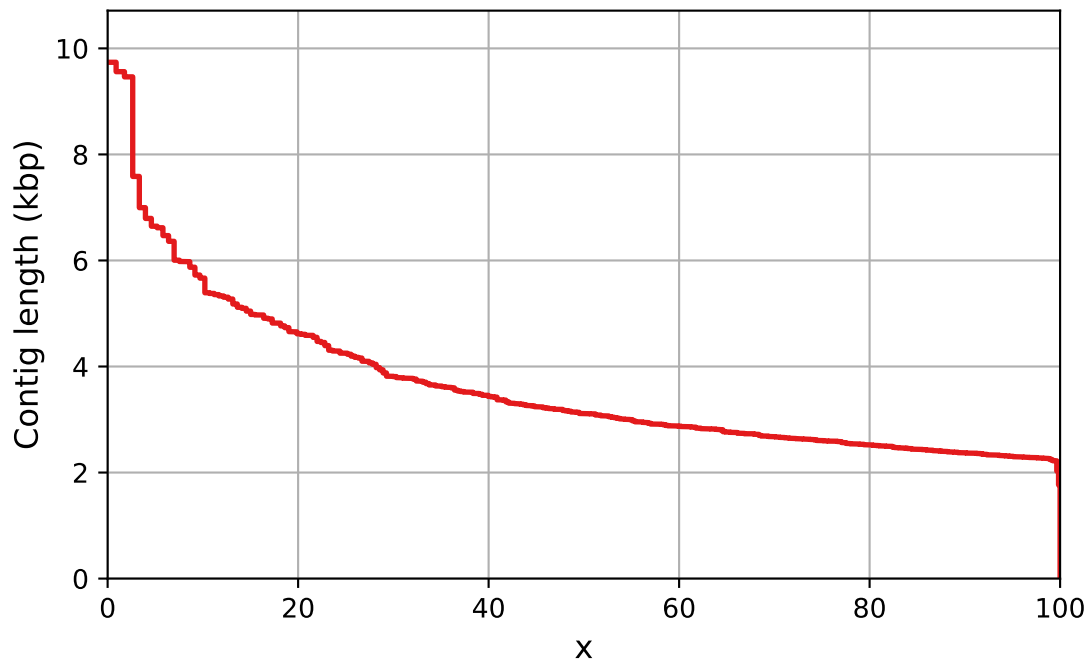


Report

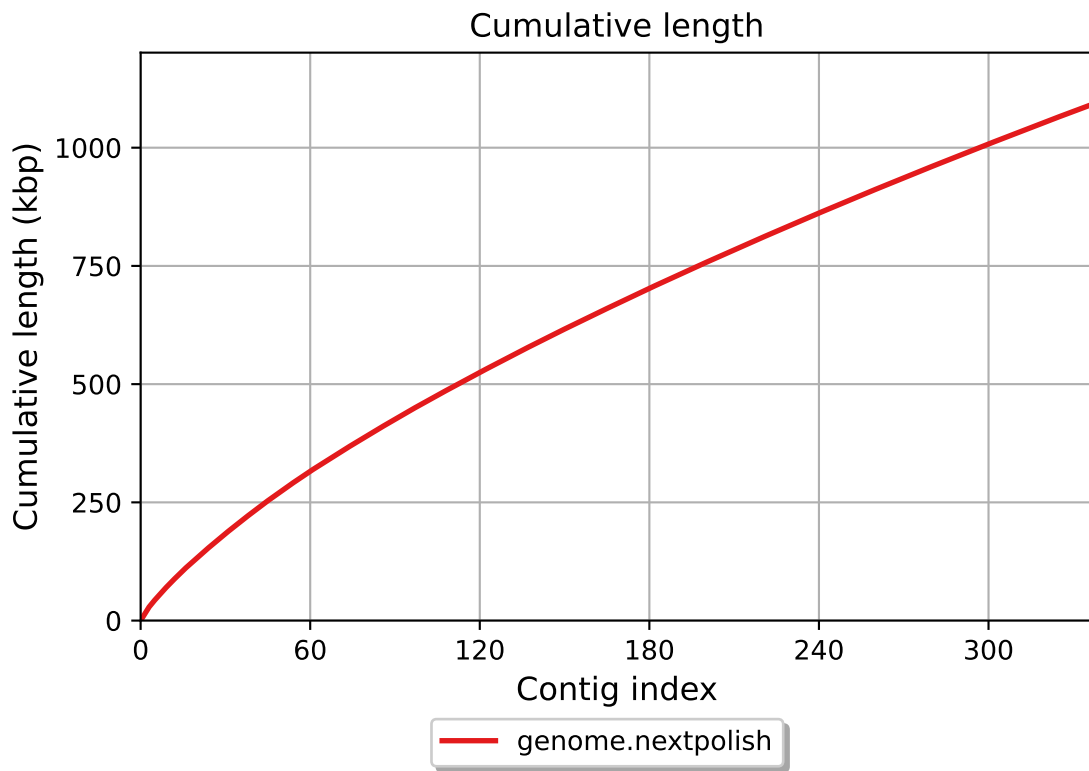
	genome.nextpolish
# contigs (>= 0 bp)	337
# contigs (>= 1000 bp)	337
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1091823
Total length (>= 1000 bp)	1091823
Total length (>= 5000 bp)	163929
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	337
Largest contig	9738
Total length	1091823
GC (%)	54.94
N50	3112
N90	2372
auN	3654.6
L50	127
L90	290
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 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).

