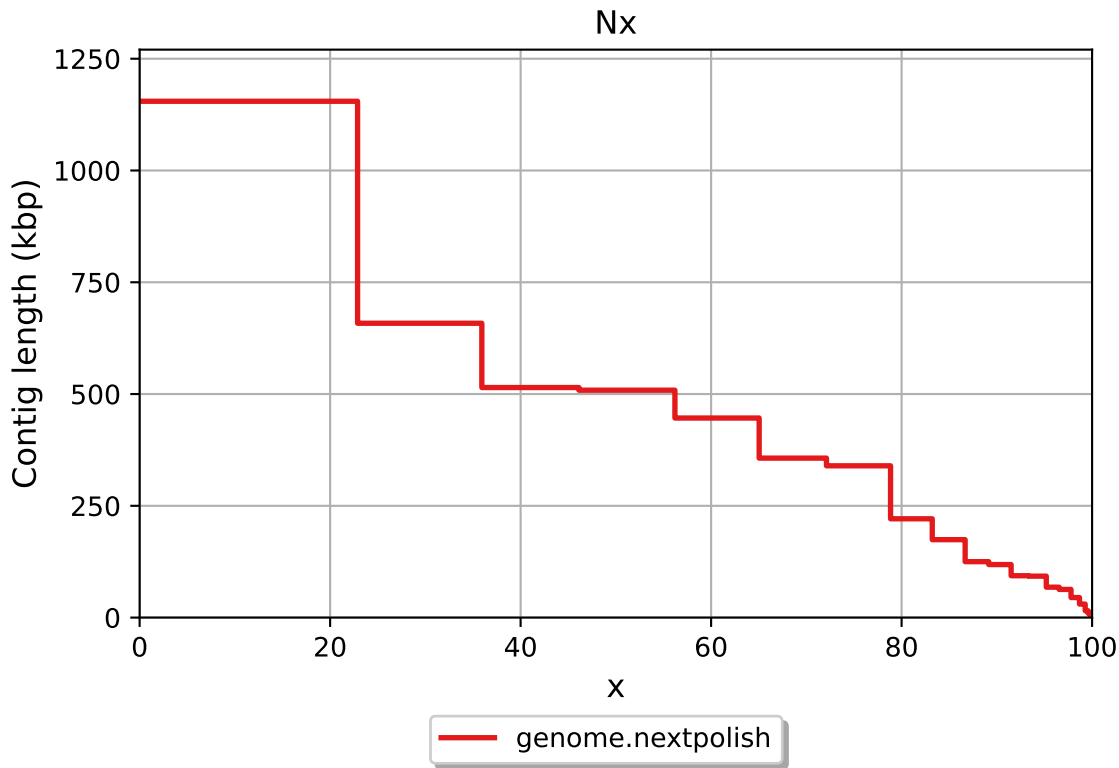


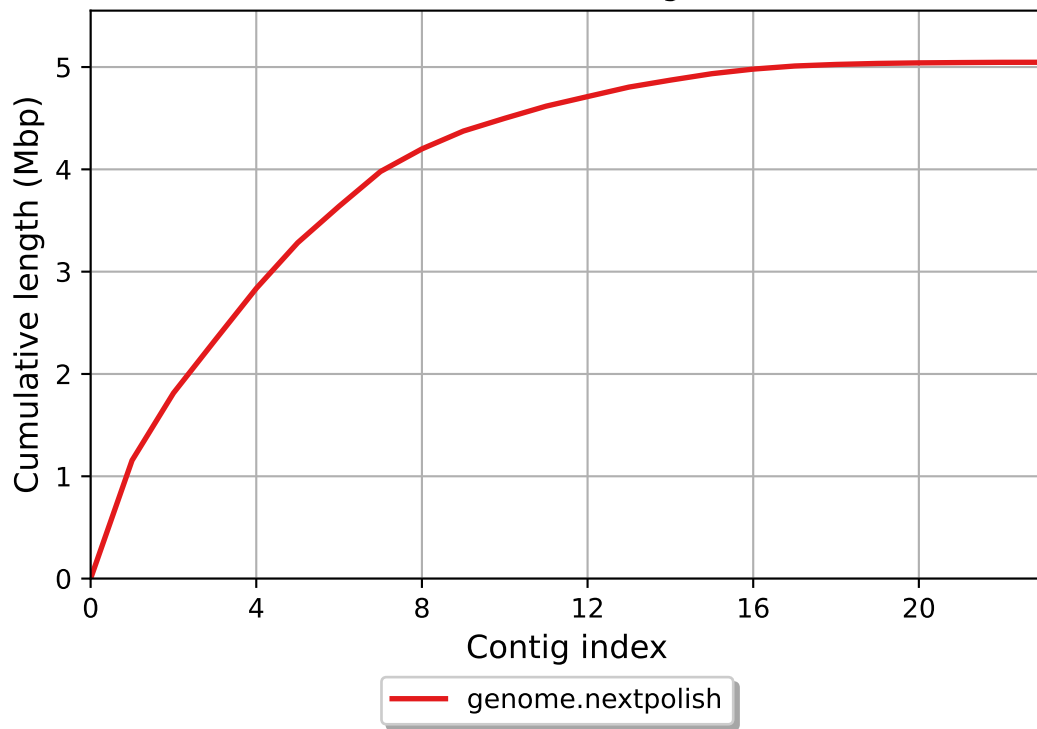
Report

	genome.nextpolish
# contigs (>= 0 bp)	23
# contigs (>= 1000 bp)	22
# contigs (>= 5000 bp)	20
# contigs (>= 10000 bp)	18
# contigs (>= 25000 bp)	17
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	5046680
Total length (>= 1000 bp)	5046128
Total length (>= 5000 bp)	5040763
