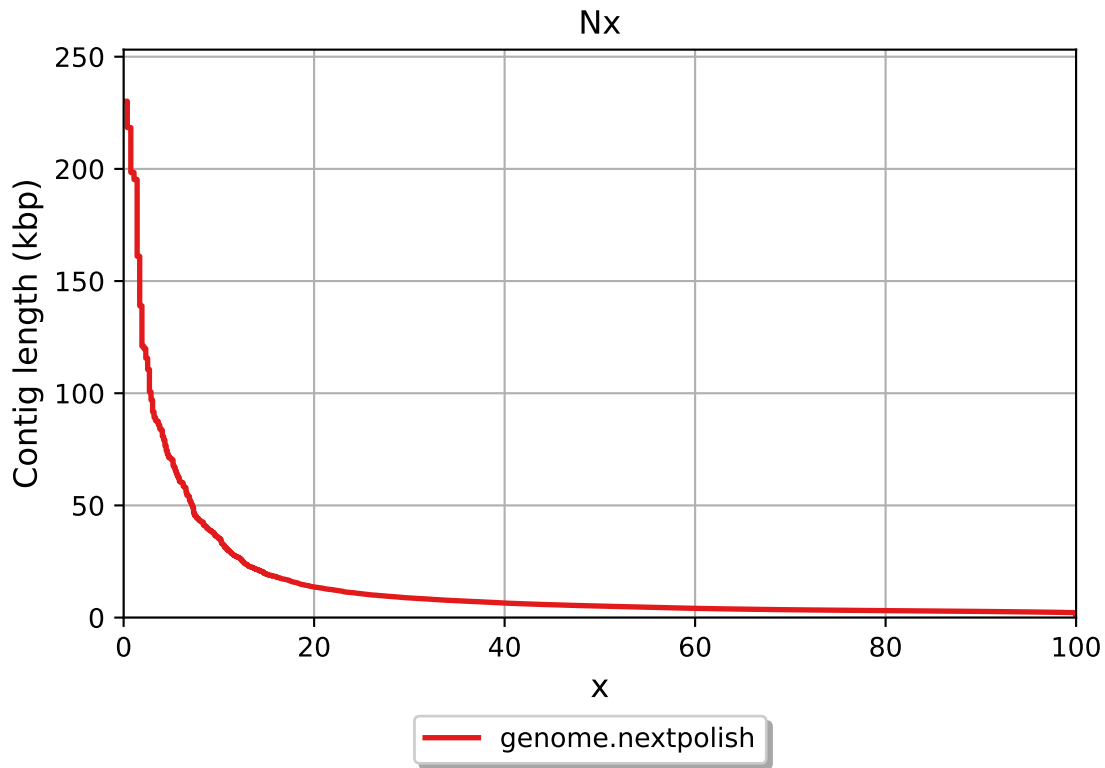


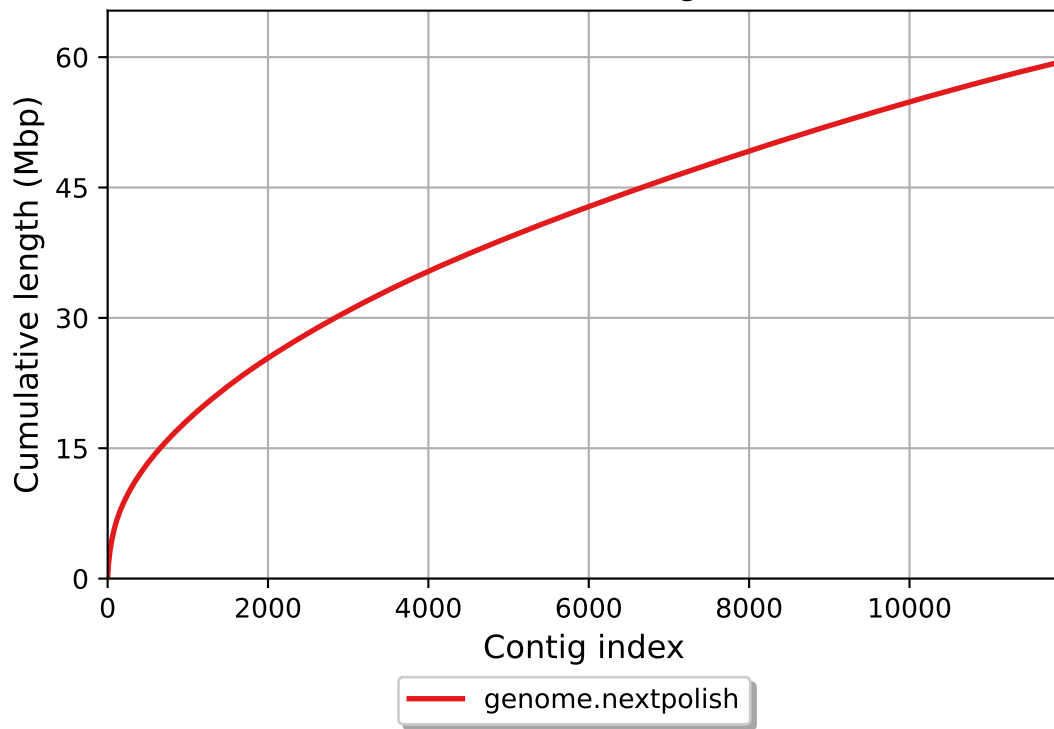
Report

	genome.nextpolish
# contigs (>= 0 bp)	11878
# contigs (>= 1000 bp)	11871
# contigs (>= 5000 bp)	2931
# contigs (>= 10000 bp)	719
# contigs (>= 25000 bp)	141
# contigs (>= 50000 bp)	48
Total length (>= 0 bp)	59411923
Total length (>= 1000 bp)	59406800
Total length (>= 5000 bp)	30476772
