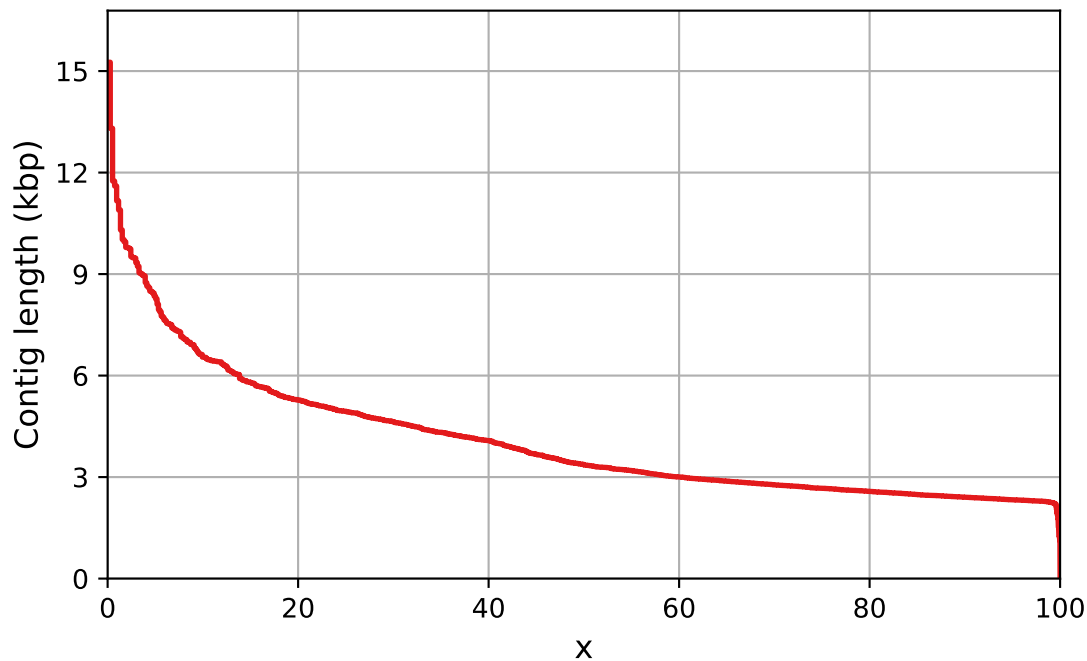


Report

	genome.nextpolish
# contigs (>= 0 bp)	1576
# contigs (>= 1000 bp)	1576
# contigs (>= 5000 bp)	202
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5484959
Total length (>= 1000 bp)	5484959
Total length (>= 5000 bp)	1314425
Total length (>= 10000 bp)	94303
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1576
Largest contig	15260
Total length	5484959
GC (%)	42.63
N50	3369
N90	2405
auN	4104.9
L50	547
L90	1336
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 4 part

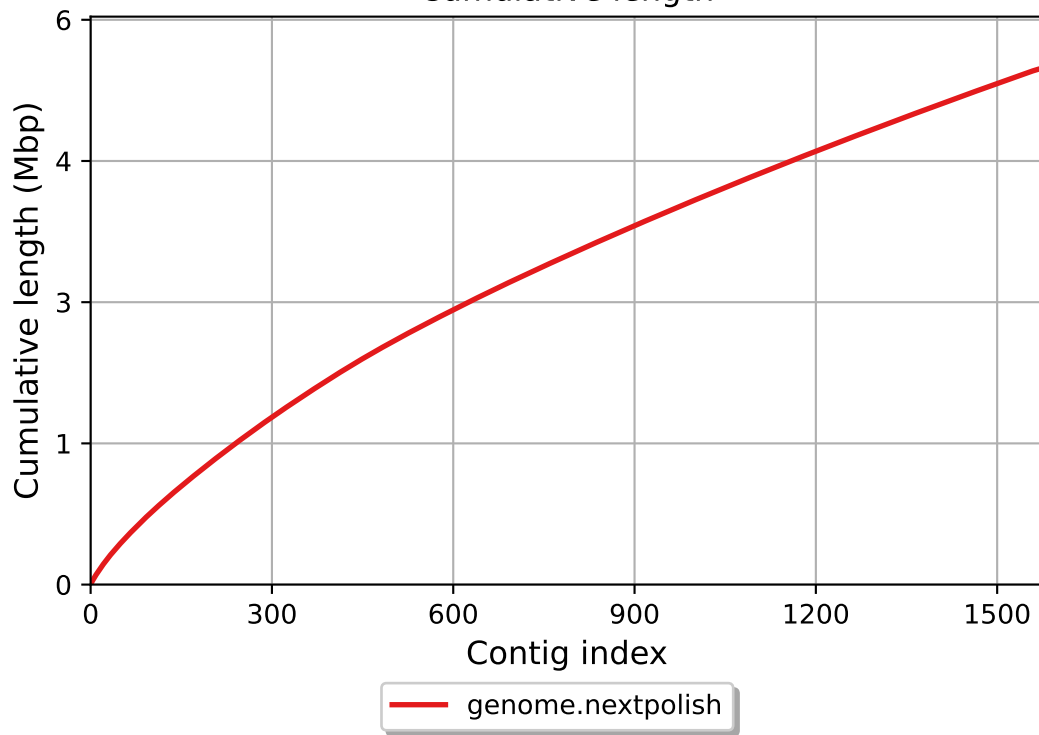
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

