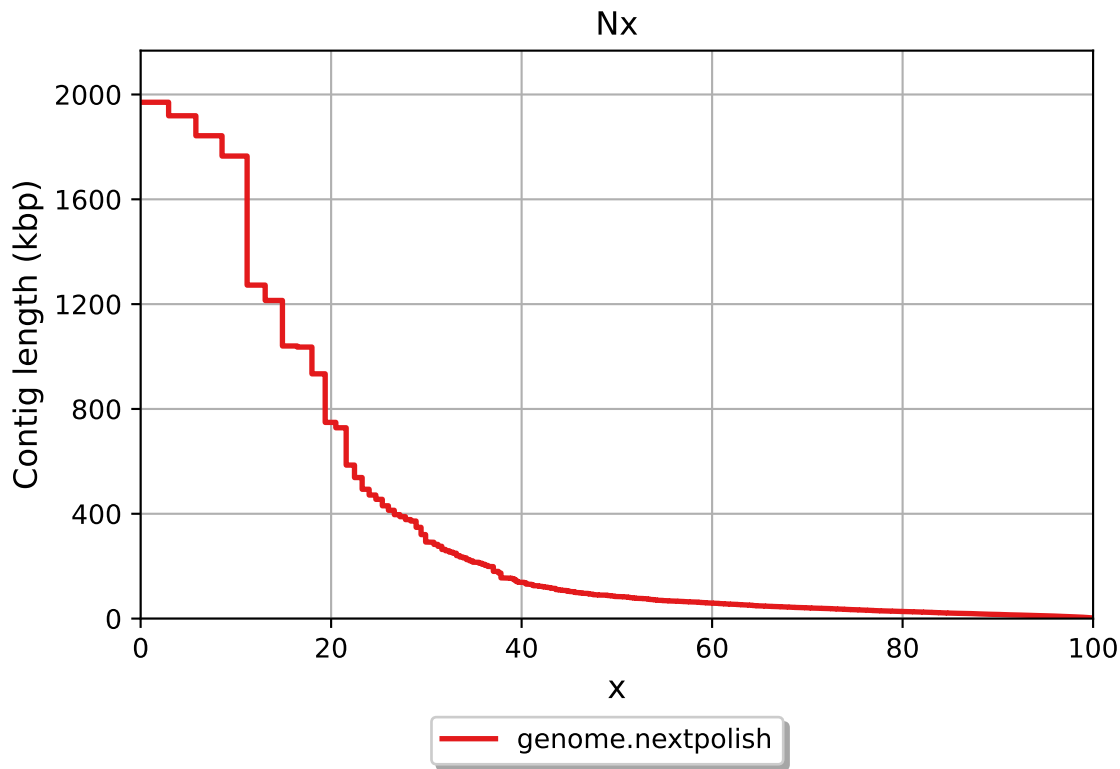


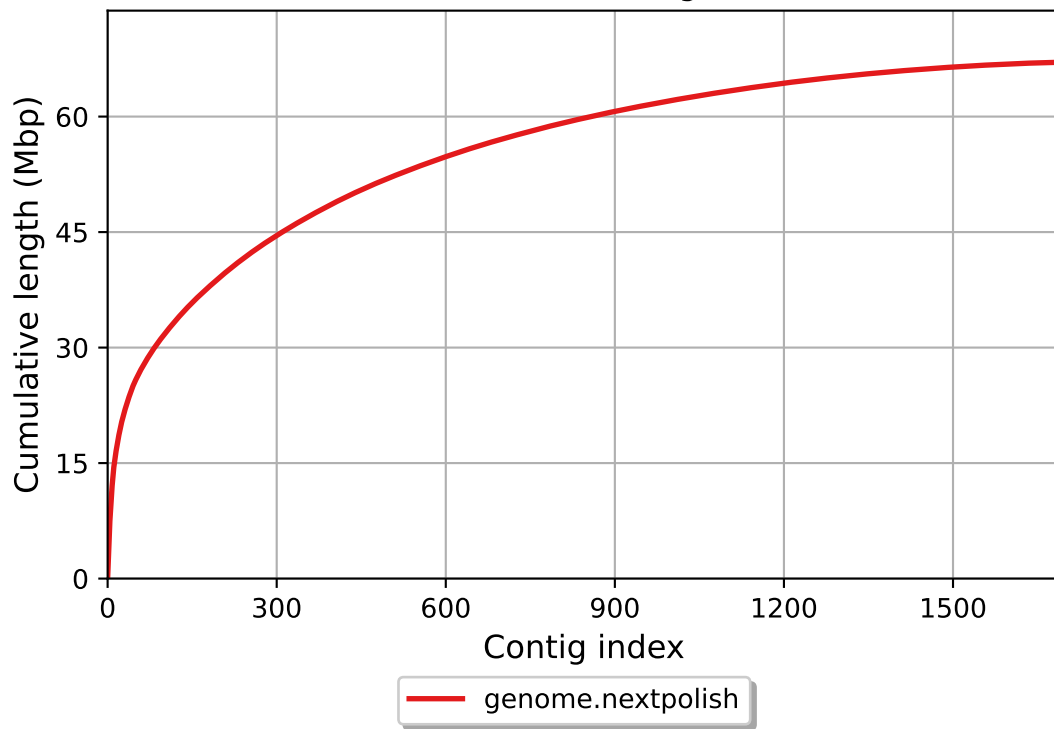
Report

	genome.nextpolish
# contigs (>= 0 bp)	1690
# contigs (>= 1000 bp)	1686
# contigs (>= 5000 bp)	1486
# contigs (>= 10000 bp)	1172
# contigs (>= 25000 bp)	607
# contigs (>= 50000 bp)	270
Total length (>= 0 bp)	67053458
Total length (>= 1000 bp)	67050693
Total length (>= 5000 bp)	66365582
