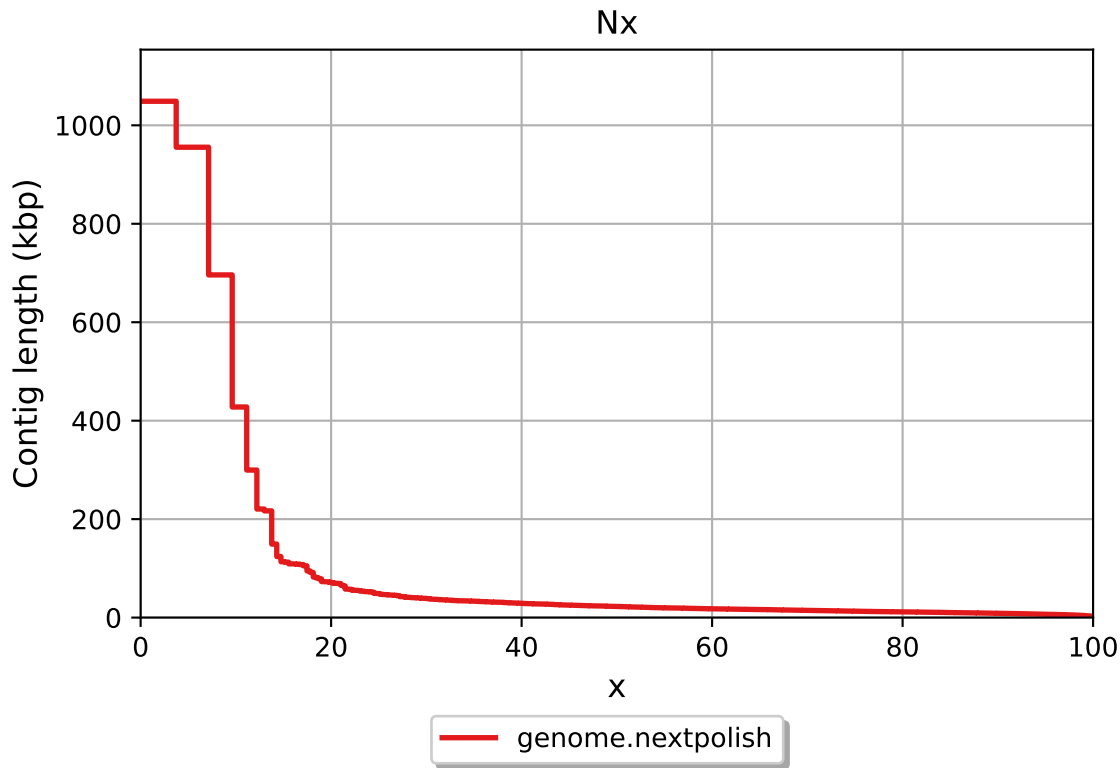


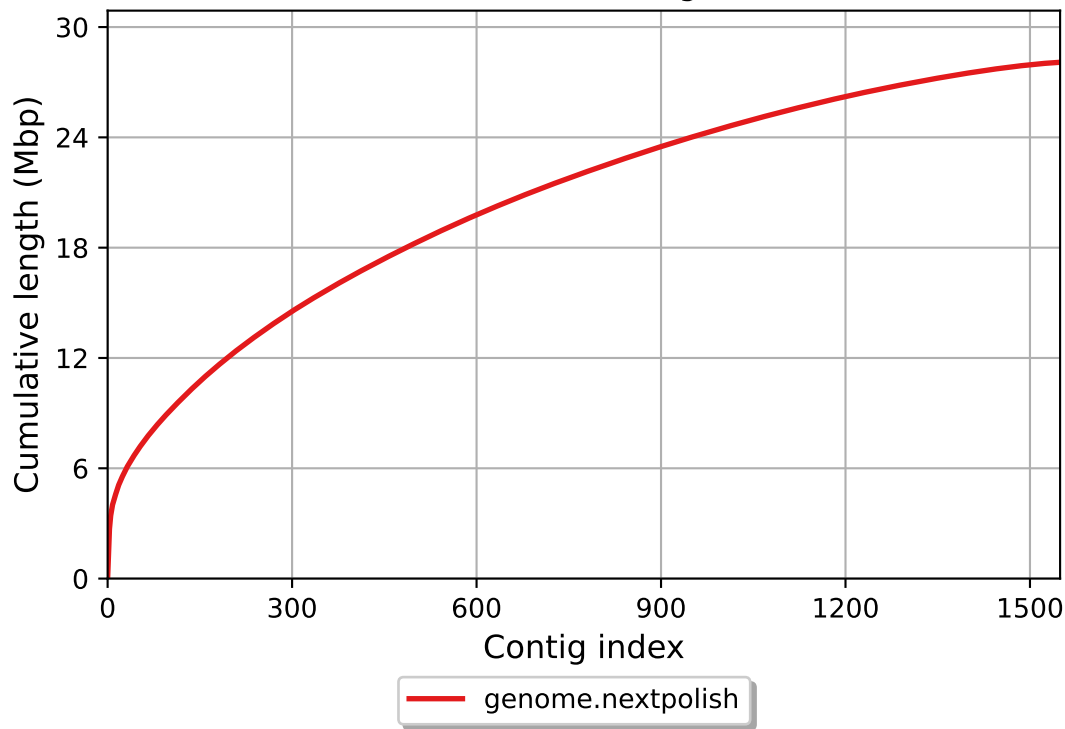
Report

	genome.nextpolish
# contigs (>= 0 bp)	1549
# contigs (>= 1000 bp)	1547
# contigs (>= 5000 bp)	1420
# contigs (>= 10000 bp)	954
# contigs (>= 25000 bp)	226
# contigs (>= 50000 bp)	47
Total length (>= 0 bp)	28084899
Total length (>= 1000 bp)	28083253
Total length (>= 5000 bp)	27605179
