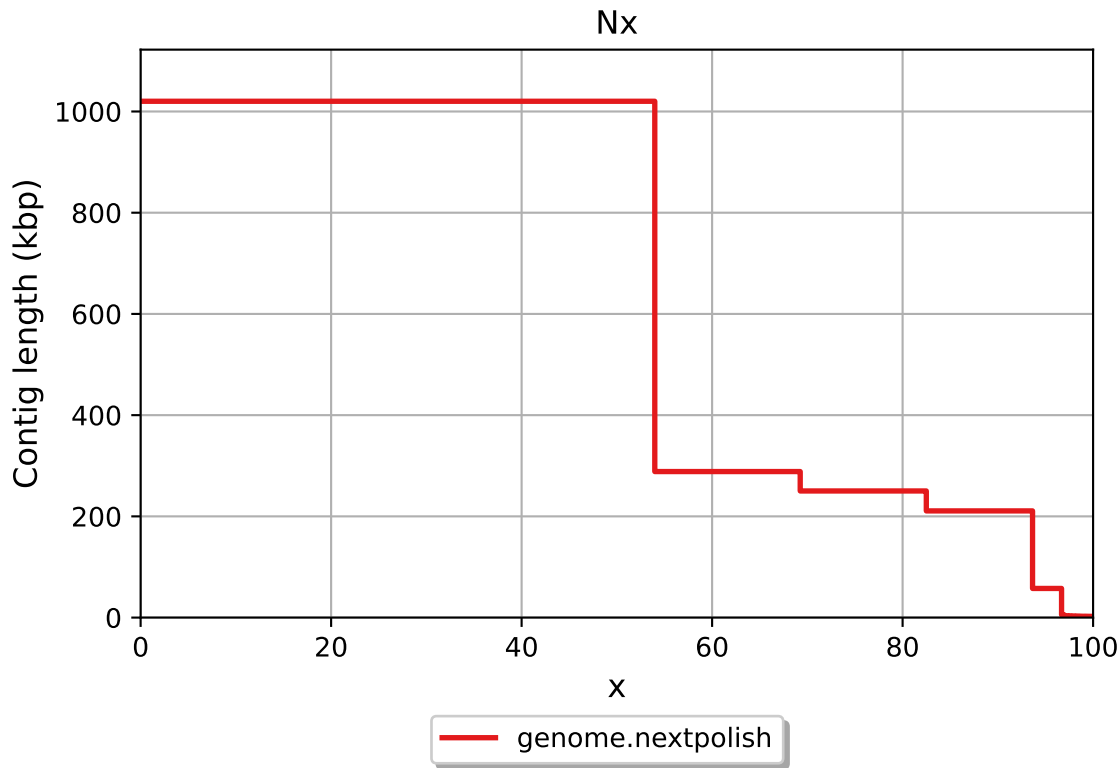
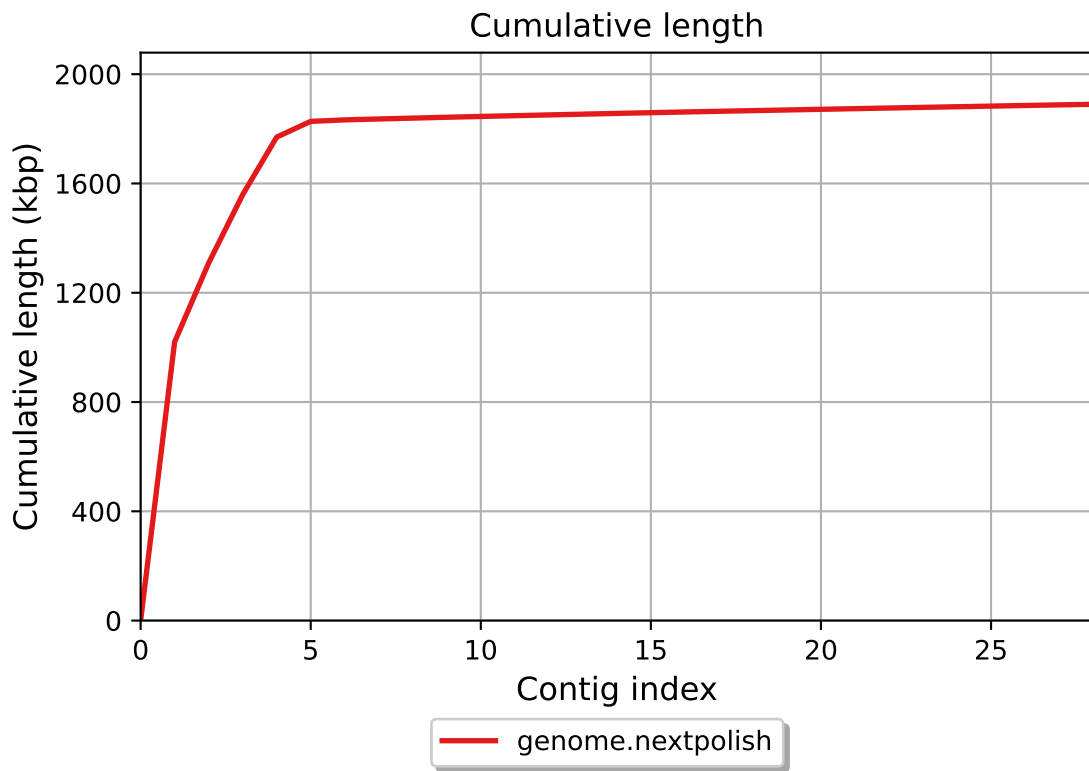


