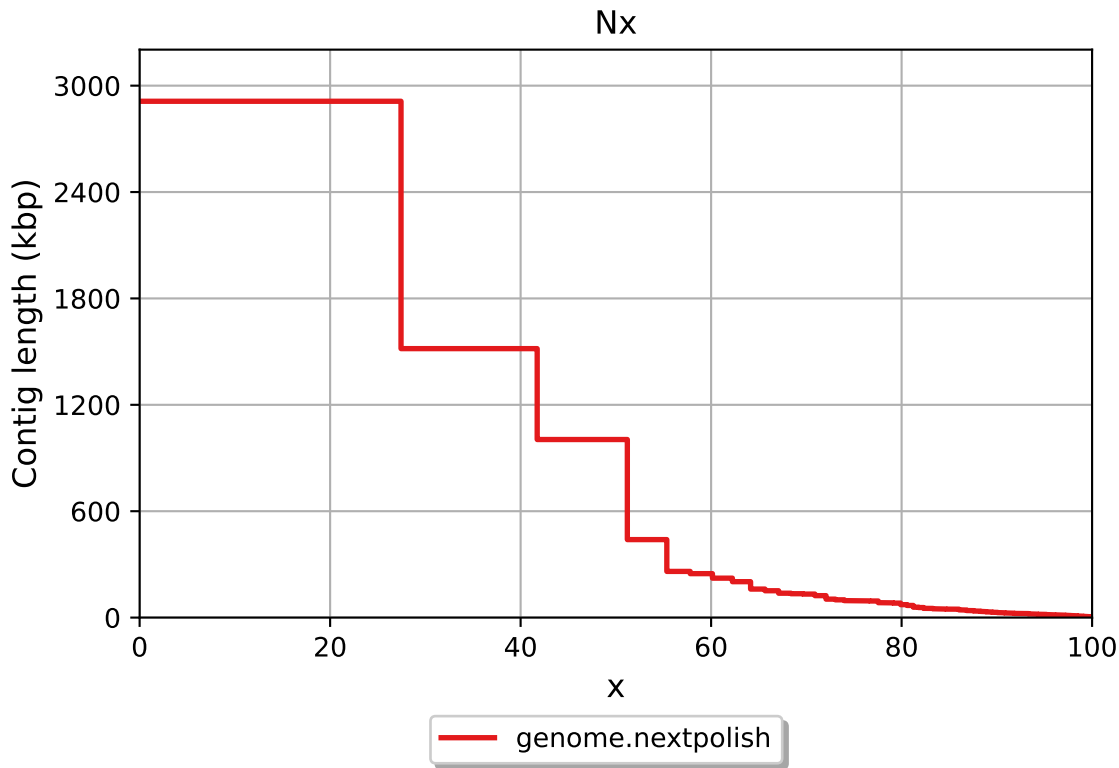


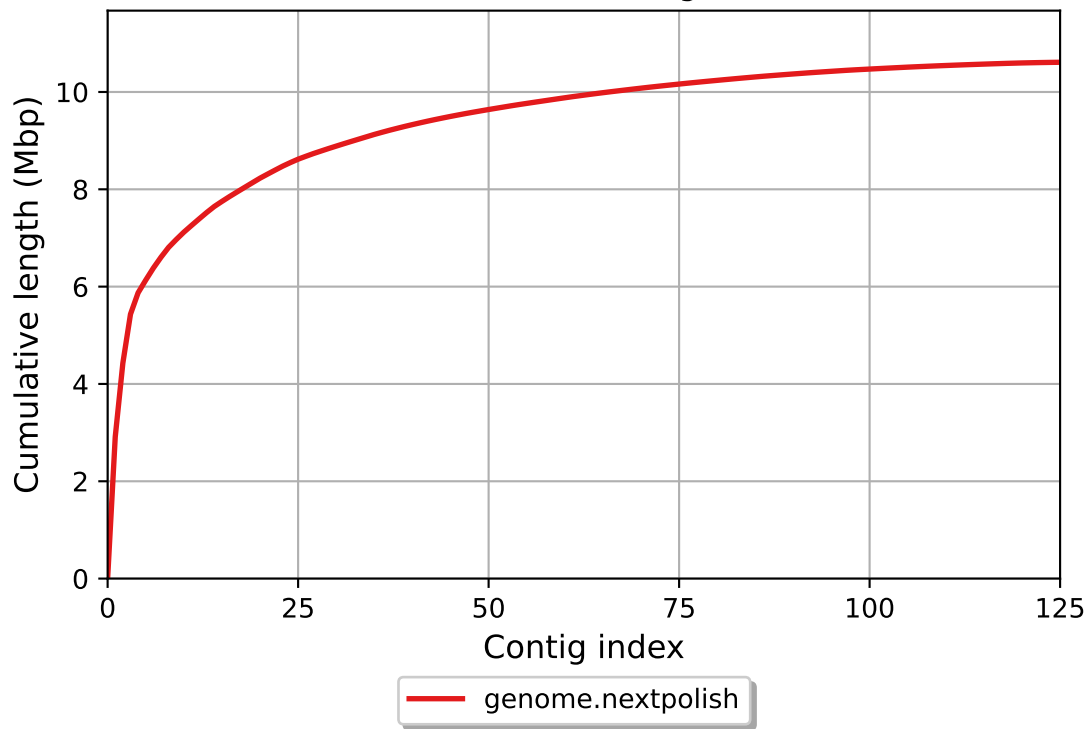
# Report

	genome.nextpolish
# contigs (>= 0 bp)	125
# contigs (>= 1000 bp)	125
# contigs (>= 5000 bp)	118
# contigs (>= 10000 bp)	97
# contigs (>= 25000 bp)	53
# contigs (>= 50000 bp)	29
Total length (>= 0 bp)	10610552
Total length (>= 1000 bp)	10610552
Total length (>= 5000 bp)	10588025