Total length (>= 10000 bp)	64063430
Total length (>= 25000 bp)	54968455
Total length (>= 50000 bp)	43117556
# contigs	1690
Largest contig	1970334
Total length	67053458
GC (%)	53.55
N50	84260
N90	15817
auN	405455.2
L50	121
L90	879
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	143 + 2 part

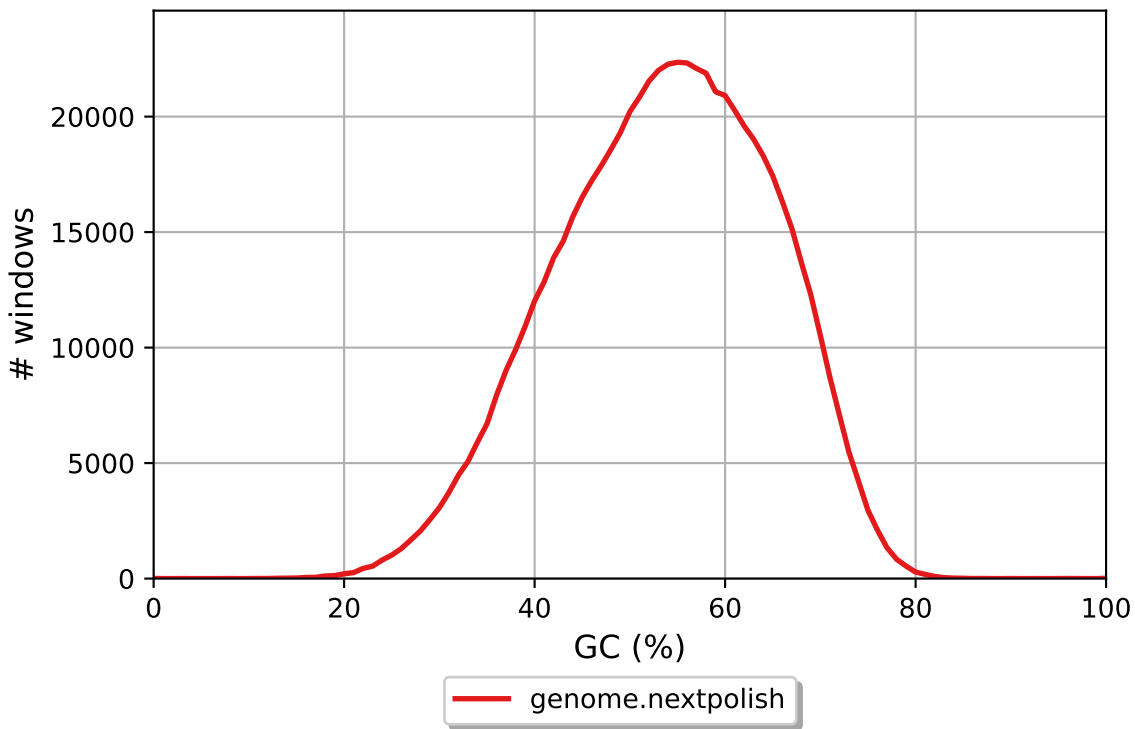
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