Total length (>= 10000 bp)	24055497
Total length (>= 25000 bp)	12821007
Total length (>= 50000 bp)	6902073
# contigs	1549
Largest contig	1048847
Total length	28084899
GC (%)	52.12
N50	22523
N90	8725
auN	125981.9
L50	278
L90	1085
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	73 + 8 part

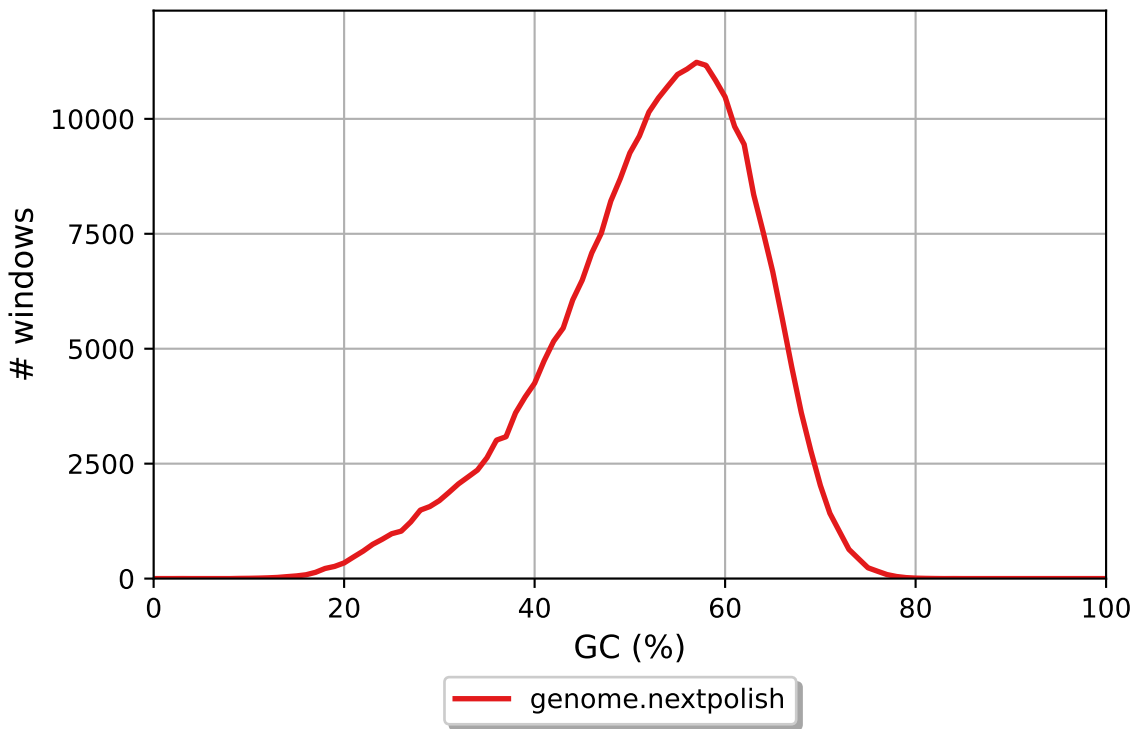
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