Total length (>= 10000 bp)	10445382
Total length (>= 25000 bp)	9717089
Total length (>= 50000 bp)	8837874
# contigs	125
Largest contig	2911968
Total length	10610552
GC (%)	40.25
N50	1004266
N90	29319
auN	1175051.0
L50	3
L90	47
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	46 + 0 part

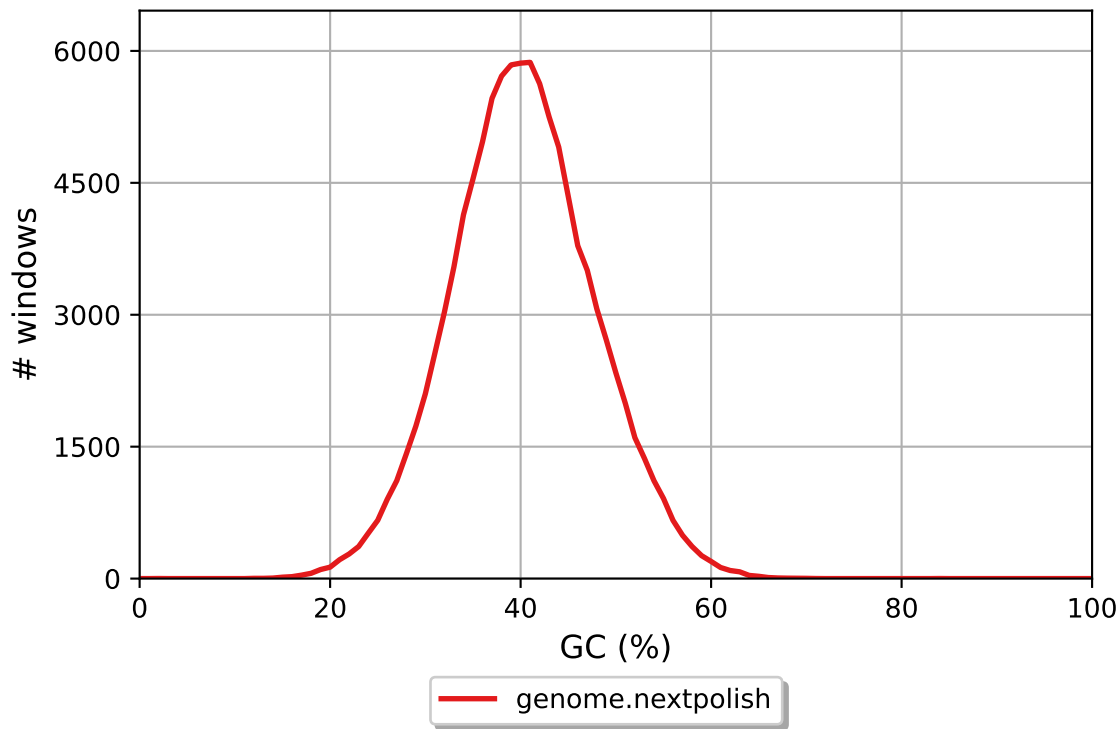
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

