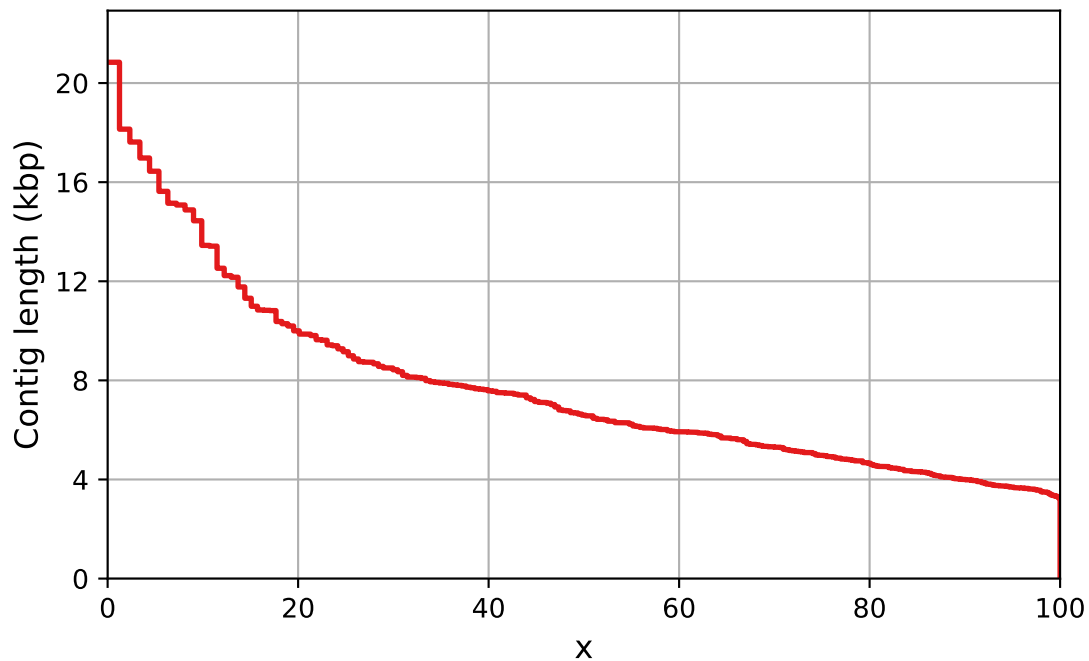


Report

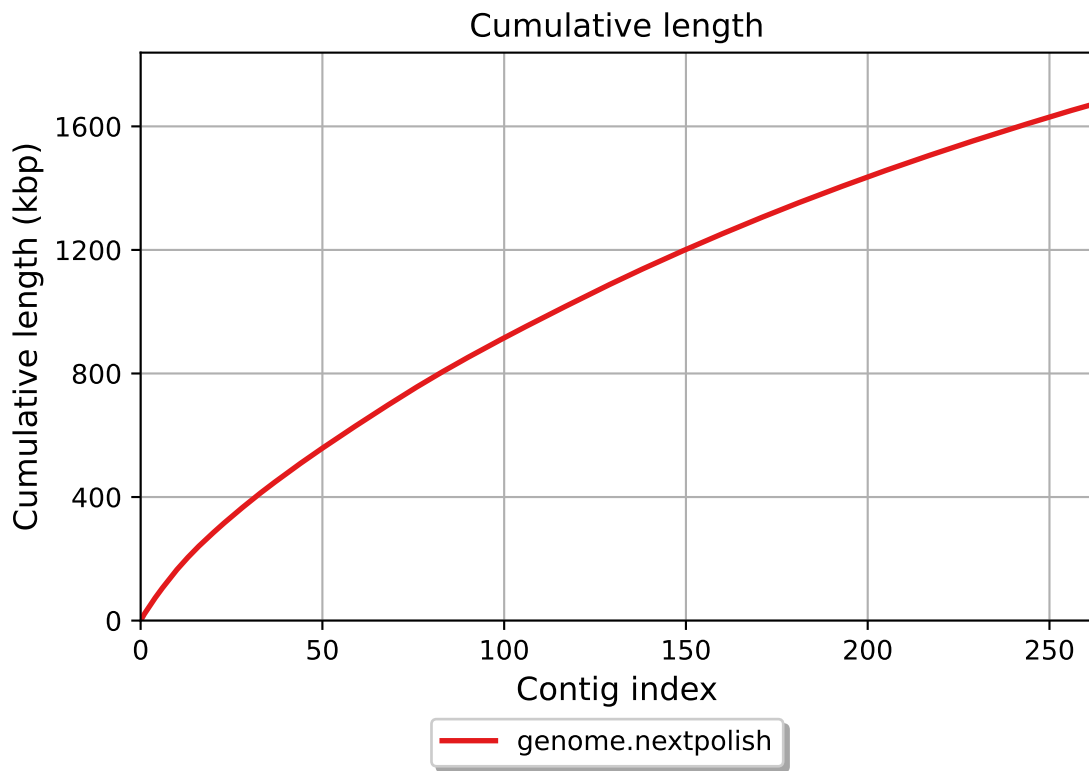
	genome.nextpolish
# contigs (>= 0 bp)	262
# contigs (>= 1000 bp)	262
# contigs (>= 5000 bp)	158
# contigs (>= 10000 bp)	25
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1671568
Total length (>= 1000 bp)	1671568
Total length (>= 5000 bp)	1242345
Total length (>= 10000 bp)	336391
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	262
Largest contig	20839
Total length	1671568
GC (%)	58.63
N50	6597
N90	3999
auN	7731.7
L50	88
L90	217
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 2 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).

