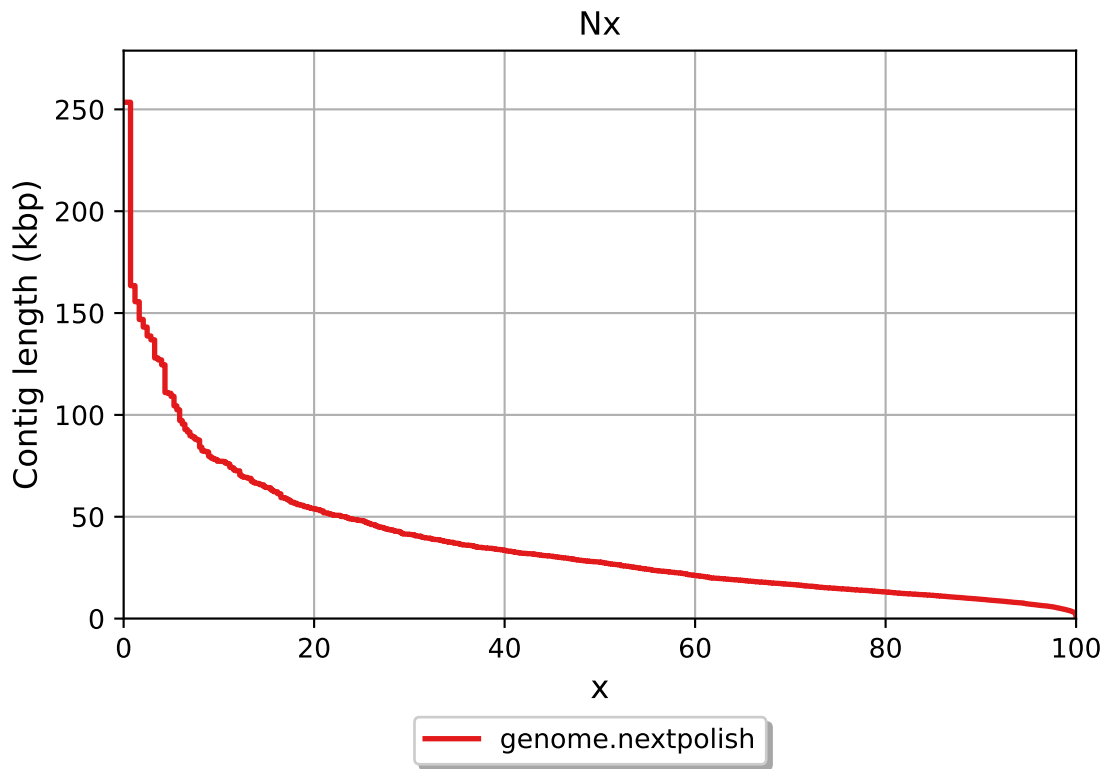


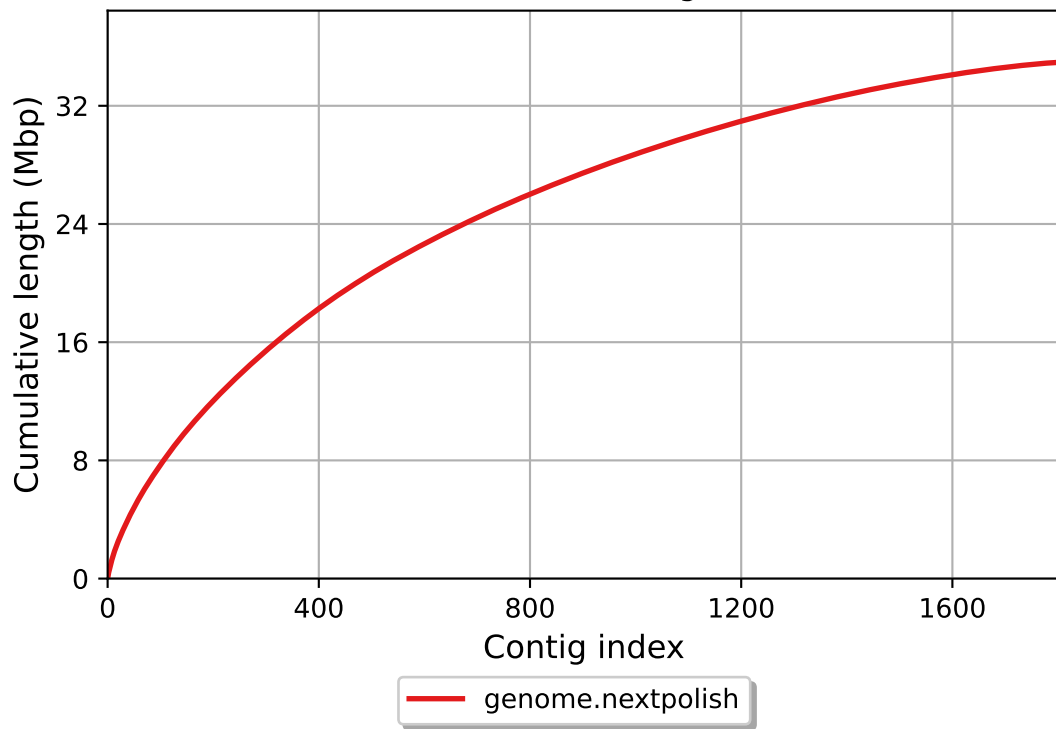
Report

	genome.nextpolish
# contigs (>= 0 bp)	1804
# contigs (>= 1000 bp)	1802
# contigs (>= 5000 bp)	1647
# contigs (>= 10000 bp)	1199
# contigs (>= 25000 bp)	422
# contigs (>= 50000 bp)	107
Total length (>= 0 bp)	34945441
Total length (>= 1000 bp)	34943562
Total length (>= 5000 bp)	34357135
