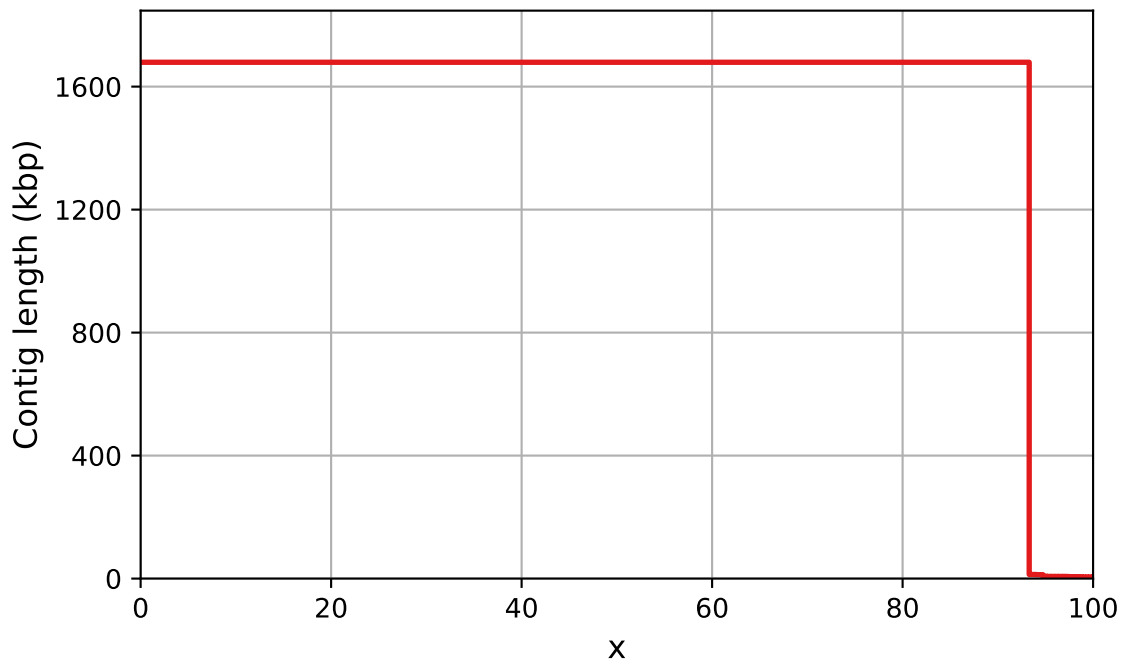


Report

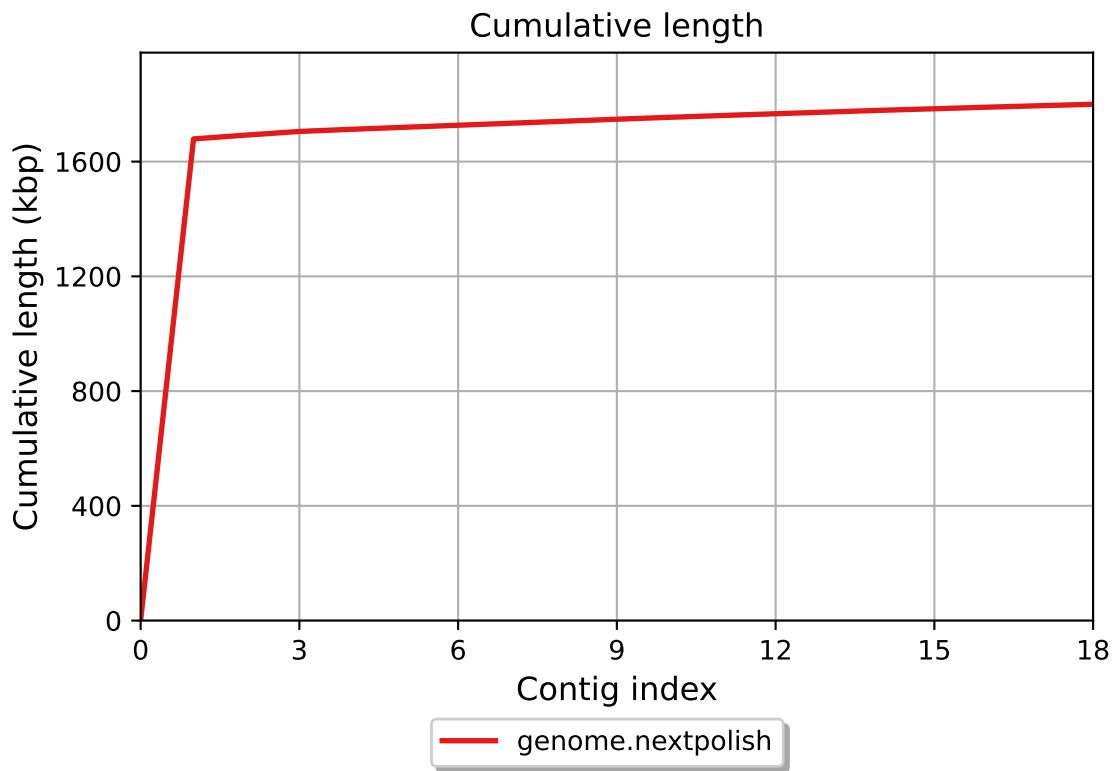
	genome.nextpolish
# contigs (>= 0 bp)	18
# contigs (>= 1000 bp)	18
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	1799923
Total length (>= 1000 bp)	1799923
Total length (>= 5000 bp)	1799923
Total length (>= 10000 bp)	1705180
Total length (>= 25000 bp)	1679171
Total length (>= 50000 bp)	1679171
# contigs	18
Largest contig	1679171
Total length	1799923
GC (%)	30.05
N50	1679171
N90	1679171
auN	1567045.4
L50	1
L90	1
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

