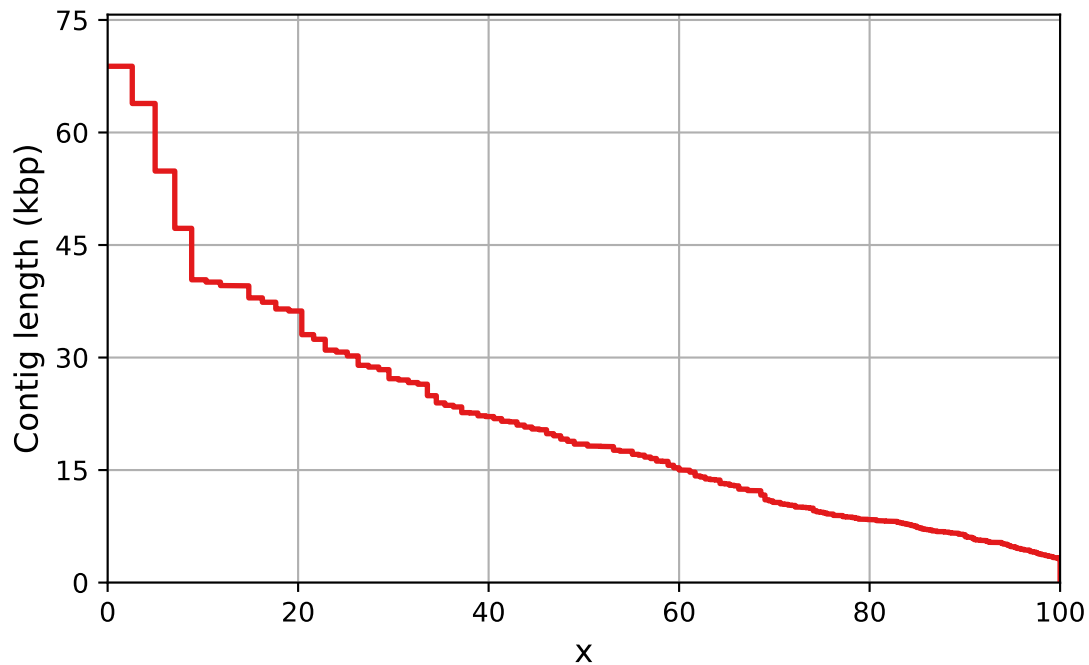


Report

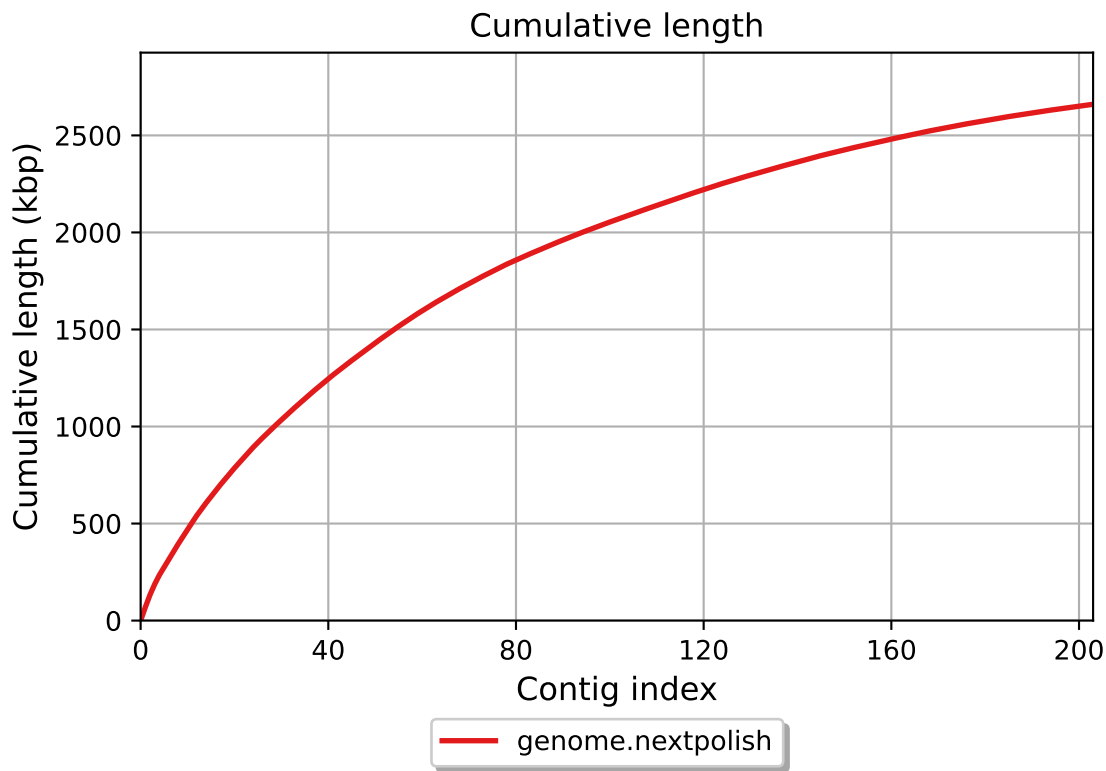
	genome.nextpolish
# contigs (>= 0 bp)	203
# contigs (>= 1000 bp)	203
# contigs (>= 5000 bp)	166
# contigs (>= 10000 bp)	89
# contigs (>= 25000 bp)	24
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	2660408
Total length (>= 1000 bp)	2660408
Total length (>= 5000 bp)	2512460
Total length (>= 10000 bp)	1951003
Total length (>= 25000 bp)	893130
Total length (>= 50000 bp)	187552
# contigs	203
Largest contig	68827
Total length	2660408
GC (%)	58.08
N50	18459
N90	6366
auN	22340.7
L50	45
L90	145
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

