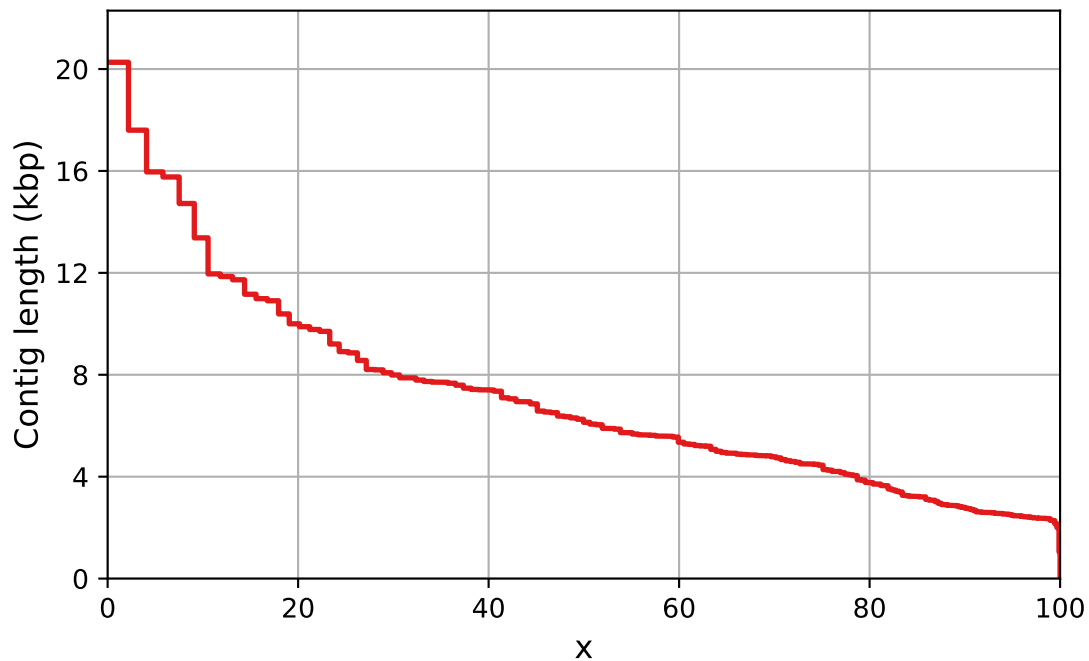


Report

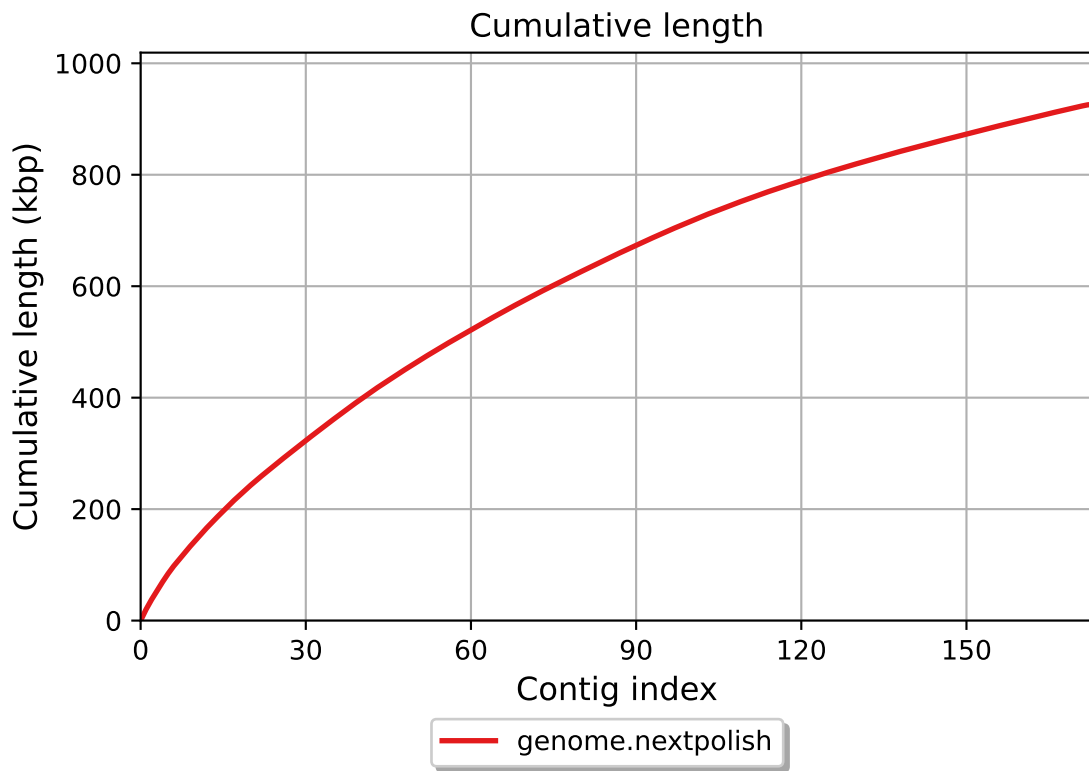
	genome.nextpolish
# contigs (>= 0 bp)	173
# contigs (>= 1000 bp)	173
# contigs (>= 5000 bp)	73
# contigs (>= 10000 bp)	14
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	926467
Total length (>= 1000 bp)	926467
Total length (>= 5000 bp)	591731
Total length (>= 10000 bp)	186669
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	173
Largest contig	20265
Total length	926467
GC (%)	43.25
N50	6134
N90	2763
auN	7228.4
L50	51
L90	136
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

