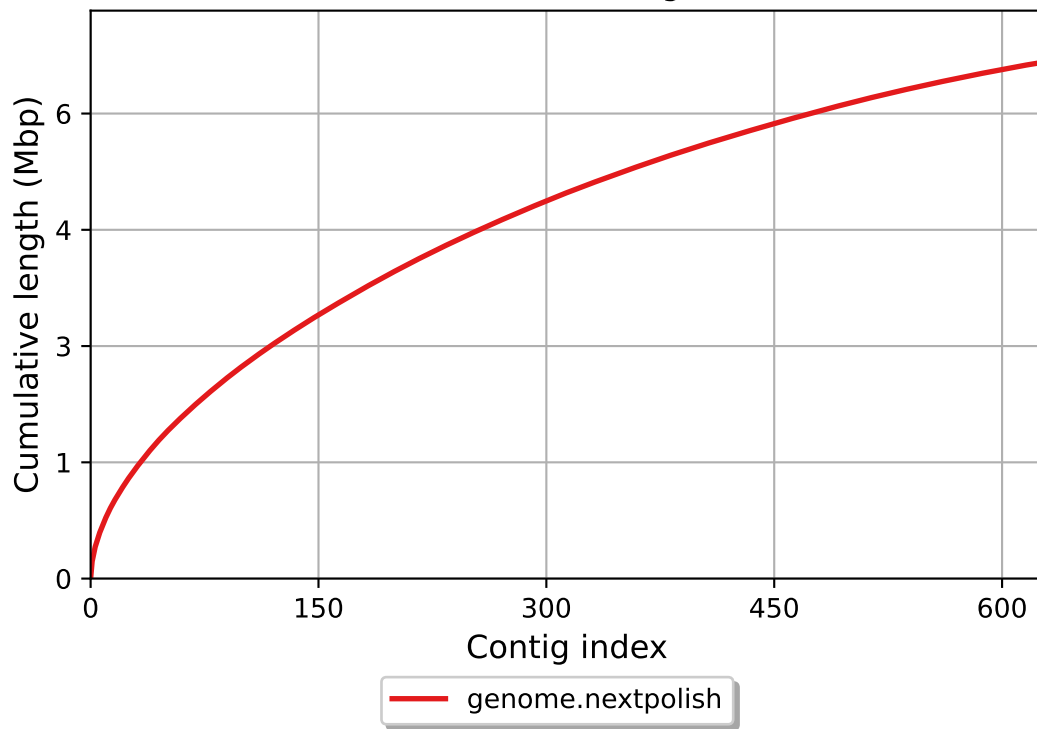
Report

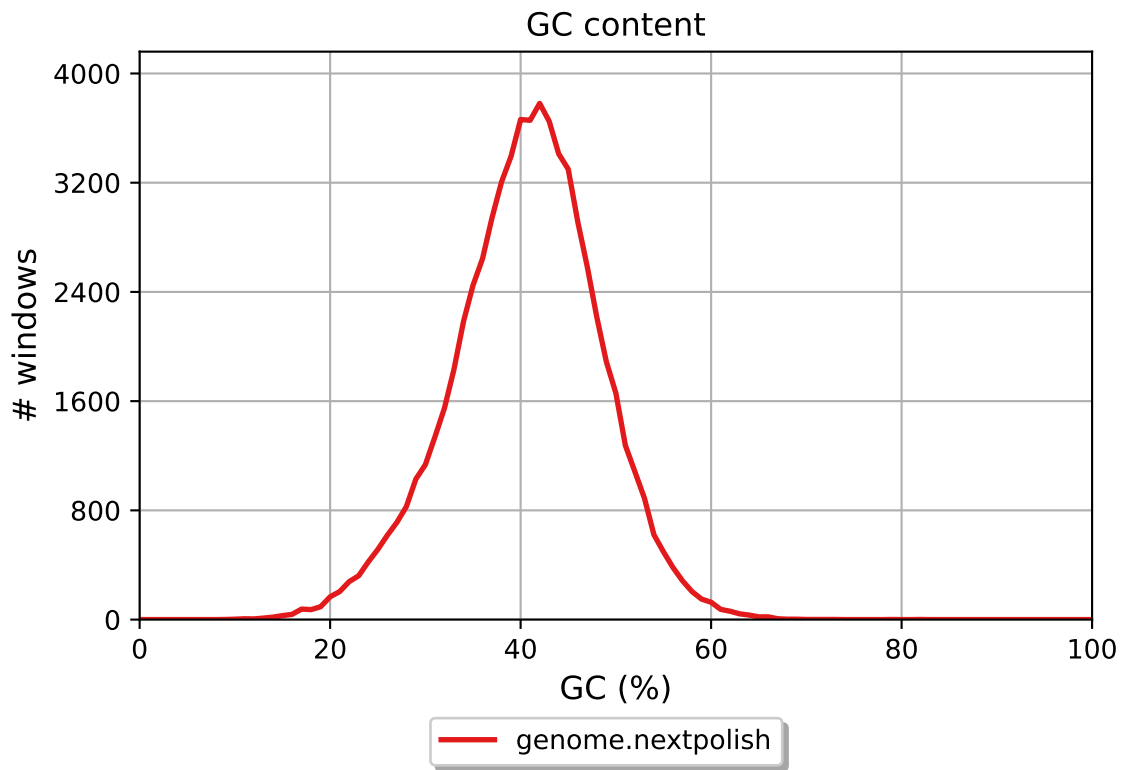
	genome.nextpolish
# contigs (>= 0 bp)	627
# contigs (>= 1000 bp)	627
# contigs (>= 5000 bp)	501
# contigs (>= 10000 bp)	214
# contigs (>= 25000 bp)	34
# contigs (>= 50000 bp)	6
Total length (>= 0 bp)	6661581
Total length (>= 1000 bp)	6661581
Total length (>= 5000 bp)	6142252
Total length (>= 10000 bp)	4106193
Total length (>= 25000 bp)	1531053
Total length (>= 50000 bp)	593257
# contigs	627
Largest contig	221801
Total length	6661581
GC (%)	40.56
N50	12139
N90	5373
auN	25153.4
L50	145
L90	473
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 1 part

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).



Cumulative length





genome.nextpolish GC content