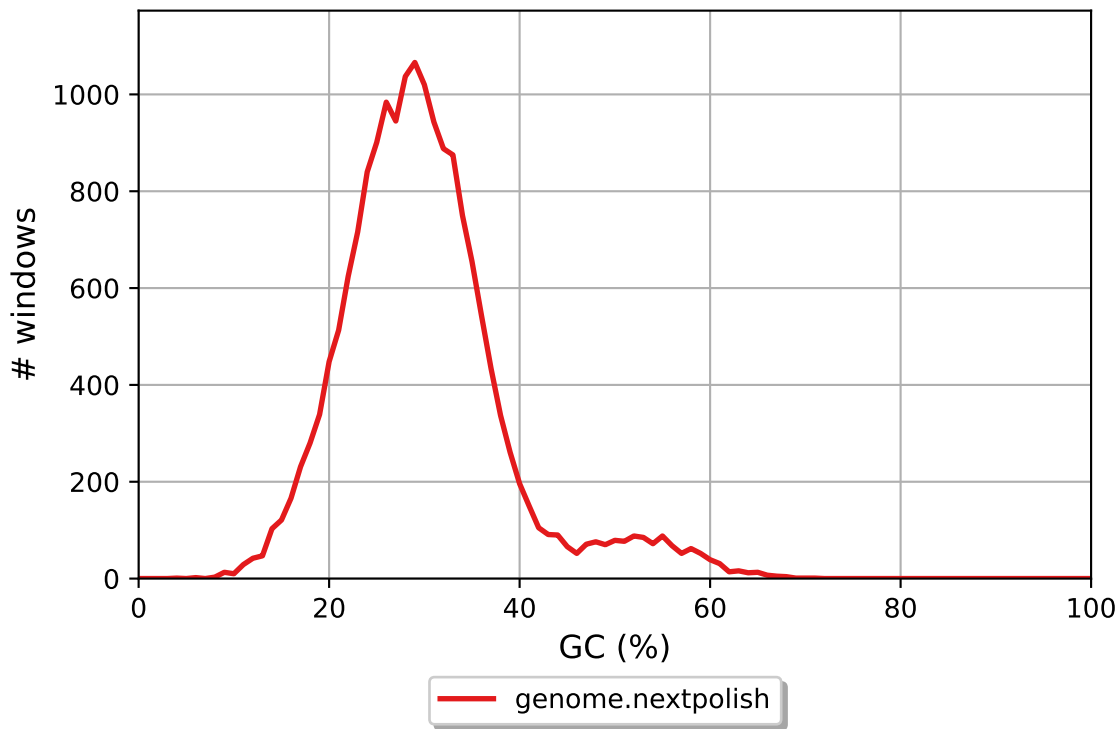
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

