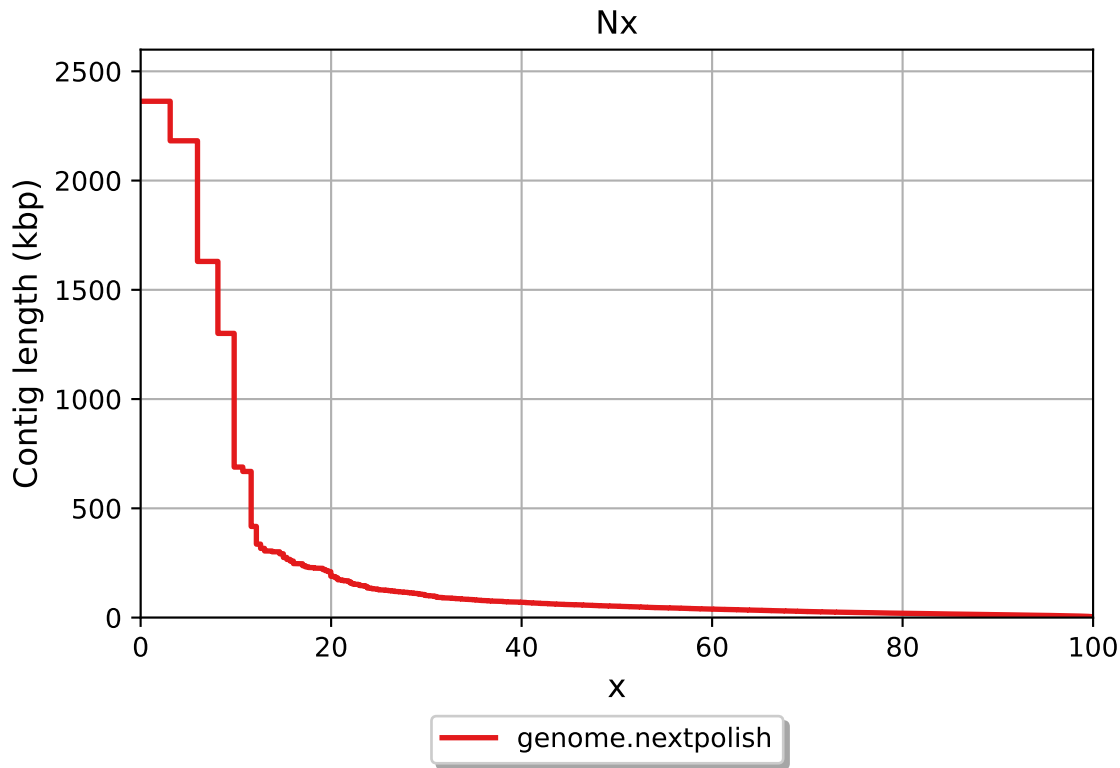


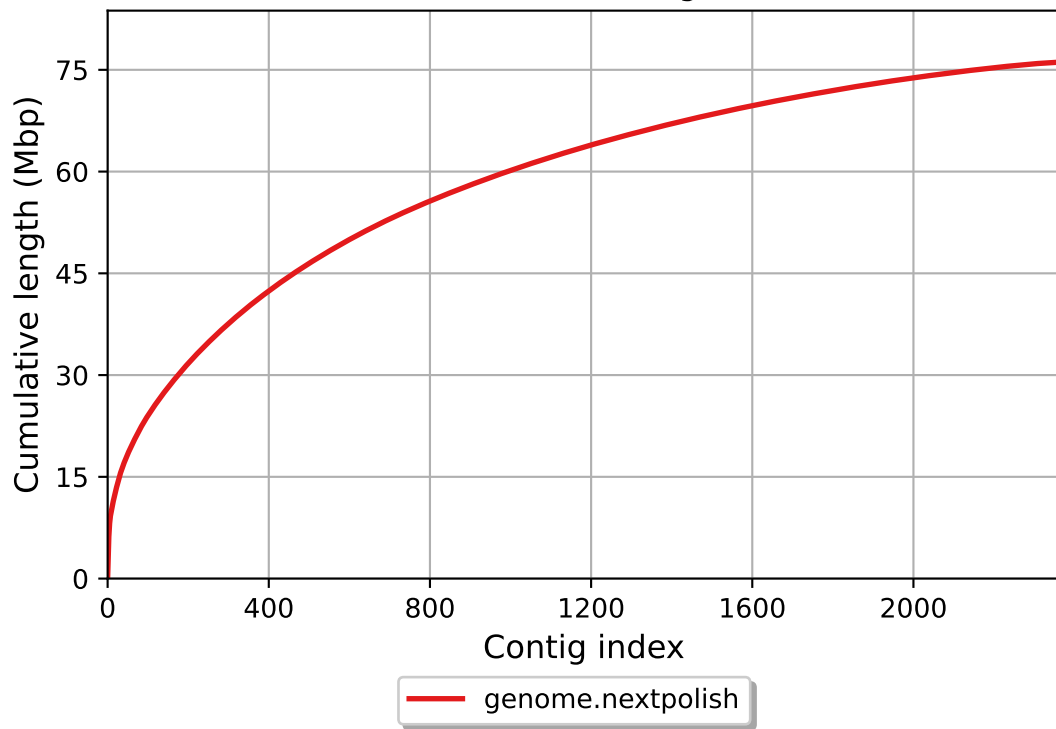
Report

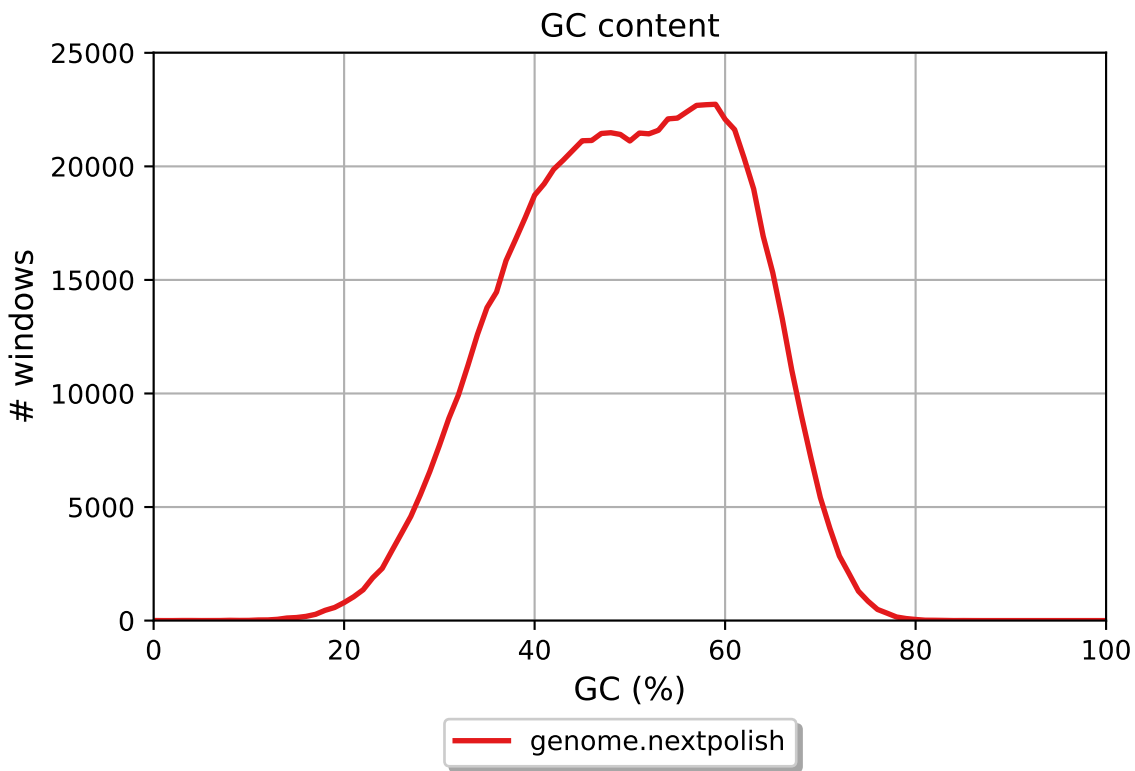
	genome.nextpolish
# contigs (>= 0 bp)	2364
# contigs (>= 1000 bp)	2356
# contigs (>= 5000 bp)	2297
# contigs (>= 10000 bp)	1835
# contigs (>= 25000 bp)	793
# contigs (>= 50000 bp)	329
Total length (>= 0 bp)	76113880
Total length (>= 1000 bp)	76107761
Total length (>= 5000 bp)	75874217
Total length (>= 10000 bp)	72309612
Total length (>= 25000 bp)	55481303
Total length (>= 50000 bp)	39082025
# contigs	2364
Largest contig	2362900
Total length	76113880
GC (%)	49.61
N50	52160
N90	13253
auN	268759.5
L50	309
L90	1505
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	176 + 12 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

