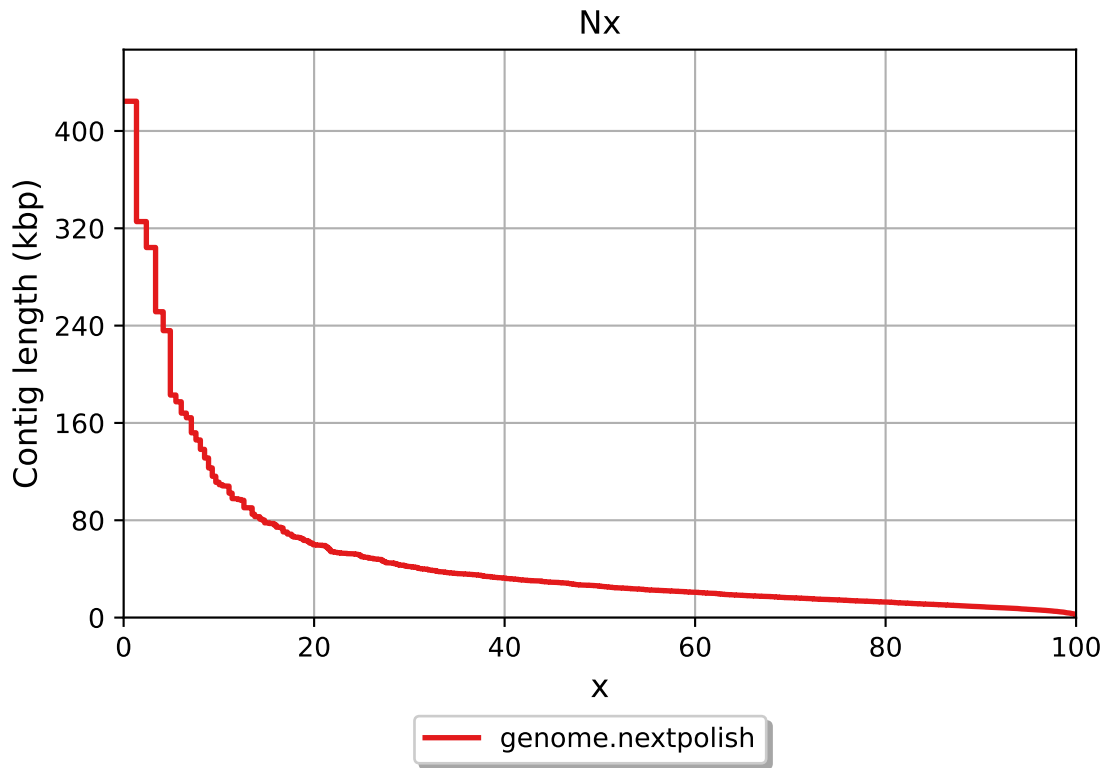


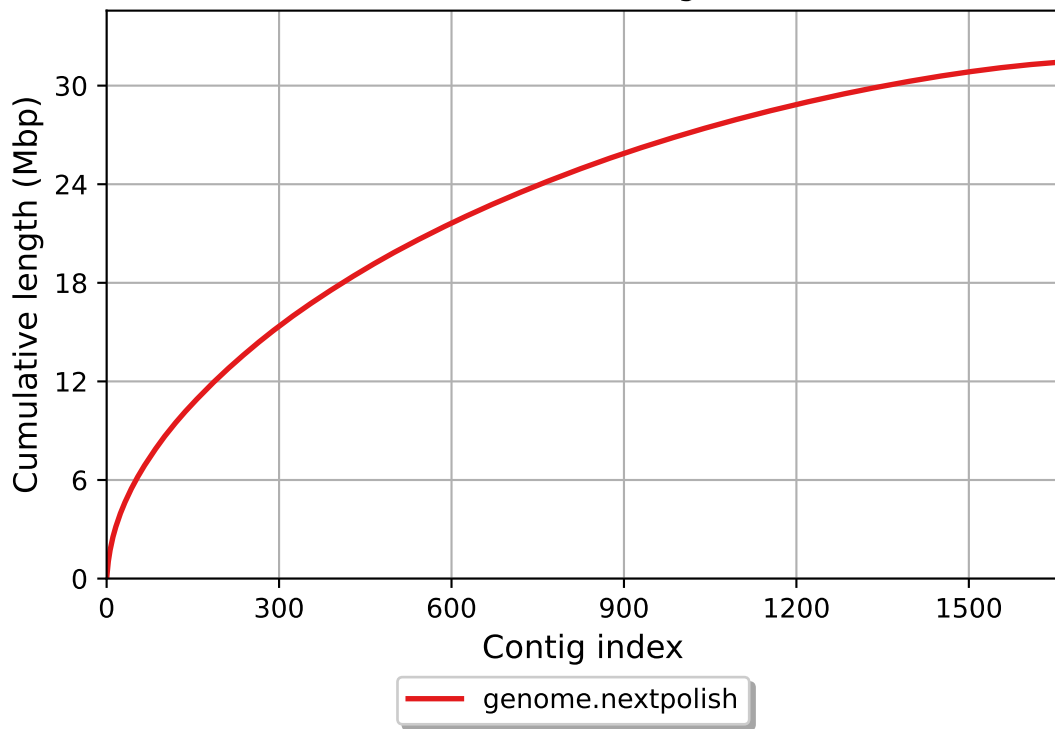
Report

	genome.nextpolish
# contigs (>= 0 bp)	1657
# contigs (>= 1000 bp)	1657
# contigs (>= 5000 bp)	1495
# contigs (>= 10000 bp)	1041
# contigs (>= 25000 bp)	327
# contigs (>= 50000 bp)	85
Total length (>= 0 bp)	31421804
Total length (>= 1000 bp)	31421804
Total length (>= 5000 bp)	30813917