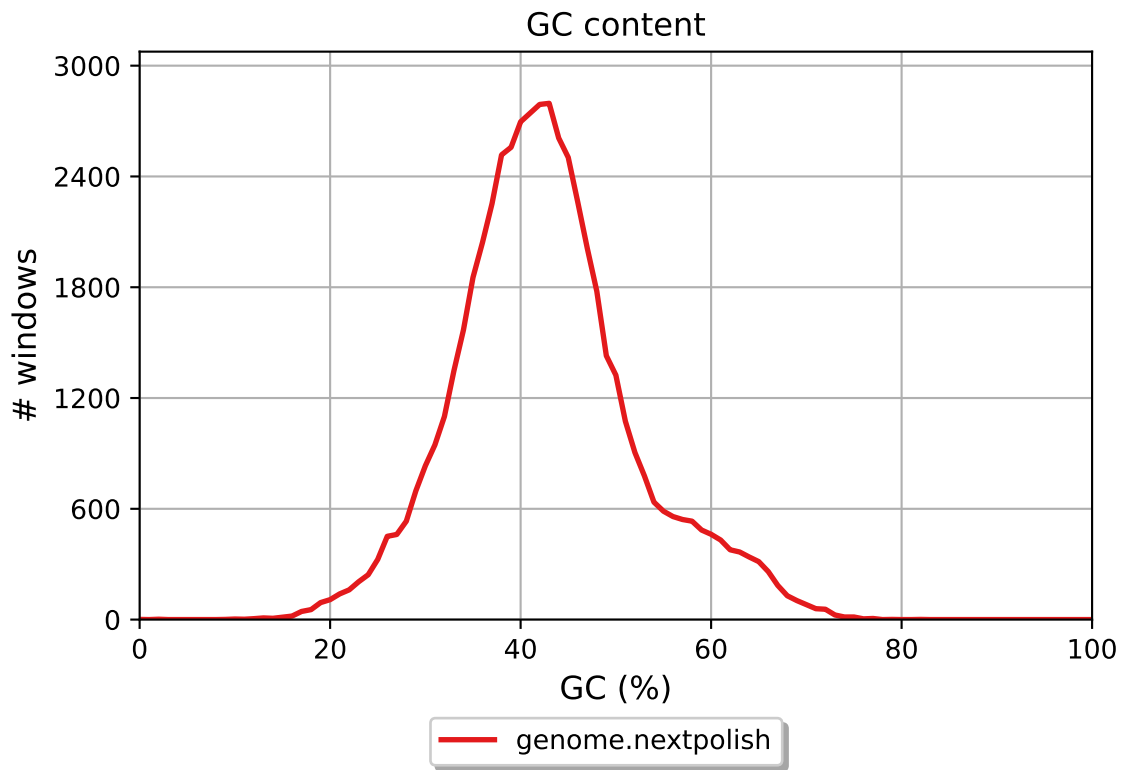
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