Report

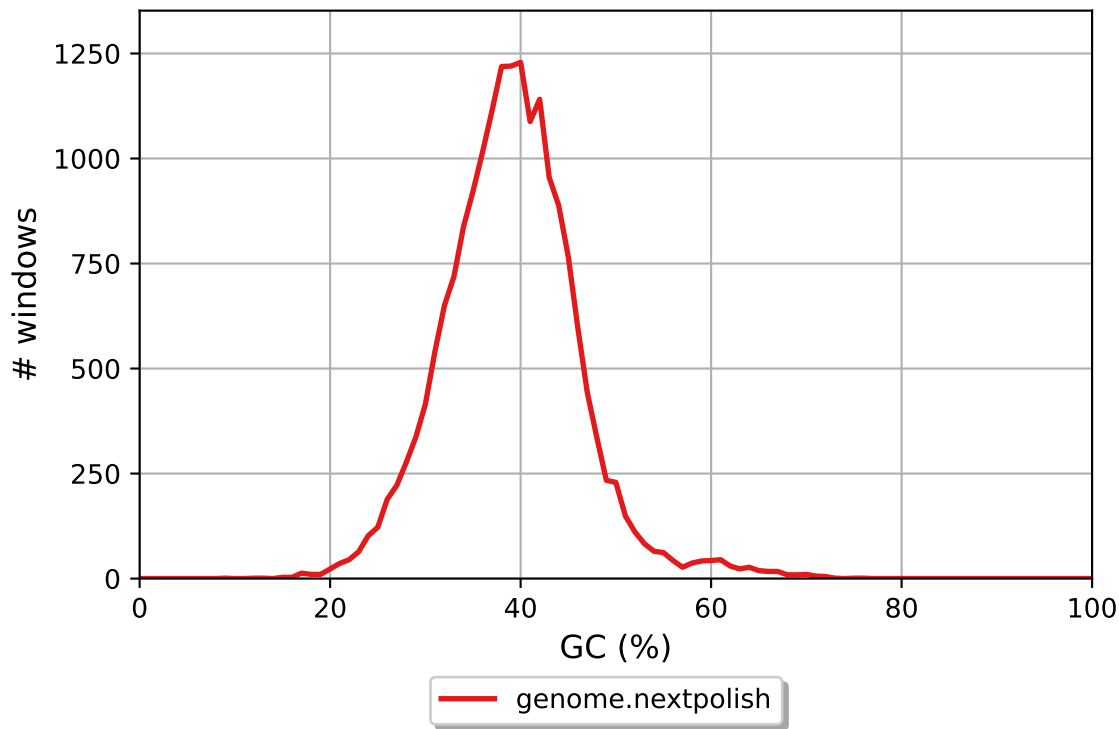
	genome.nextpolish
# contigs (>= 0 bp)	28
# contigs (>= 1000 bp)	28
# contigs (>= 5000 bp)	6
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	1889888
Total length (>= 1000 bp)	1889888
Total length (>= 5000 bp)	1832731
Total length (>= 10000 bp)	1827317
Total length (>= 25000 bp)	1827317
Total length (>= 50000 bp)	1827317
# contigs	28
Largest contig	1020198
Total length	1889888
GC (%)	39.06
N50	1020198
N90	210801
auN	653263.6
L50	1
L90	4
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 2 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).





GC content



genome.nextpolish GC content

