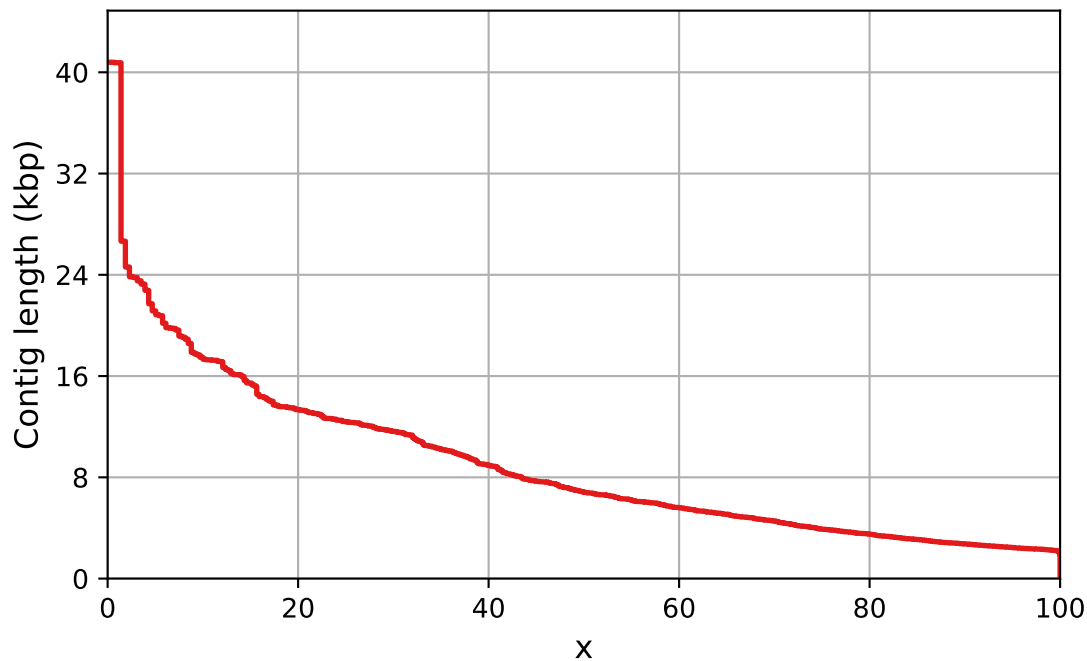


Report

	genome.nextpolish
# contigs (>= 0 bp)	1024
# contigs (>= 1000 bp)	1024
# contigs (>= 5000 bp)	396
# contigs (>= 10000 bp)	145
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5799509
Total length (>= 1000 bp)	5799509
Total length (>= 5000 bp)	3797470
Total length (>= 10000 bp)	2106733
Total length (>= 25000 bp)	108204
Total length (>= 50000 bp)	0
# contigs	1024
Largest contig	40791
Total length	5799509
GC (%)	46.64
N50	6831
N90	2736
auN	9044.7
L50	243
L90	788
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 1 part

