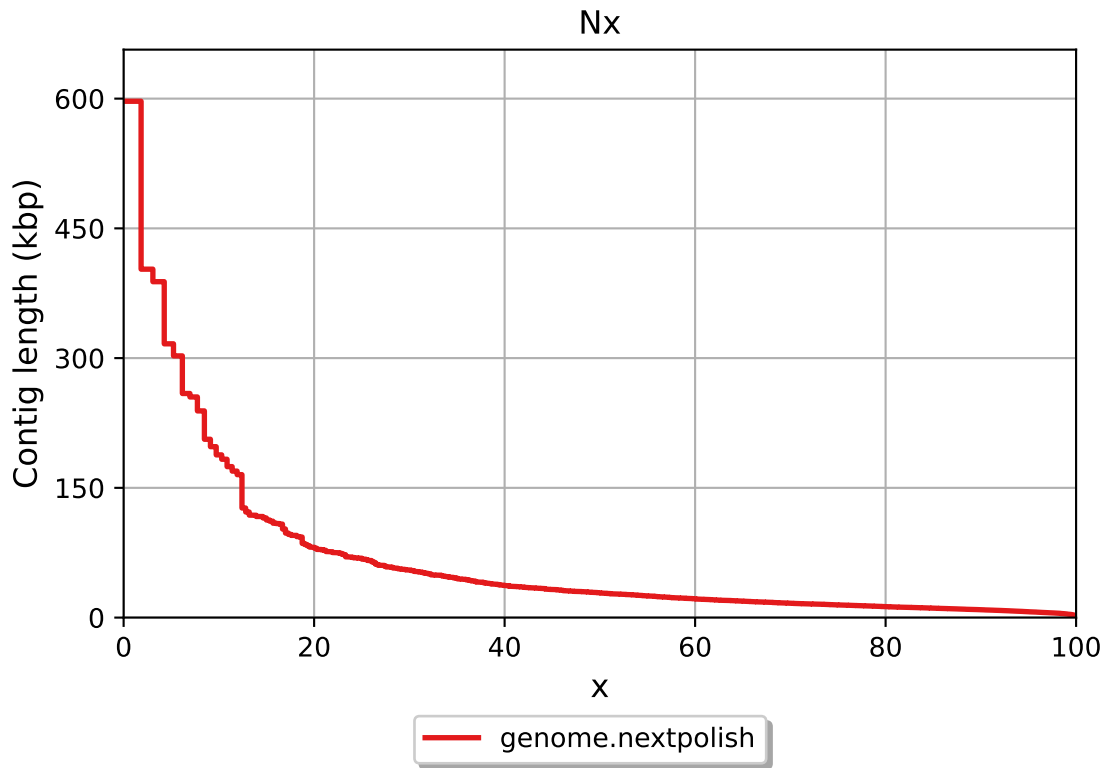


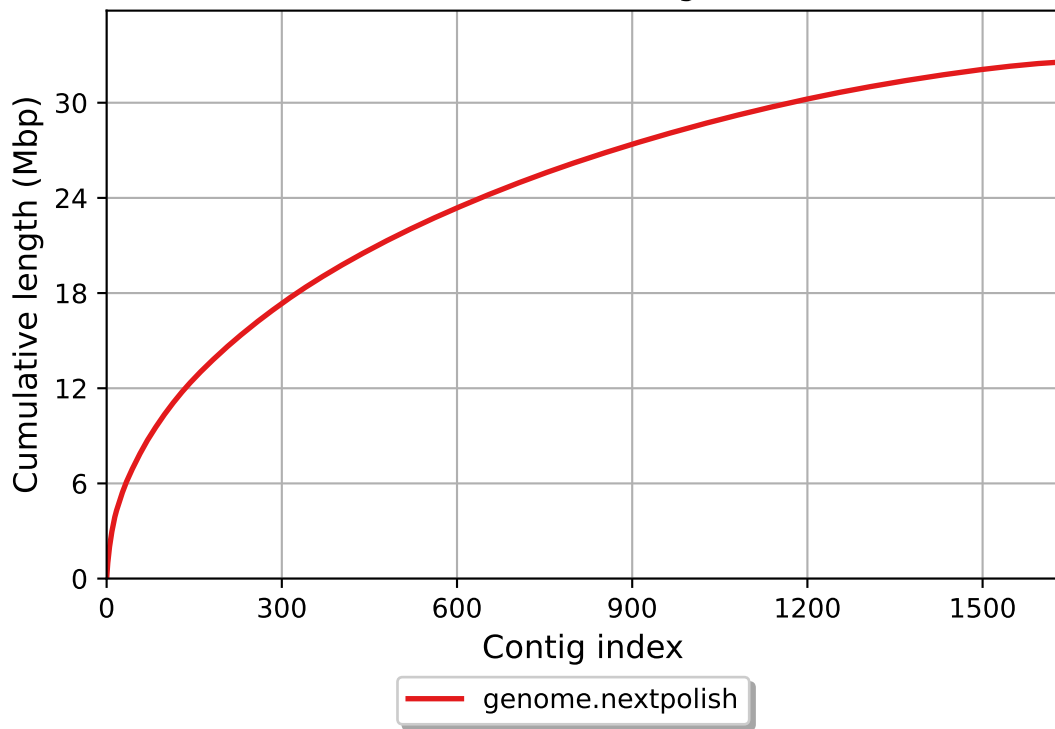
Report

	genome.nextpolish
# contigs (>= 0 bp)	1631
# contigs (>= 1000 bp)	1625
# contigs (>= 5000 bp)	1474
# contigs (>= 10000 bp)	1006
# contigs (>= 25000 bp)	322
# contigs (>= 50000 bp)	102
Total length (>= 0 bp)	32551005
Total length (>= 1000 bp)	32546101
Total length (>= 5000 bp)	31976756
