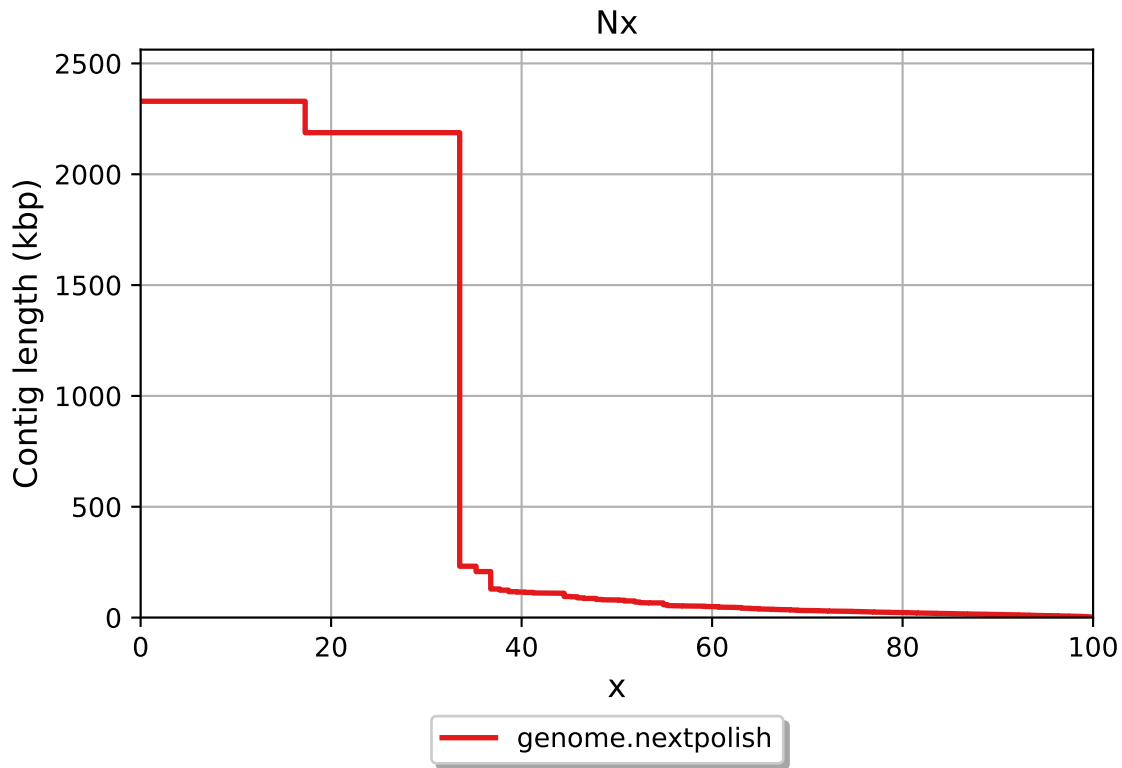


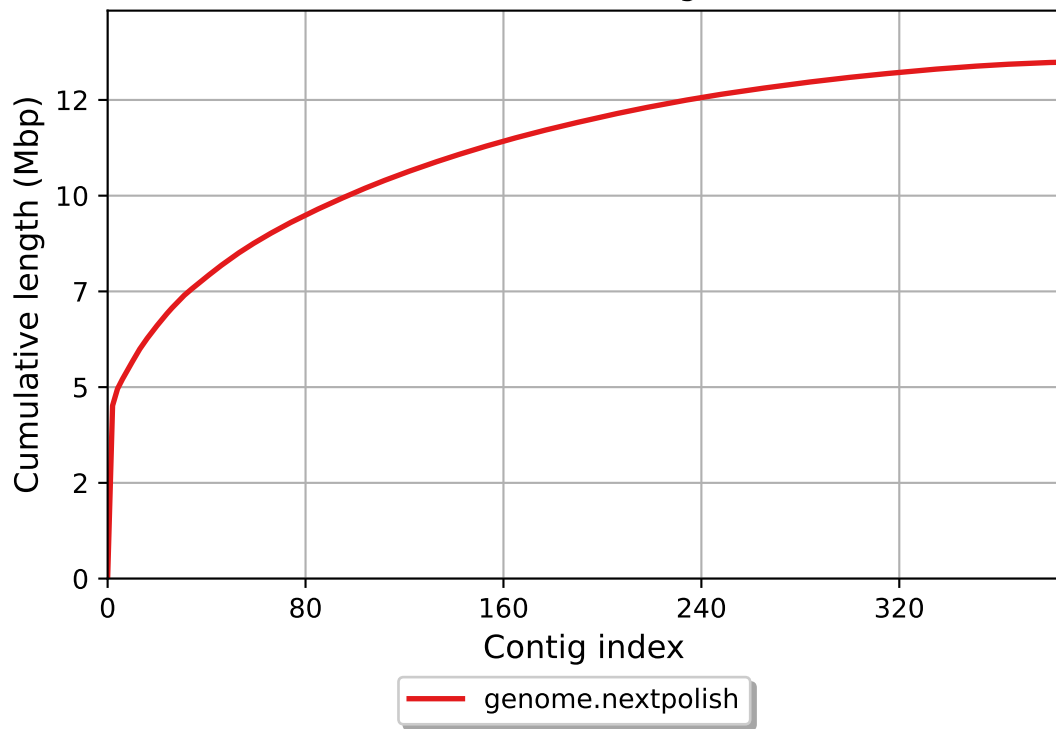
Report

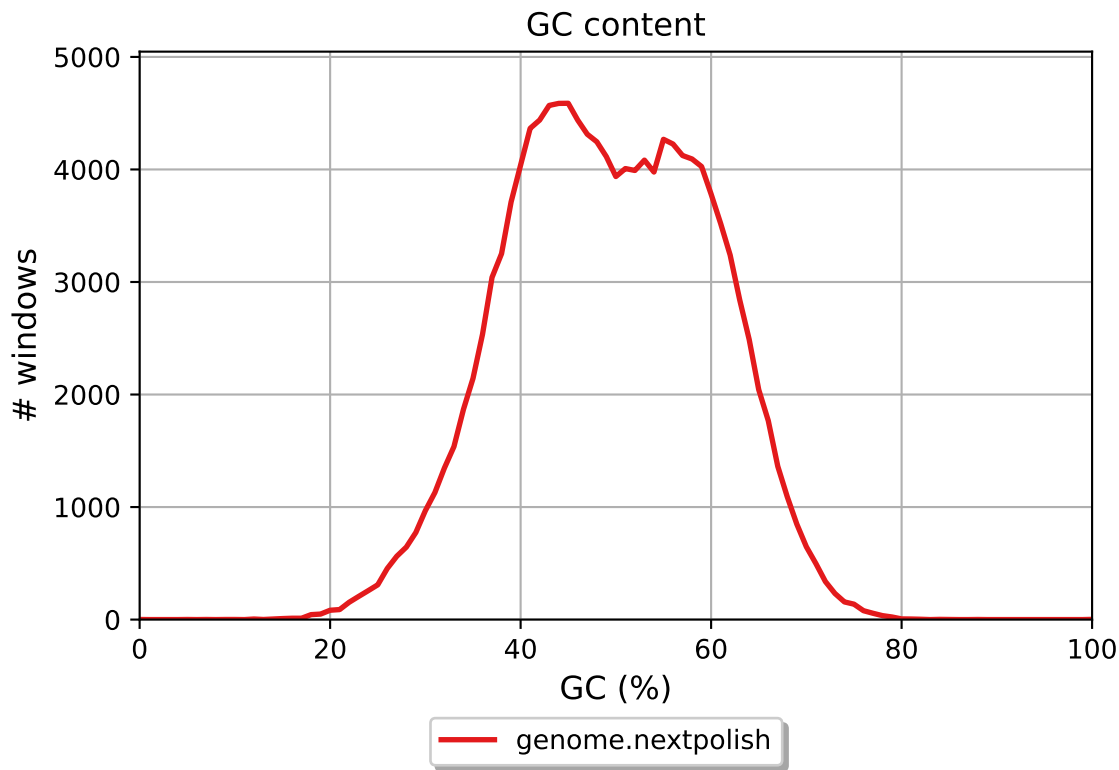
	genome.nextpolish
# contigs (>= 0 bp)	385
# contigs (>= 1000 bp)	385
# contigs (>= 5000 bp)	339
# contigs (>= 10000 bp)	247
# contigs (>= 25000 bp)	111
# contigs (>= 50000 bp)	43
Total length (>= 0 bp)	13483973
Total length (>= 1000 bp)	13483973
Total length (>= 5000 bp)	13326281
Total length (>= 10000 bp)	12637448
Total length (>= 25000 bp)	10374886
Total length (>= 50000 bp)	8033260
# contigs	385
Largest contig	2329305
Total length	13483973
GC (%)	49.27
N50	79333
N90	13902
auN	793757.5
L50	22
L90	206
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	43 + 0 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

