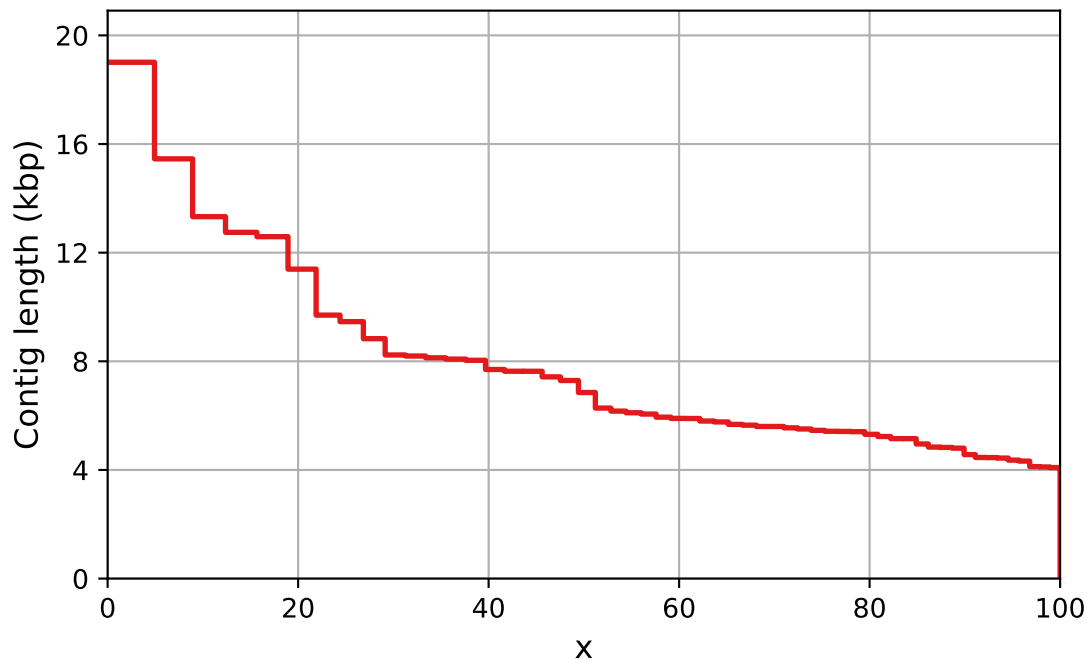


Report

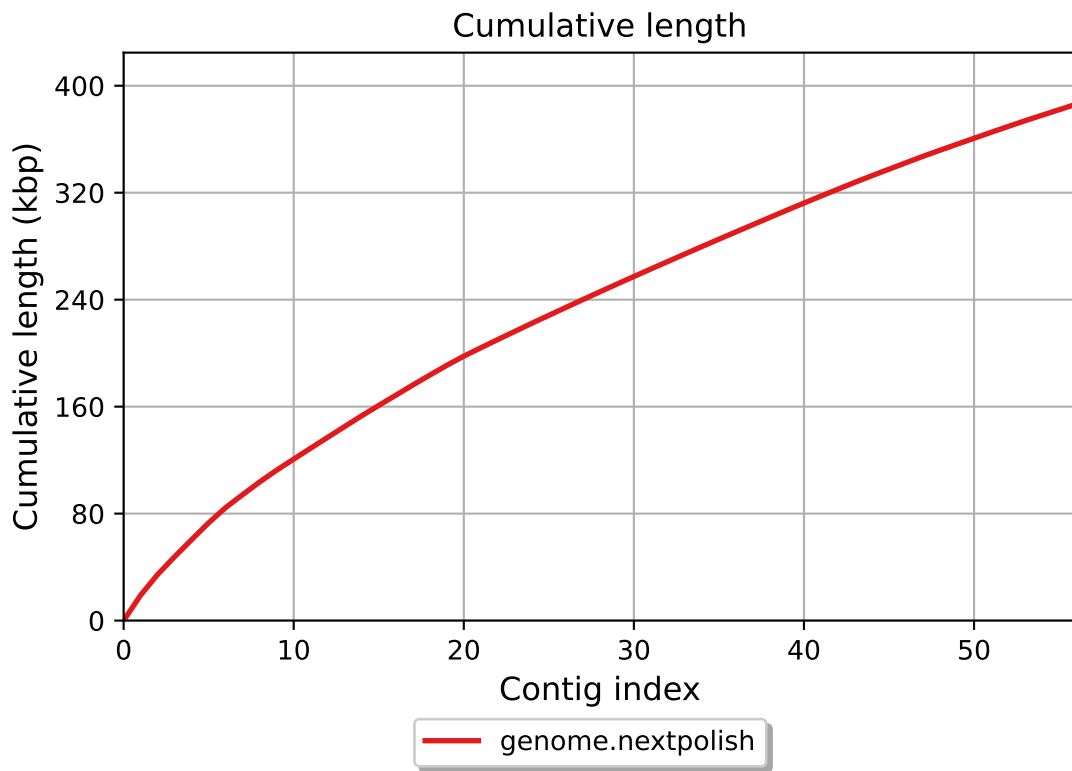
	genome.nextpolish
# contigs (>= 0 bp)	56
# contigs (>= 1000 bp)	56
# contigs (>= 5000 bp)	43
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	386133
Total length (>= 1000 bp)	386133
Total length (>= 5000 bp)	327781
Total length (>= 10000 bp)	84520
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	56
Largest contig	19010
Total length	386133
GC (%)	34.12
N50	6851
N90	4565
auN	8165.3
L50	20
L90	48
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

