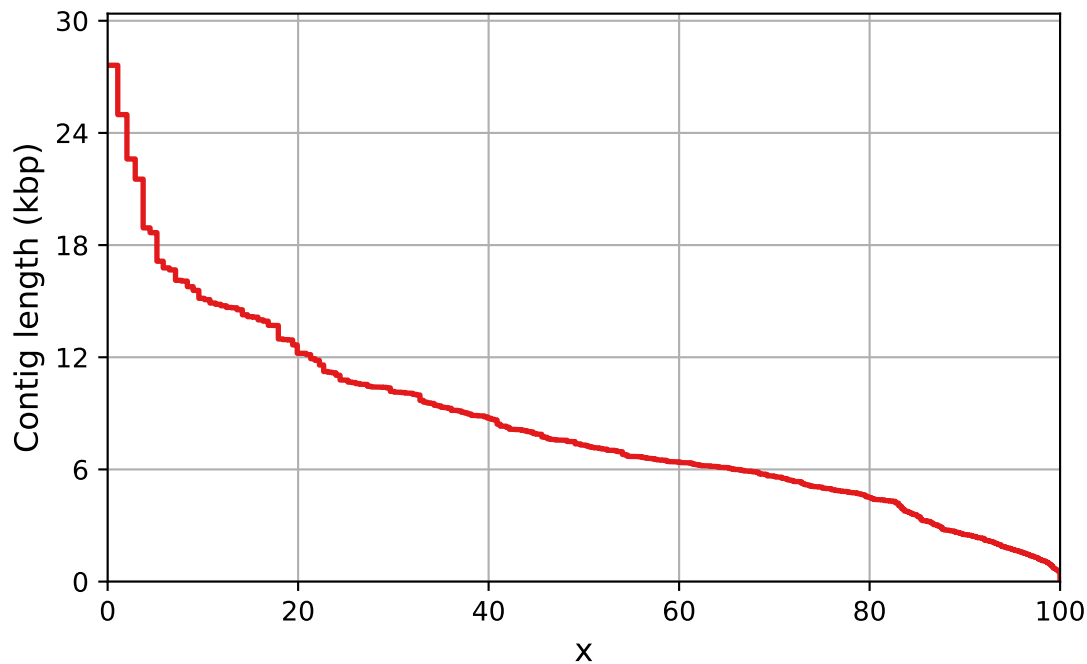


Report

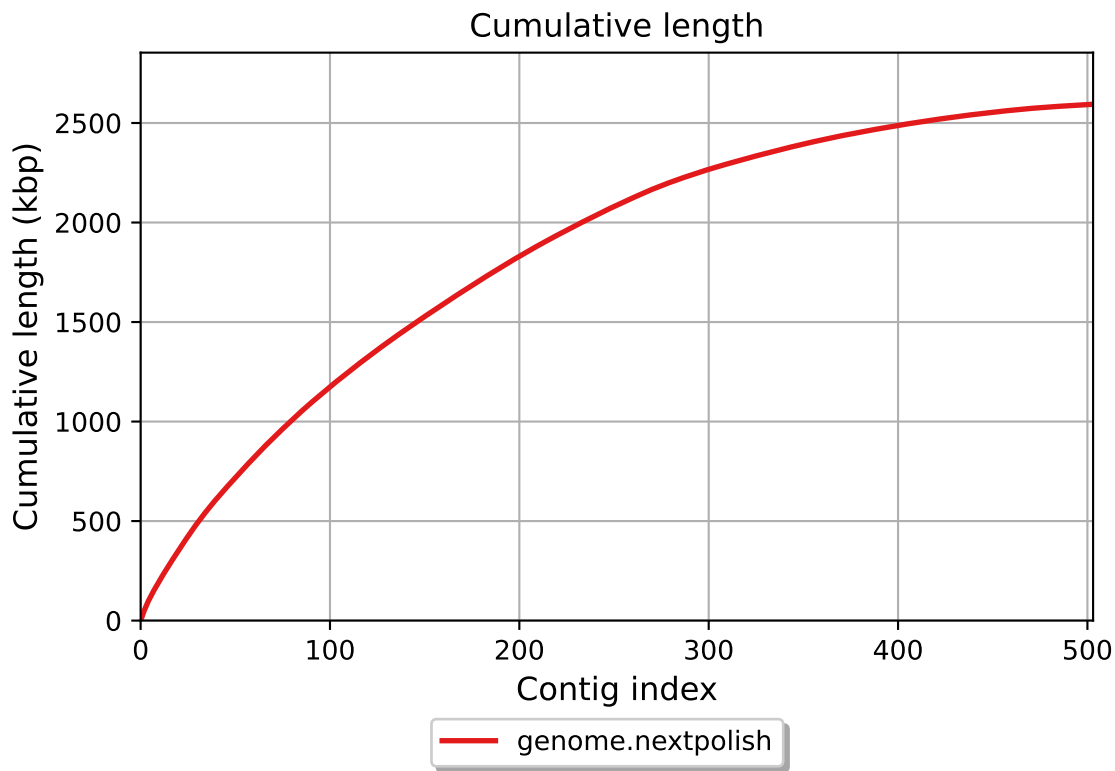
	genome.nextpolish
# contigs (>= 0 bp)	503
# contigs (>= 1000 bp)	458
# contigs (>= 5000 bp)	222
# contigs (>= 10000 bp)	62
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2594127
Total length (>= 1000 bp)	2562284
Total length (>= 5000 bp)	1946505
Total length (>= 10000 bp)	840712
Total length (>= 25000 bp)	27619
Total length (>= 50000 bp)	0
# contigs	503
Largest contig	27619
Total length	2594127
GC (%)	54.13
N50	7301
N90	2515
auN	8523.4
L50	117
L90	326
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 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).