Total length (>= 10000 bp)	27411659
Total length (>= 25000 bp)	16056044
Total length (>= 50000 bp)	7886134
# contigs	1657
Largest contig	424408
Total length	31421804
GC (%)	51.83
N50	25909
N90	8911
auN	50607.4
L50	314
L90	1133
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	63 + 8 part

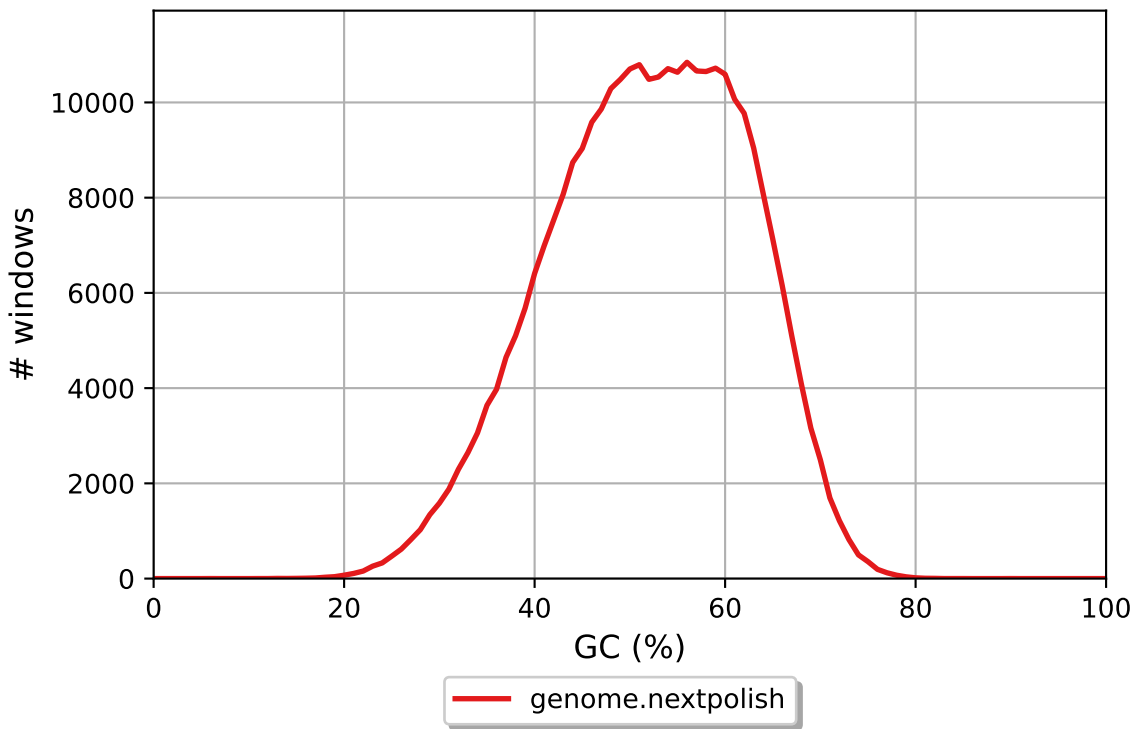
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

