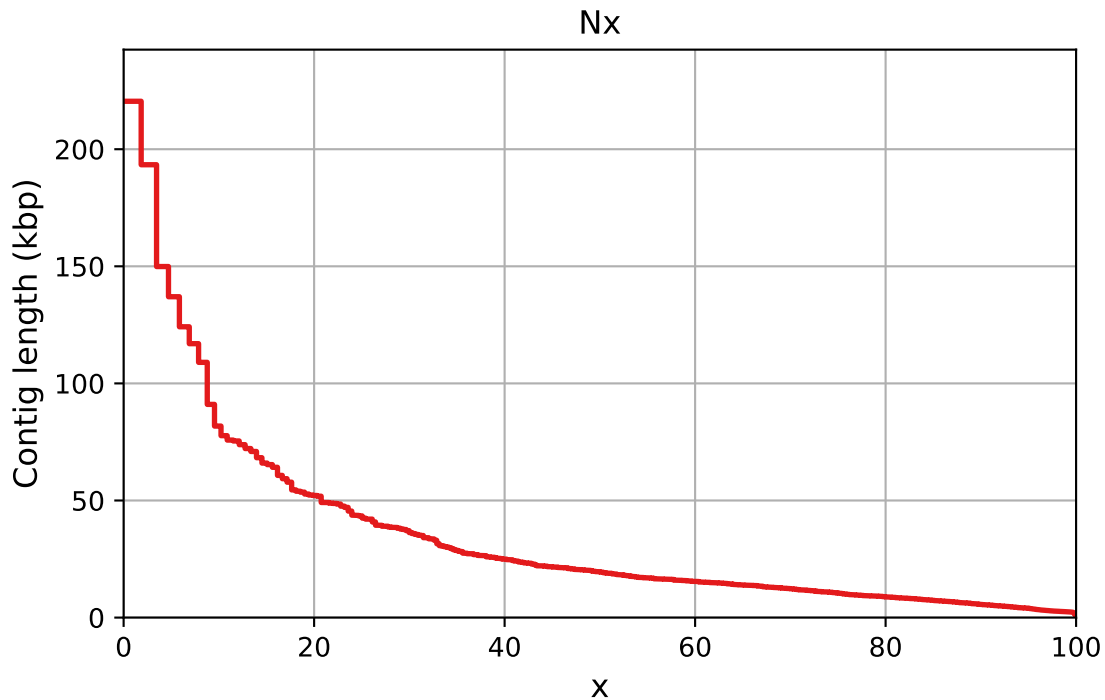


Report

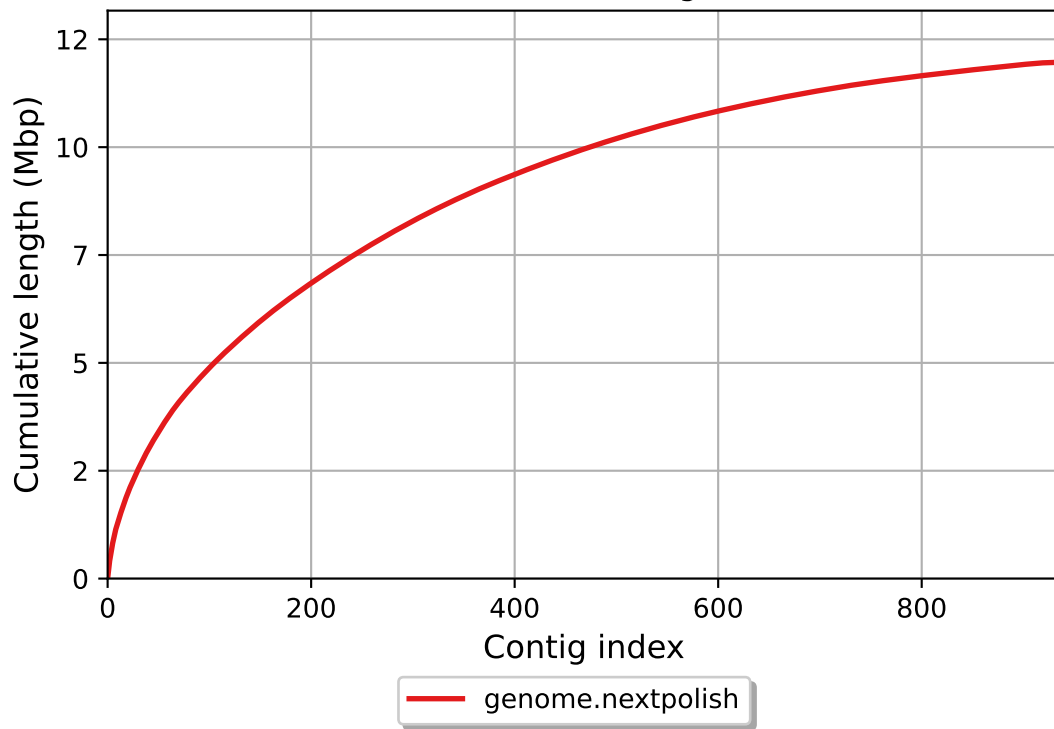
	genome.nextpolish
# contigs (>= 0 bp)	936
# contigs (>= 1000 bp)	923
# contigs (>= 5000 bp)	633
# contigs (>= 10000 bp)	370
# contigs (>= 25000 bp)	95
# contigs (>= 50000 bp)	29
Total length (>= 0 bp)	11967754
Total length (>= 1000 bp)	11959333
Total length (>= 5000 bp)	11008852
Total length (>= 10000 bp)	9080156
Total length (>= 25000 bp)	4762067
Total length (>= 50000 bp)	2482267
# contigs	936
Largest contig	220471
Total length	11967754
GC (%)	44.50
N50	19574
N90	5583
auN	36465.9
L50	151
L90	589
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 4 part

