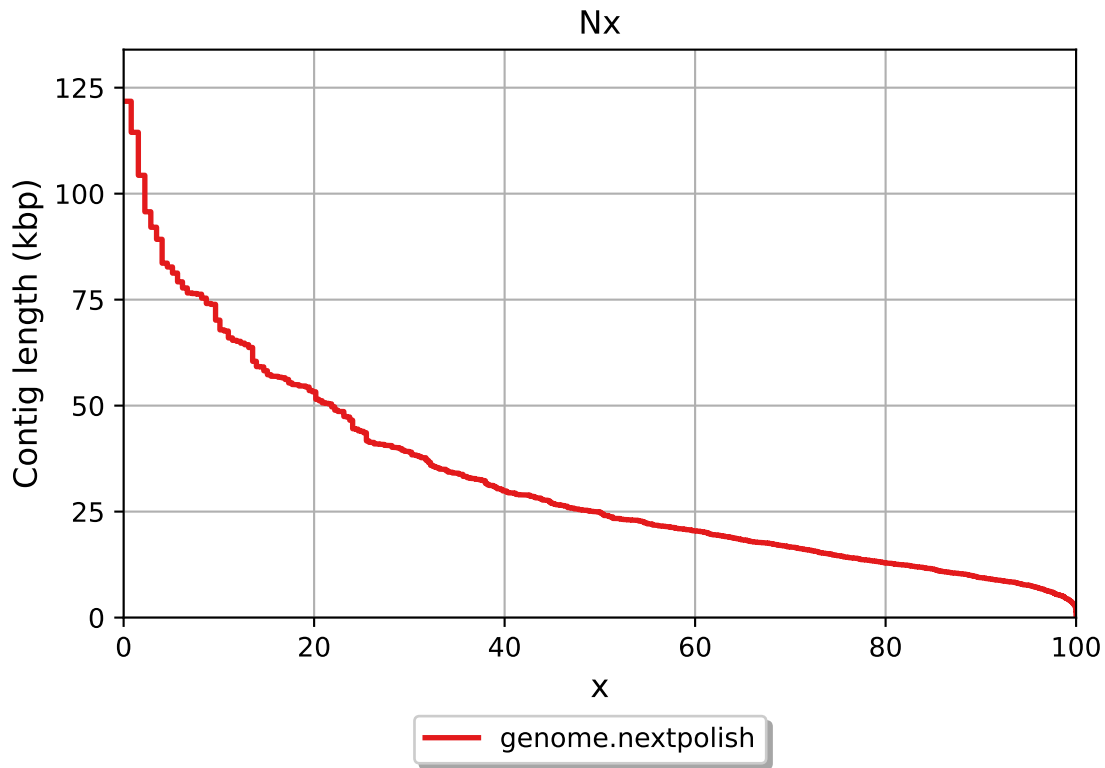


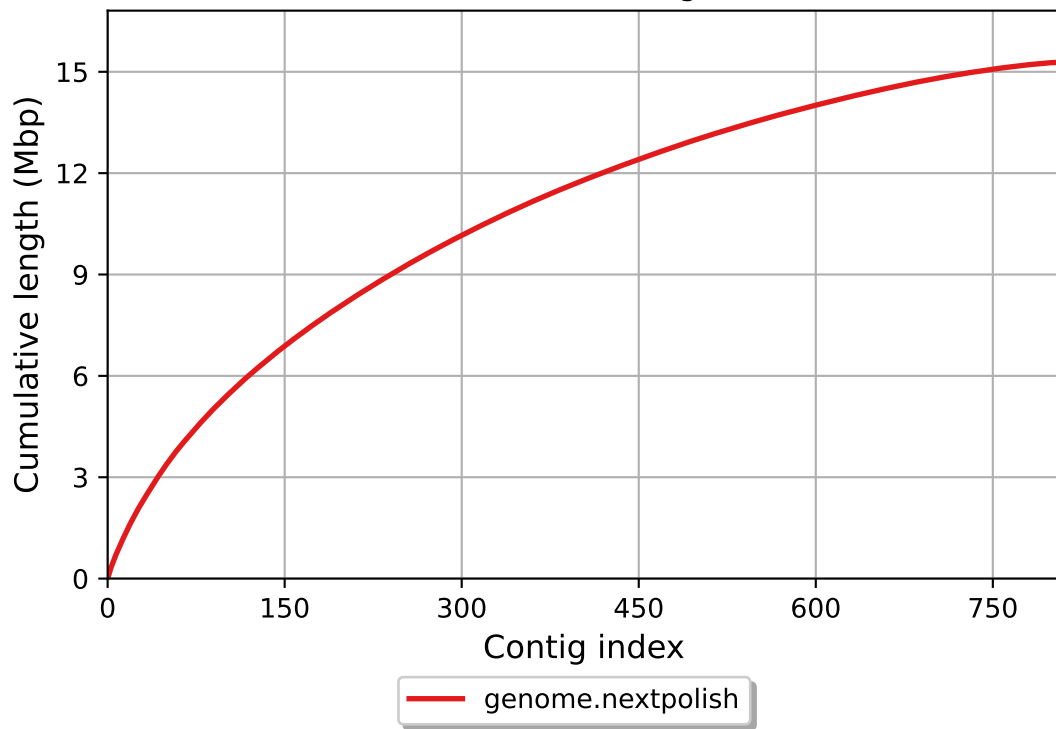
Report

	genome.nextpolish
# contigs (>= 0 bp)	807
# contigs (>= 1000 bp)	805
# contigs (>= 5000 bp)	750
# contigs (>= 10000 bp)	553
# contigs (>= 25000 bp)	176
# contigs (>= 50000 bp)	49
Total length (>= 0 bp)	15281814
Total length (>= 1000 bp)	15280253
Total length (>= 5000 bp)	15077606
