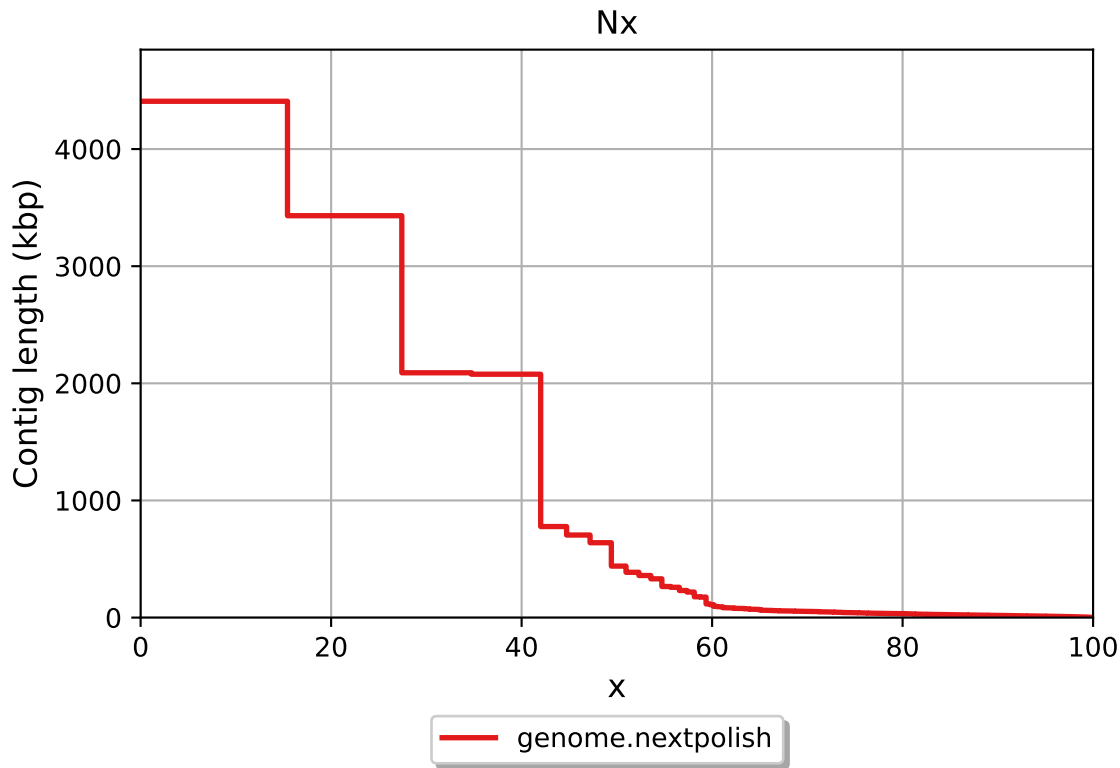


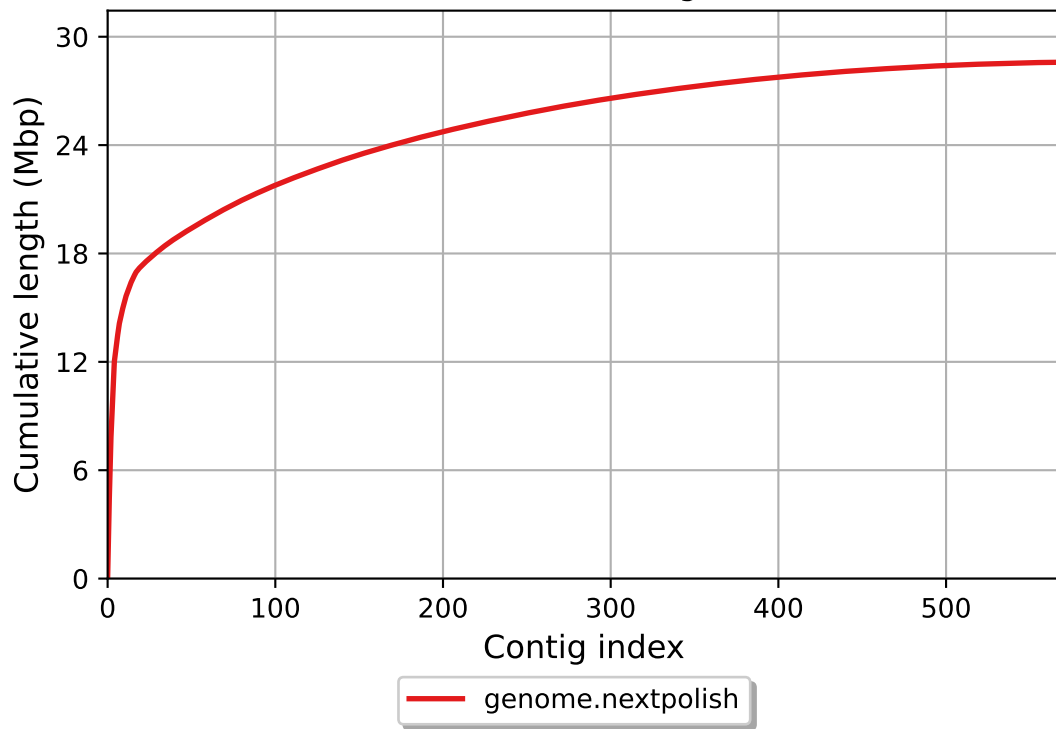
Report

	genome.nextpolish
# contigs (>= 0 bp)	568
# contigs (>= 1000 bp)	564
# contigs (>= 5000 bp)	483
# contigs (>= 10000 bp)	380
# contigs (>= 25000 bp)	180
# contigs (>= 50000 bp)	68
Total length (>= 0 bp)	28586460
Total length (>= 1000 bp)	28583669
Total length (>= 5000 bp)	28331345