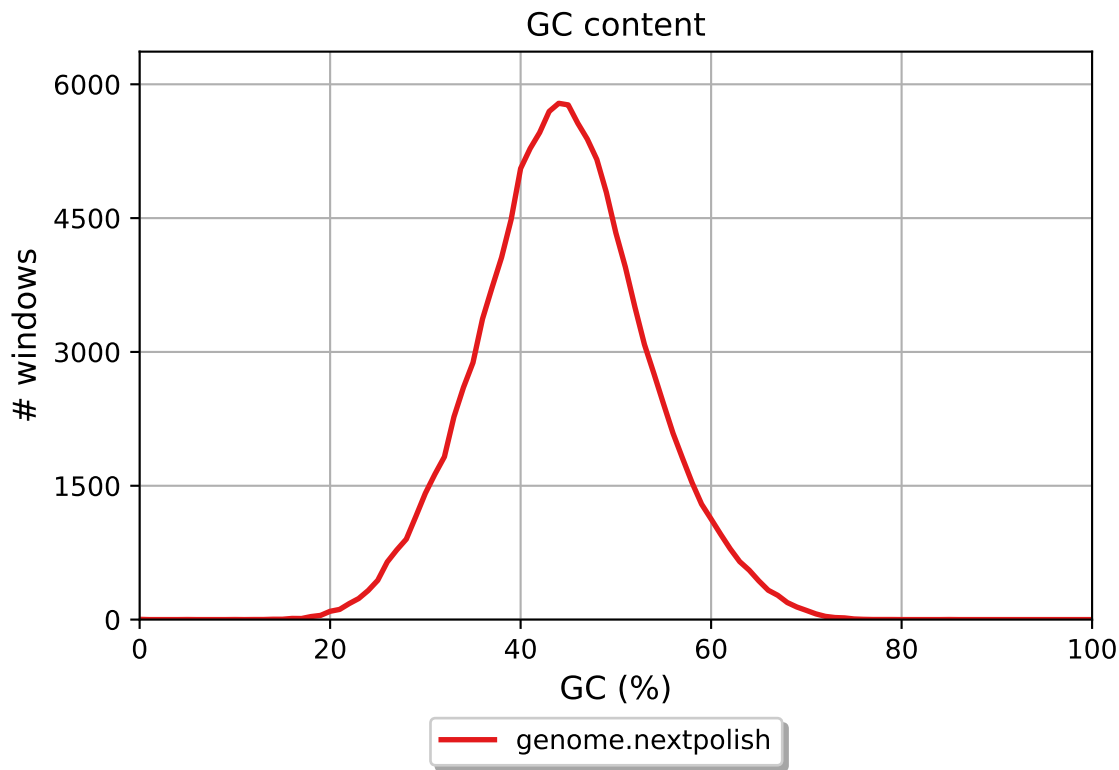
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