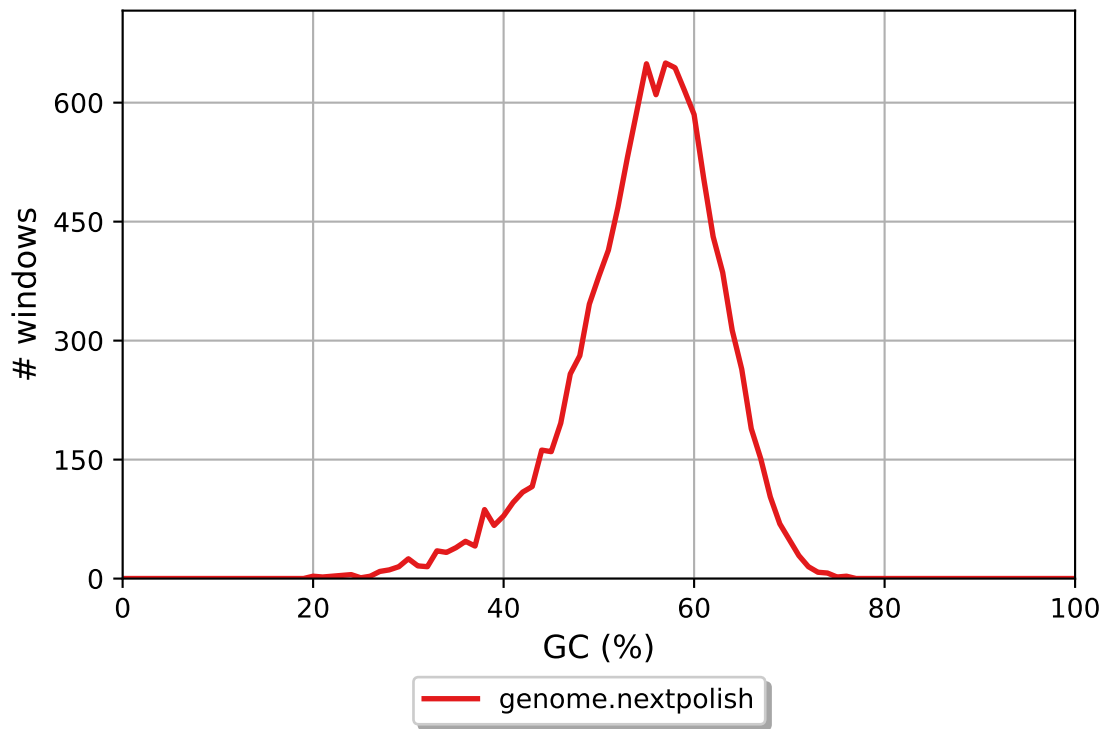
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