Total length (>= 10000 bp)	27576403
Total length (>= 25000 bp)	24270607
Total length (>= 50000 bp)	20367523
# contigs	568
Largest contig	4408013
Total length	28586460
GC (%)	40.68
N50	439656
N90	18878
auN	1494537.3
L50	8
L90	248
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	97 + 5 part

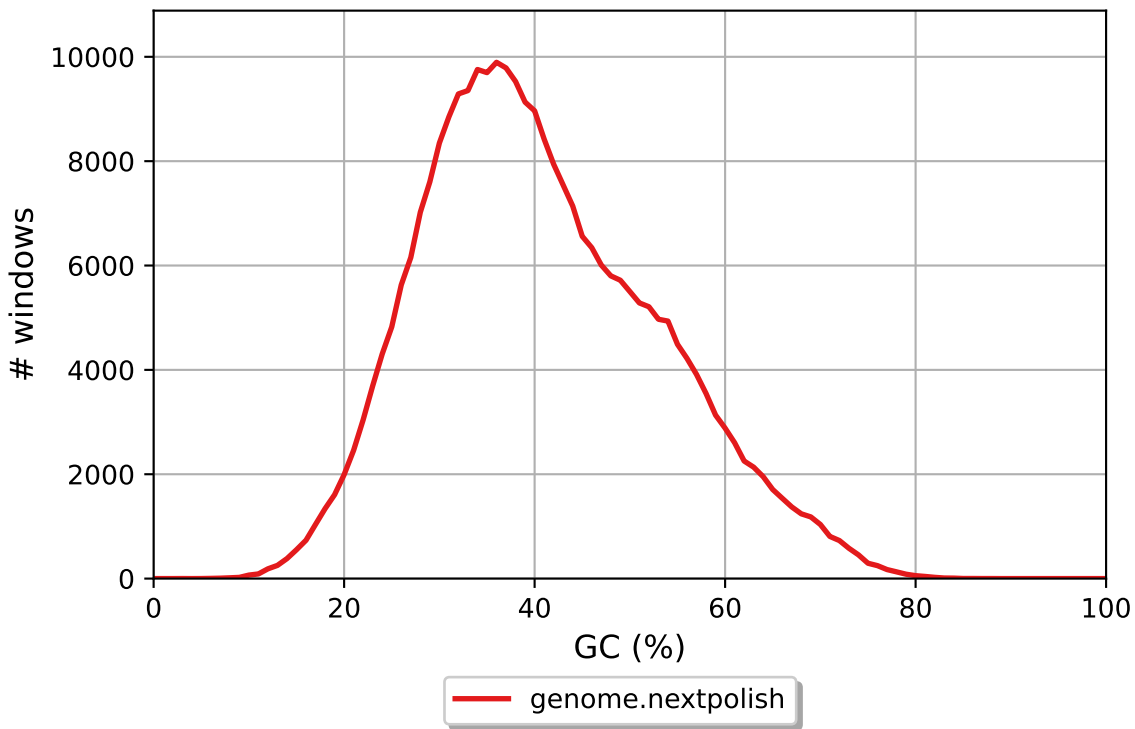
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