Total length (>= 10000 bp)	13571491
Total length (>= 25000 bp)	7556217
Total length (>= 50000 bp)	3337255
# contigs	807
Largest contig	121793
Total length	15281814
GC (%)	44.63
N50	24856
N90	9439
auN	32810.1
L50	180
L90	572
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	32 + 5 part

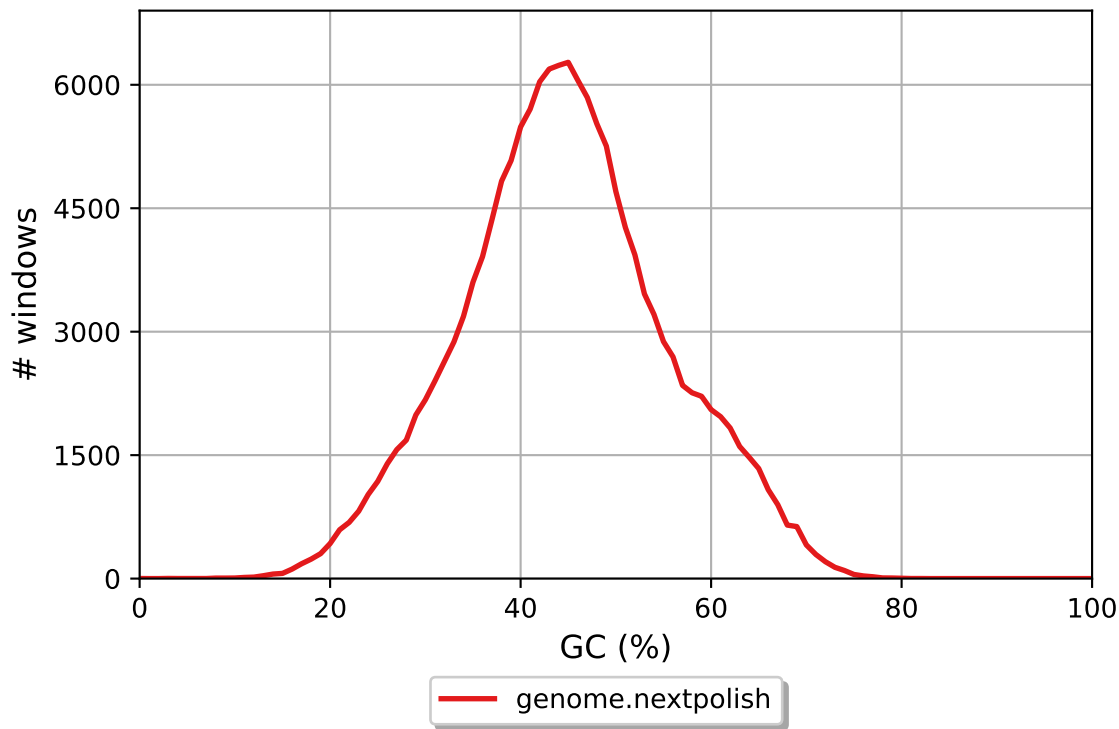
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