Total length (>= 10000 bp)	15670352
Total length (>= 25000 bp)	7455813
Total length (>= 50000 bp)	4262904
# contigs	11878
Largest contig	230138
Total length	59411923
GC (%)	38.11
N50	5144
N90	2744
auN	15309.2
L50	2779
L90	9488
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 8 part

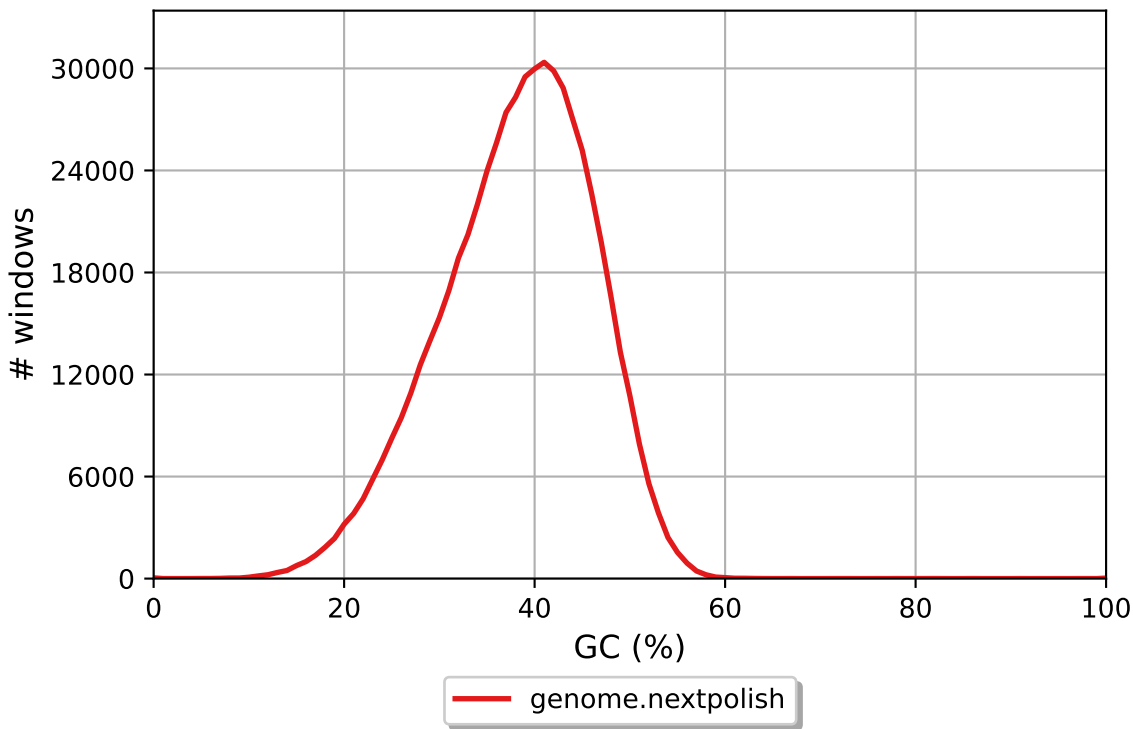
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