Total length (>= 10000 bp)	30950030
Total length (>= 25000 bp)	18836370
Total length (>= 50000 bp)	8047053
# contigs	1804
Largest contig	253483
Total length	34945441
GC (%)	50.49
N50	27740
N90	9467
auN	37749.6
L50	371
L90	1251
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	64 + 8 part

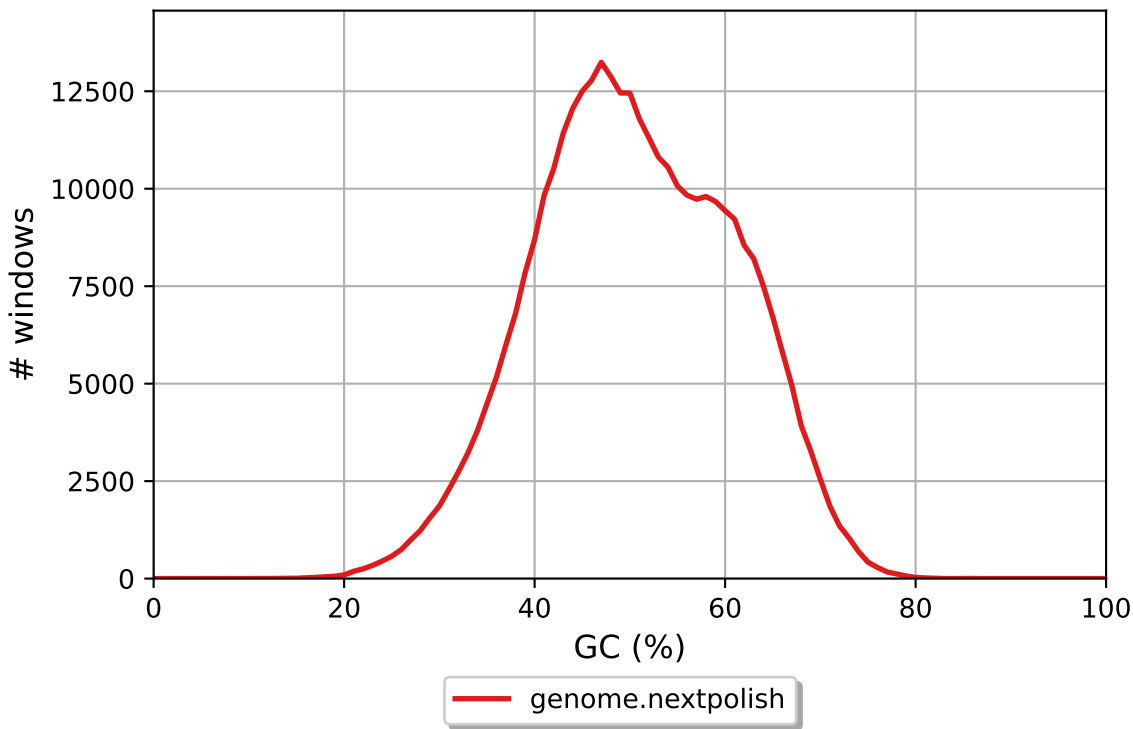
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