Total length (>= 10000 bp)	5025406
Total length (>= 25000 bp)	5010080
Total length (>= 50000 bp)	4935000
# contigs	23
Largest contig	1154831
Total length	5046680
GC (%)	55.19
N50	508567
N90	118603
auN	568746.9
L50	4
L90	11
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 0 part

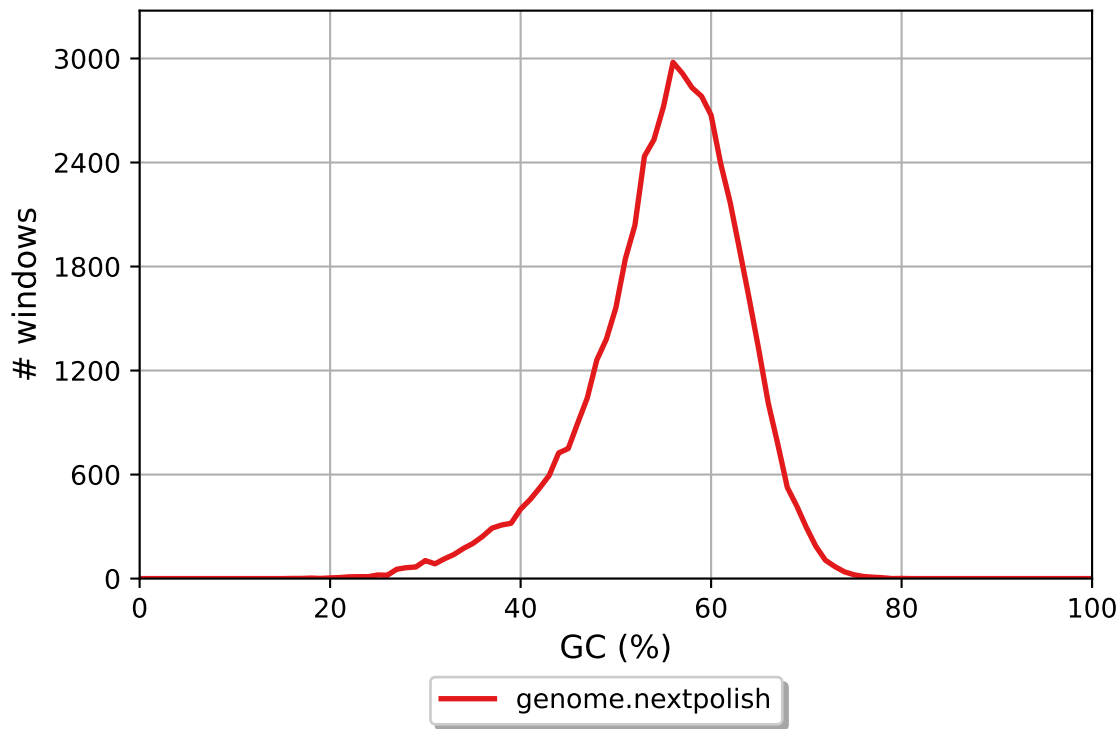
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