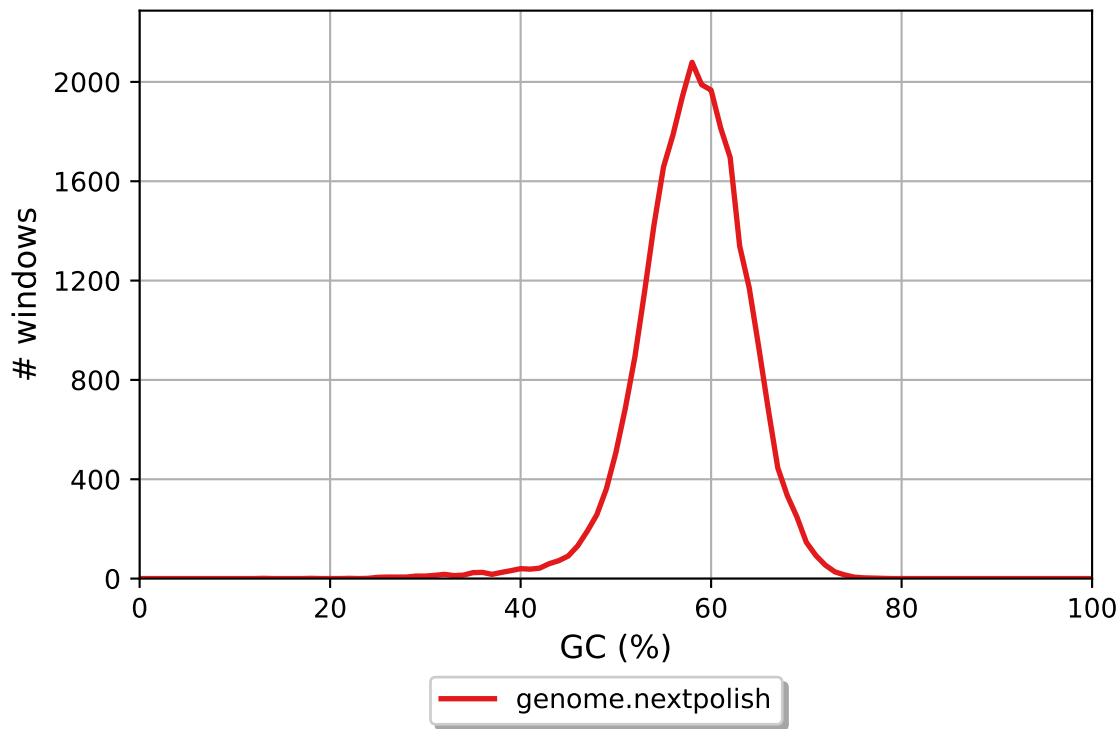
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

