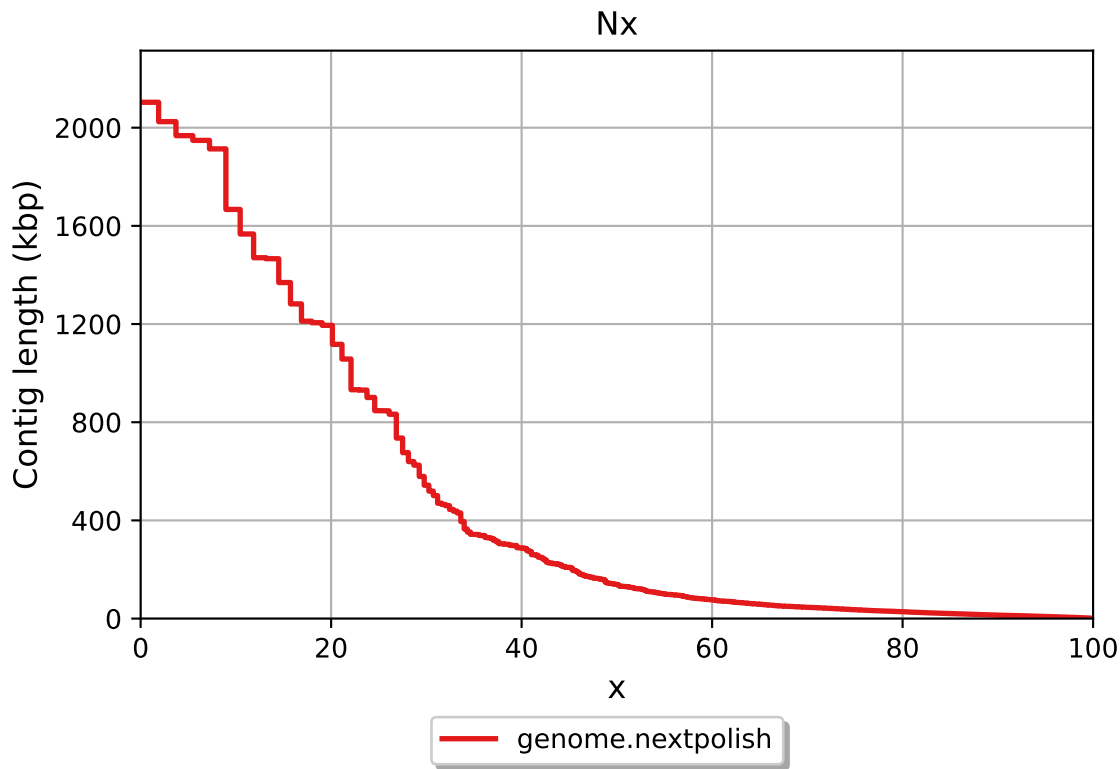


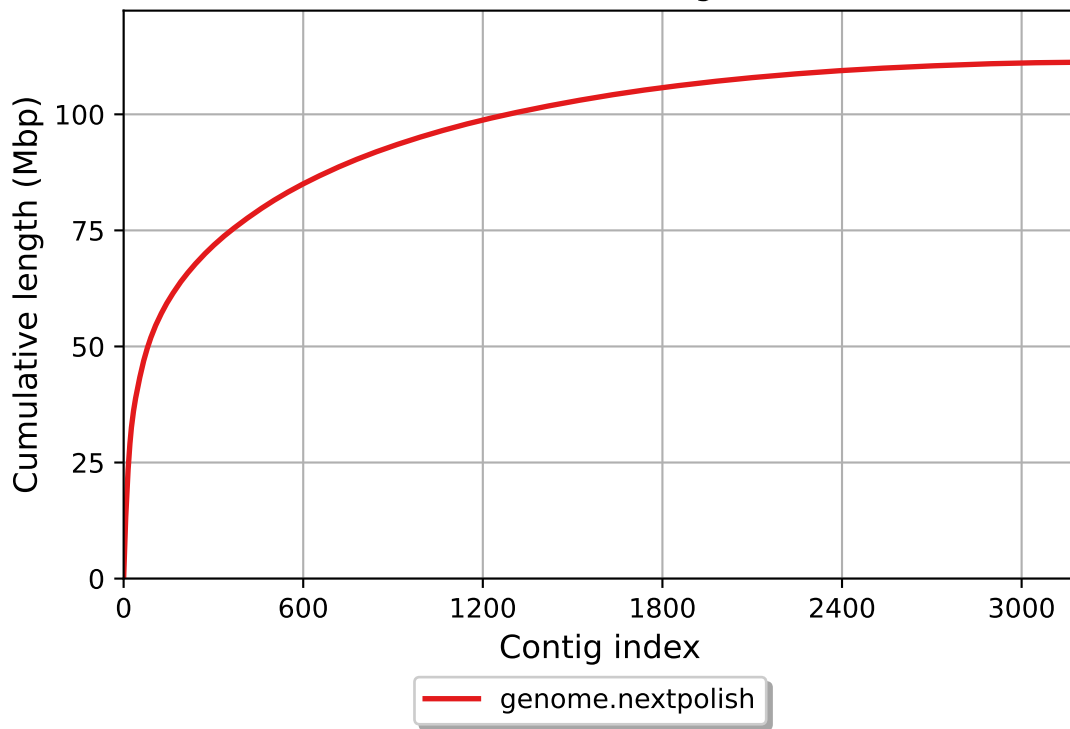
Report

	genome.nextpolish
# contigs (>= 0 bp)	3188
# contigs (>= 1000 bp)	3051
# contigs (>= 5000 bp)	2263
# contigs (>= 10000 bp)	1641
# contigs (>= 25000 bp)	804
# contigs (>= 50000 bp)	369
Total length (>= 0 bp)	111215626
Total length (>= 1000 bp)	111119304
Total length (>= 5000 bp)	108803618