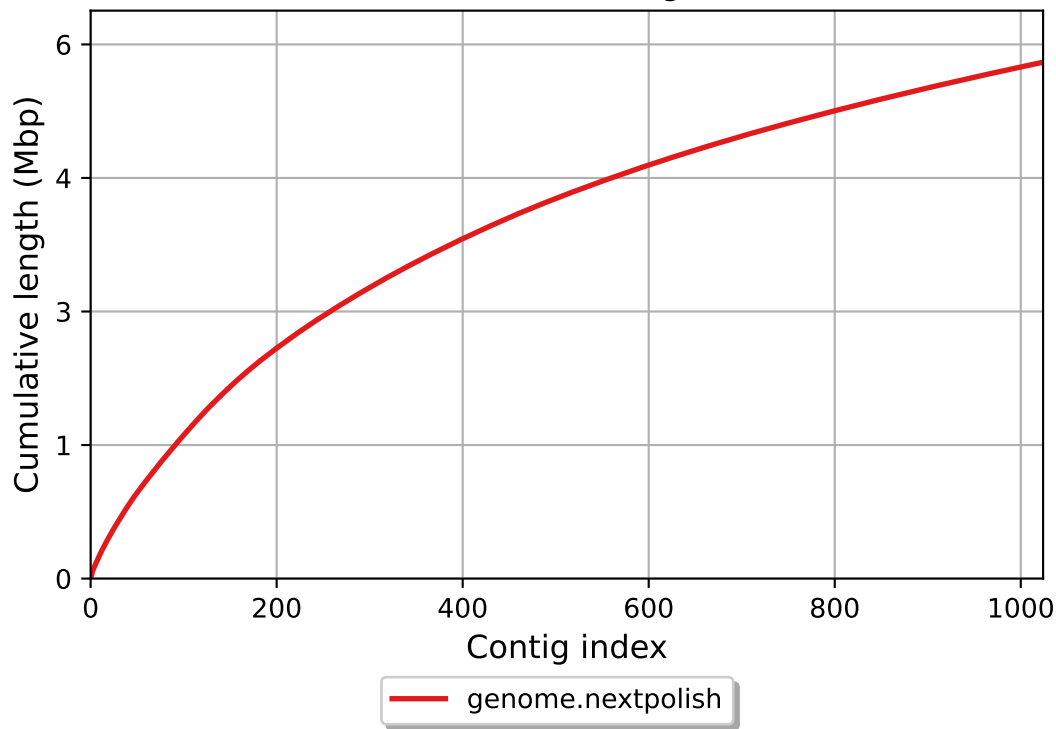
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

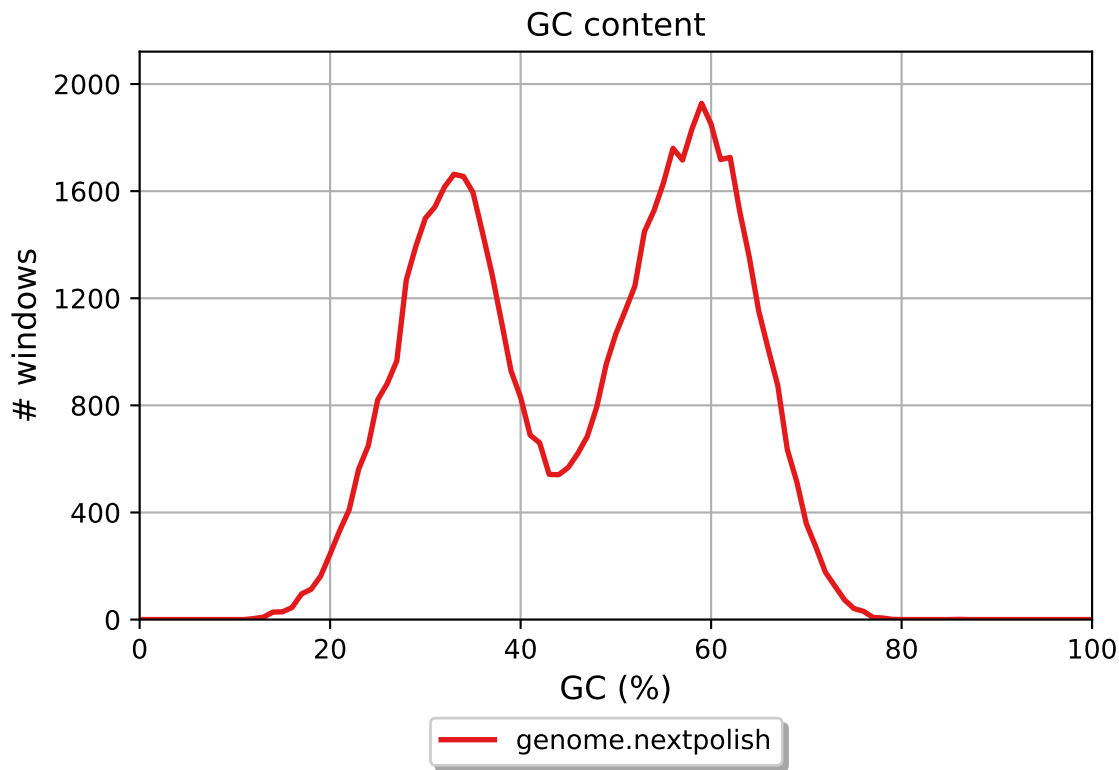
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