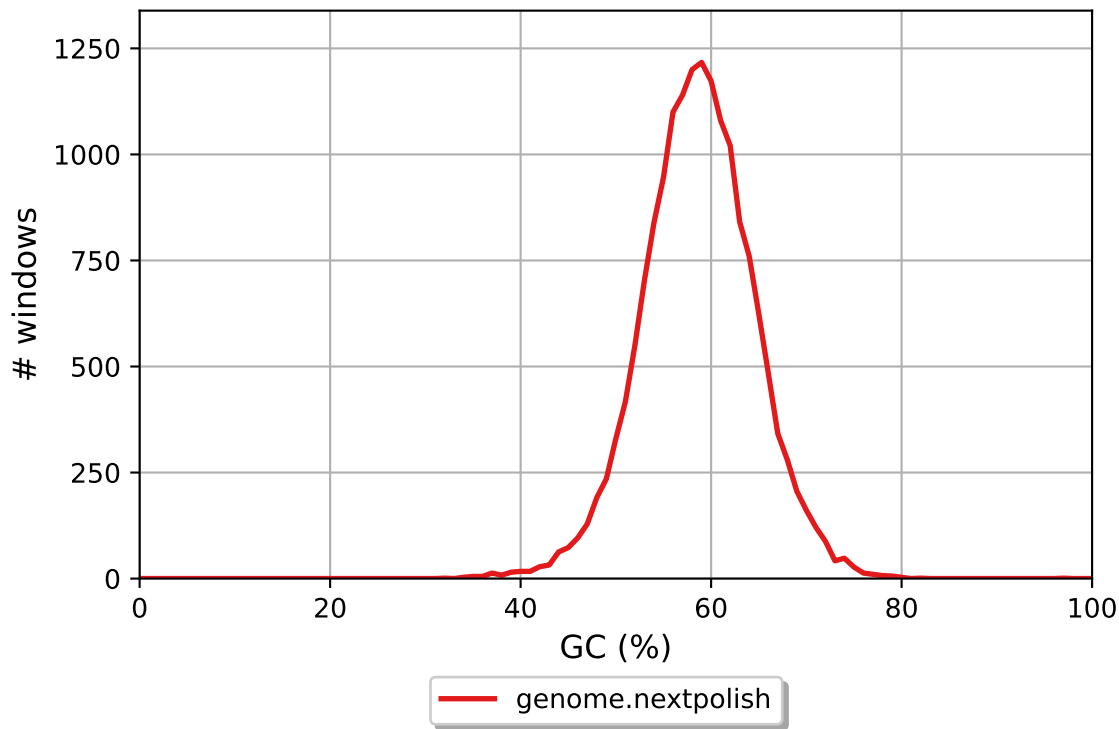
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

