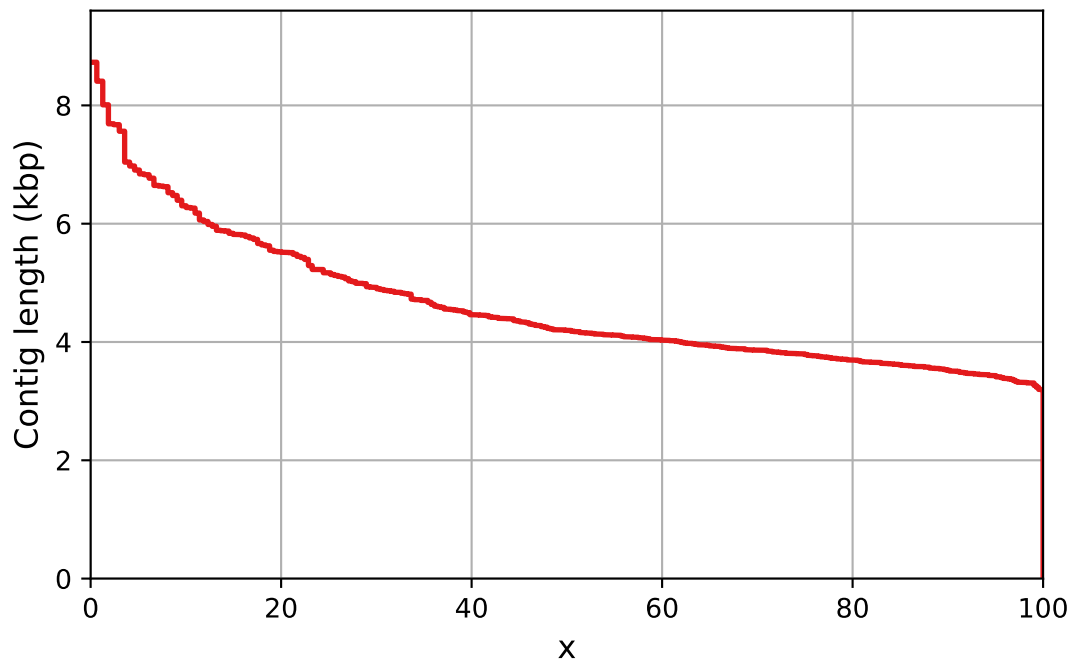


Report

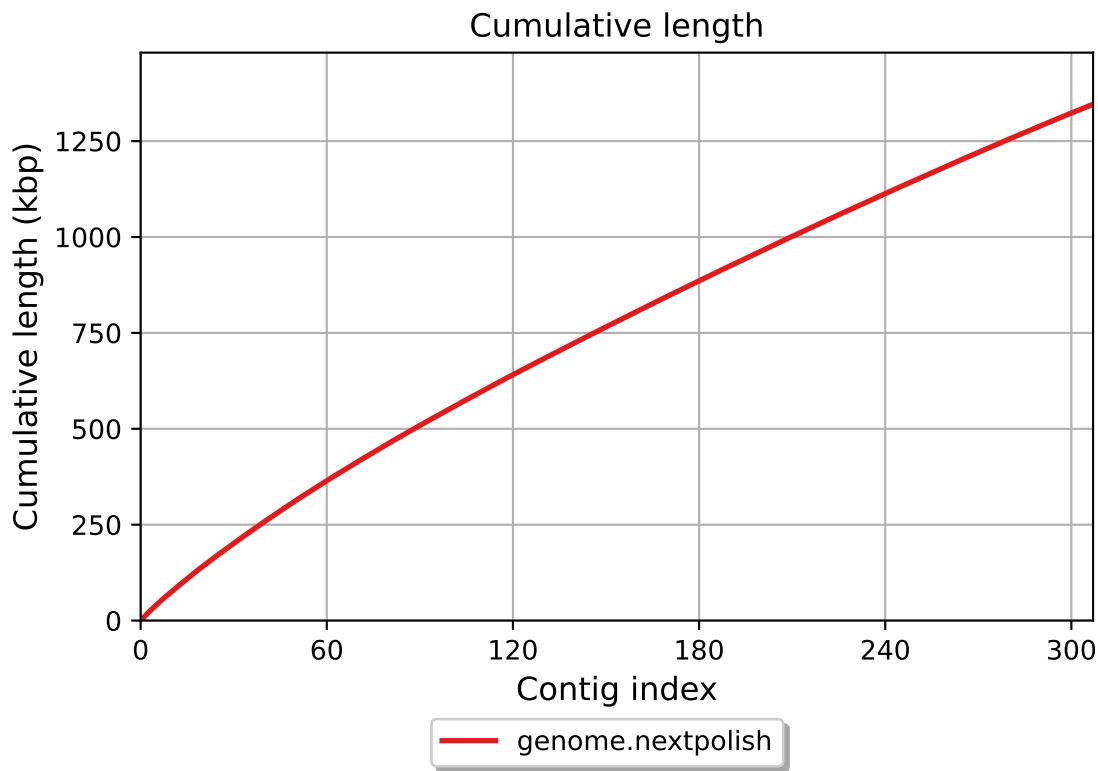
	genome.nextpolish
# contigs (>= 0 bp)	307
# contigs (>= 1000 bp)	307
# contigs (>= 5000 bp)	62
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1346091
Total length (>= 1000 bp)	1346091
Total length (>= 5000 bp)	374739
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	307
Largest contig	8729
Total length	1346091
GC (%)	57.69
N50	4197
N90	3523
auN	4611.3
L50	128
L90	268
# 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).