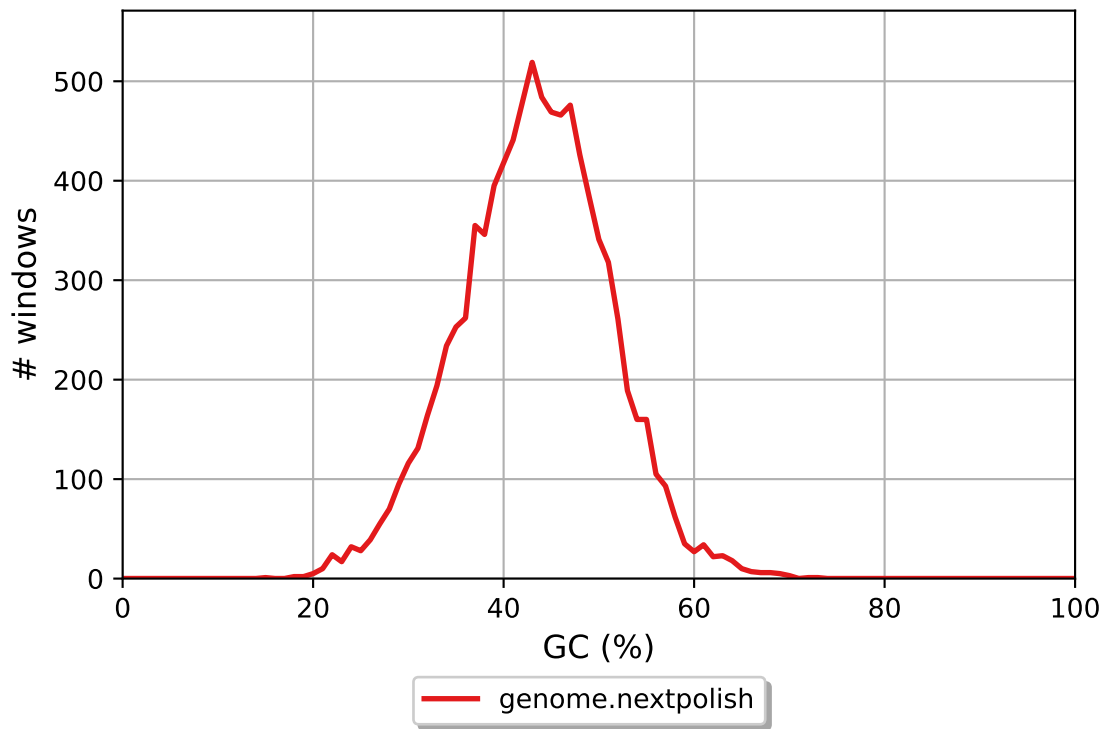
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