Total length (>= 10000 bp)	104297576
Total length (>= 25000 bp)	90947179
Total length (>= 50000 bp)	75499047
# contigs	3182
Largest contig	2103594
Total length	111213701
GC (%)	47.64
N50	139360
N90	13926
auN	500069.5
L50	115
L90	1290
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	244 + 12 part

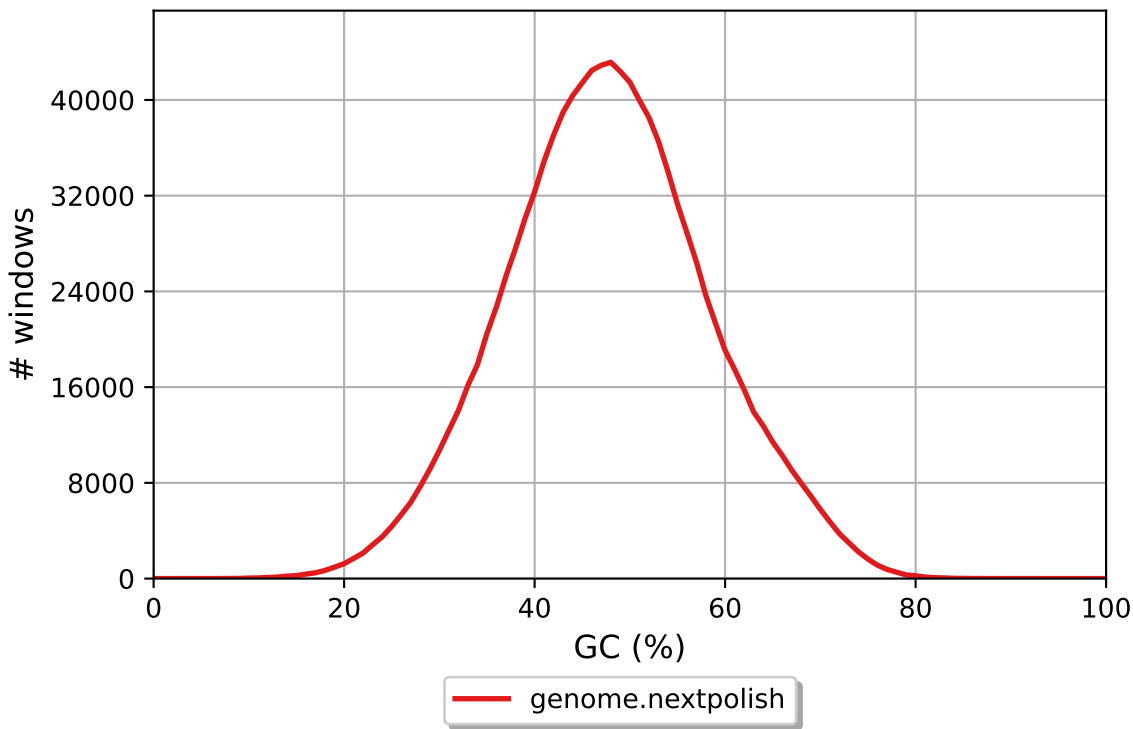
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