Total length (>= 10000 bp)	28506852
Total length (>= 25000 bp)	17909843
Total length (>= 50000 bp)	10499401
# contigs	1631
Largest contig	596940
Total length	32551005
GC (%)	51.33
N50	28636
N90	9123
auN	69650.8
L50	261
L90	1089
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	65 + 4 part

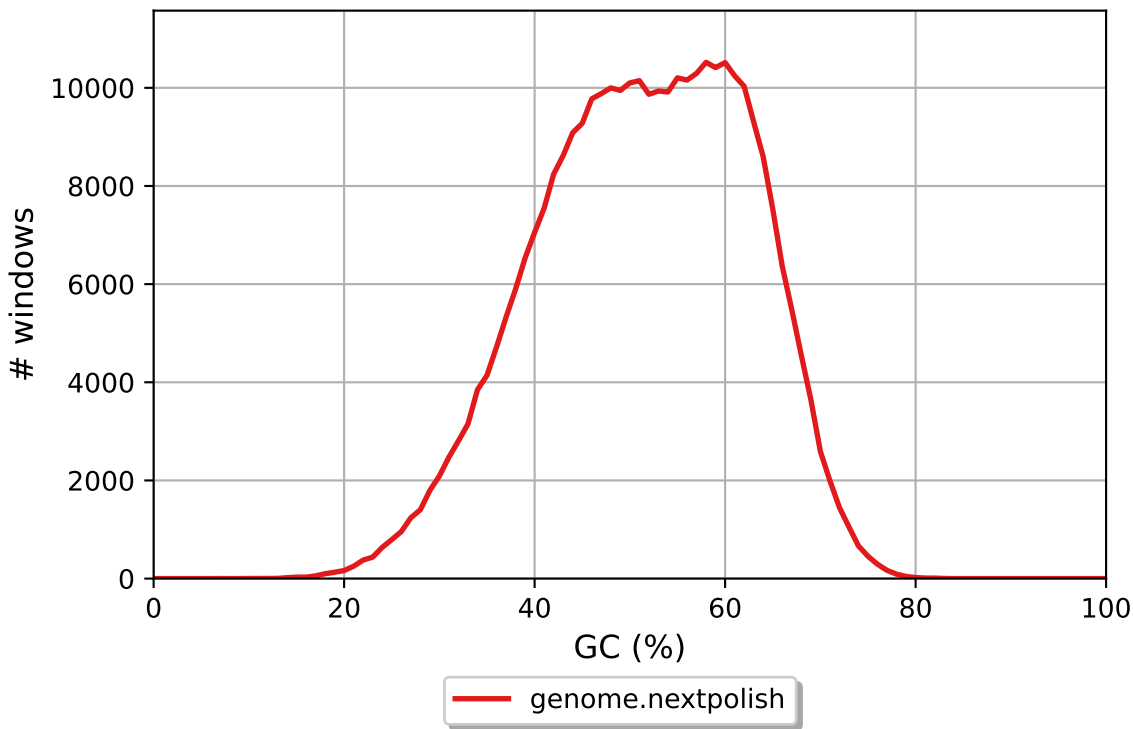
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