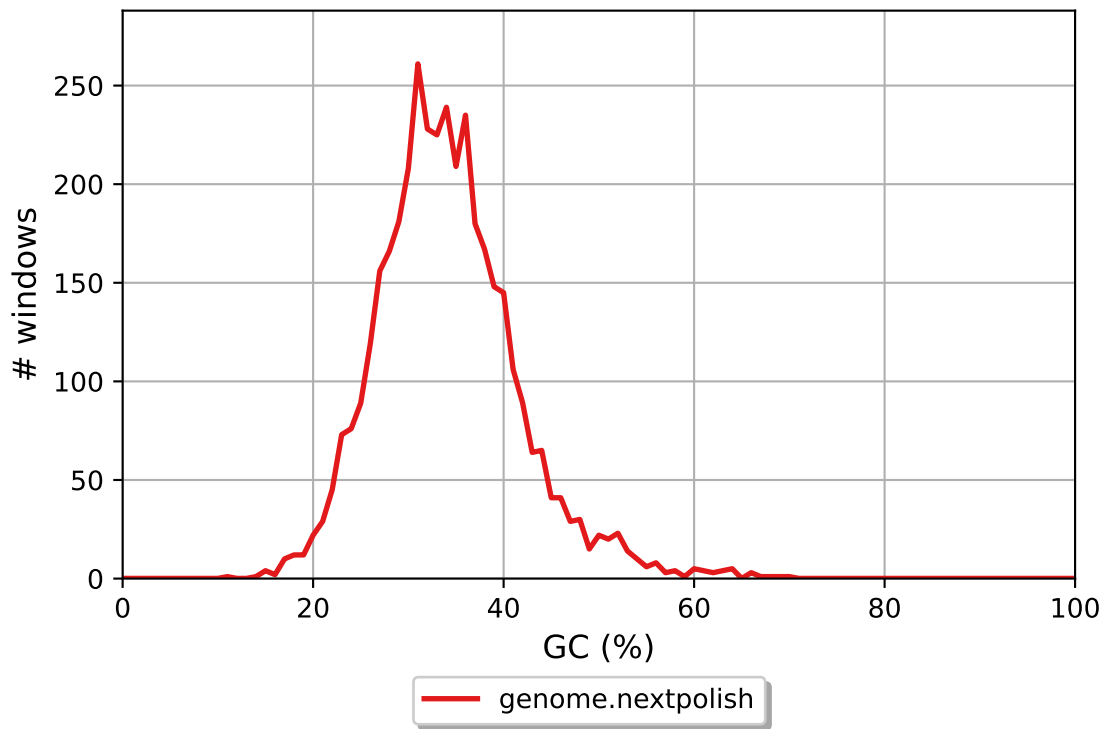
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