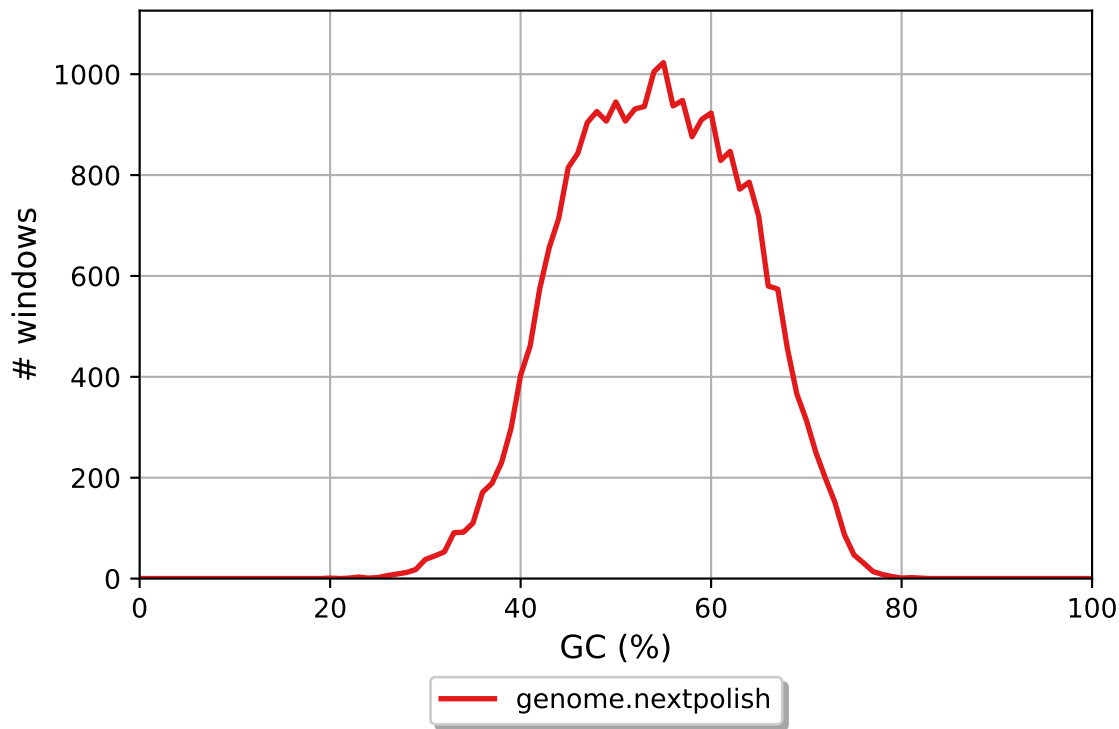
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