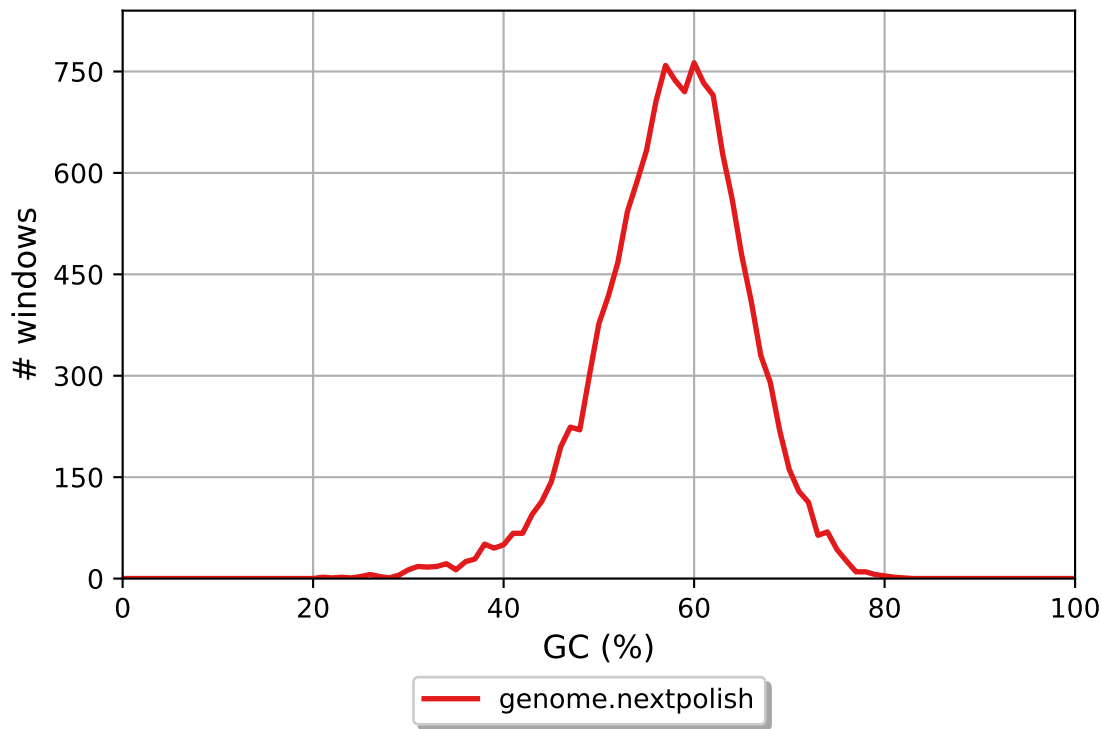
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

