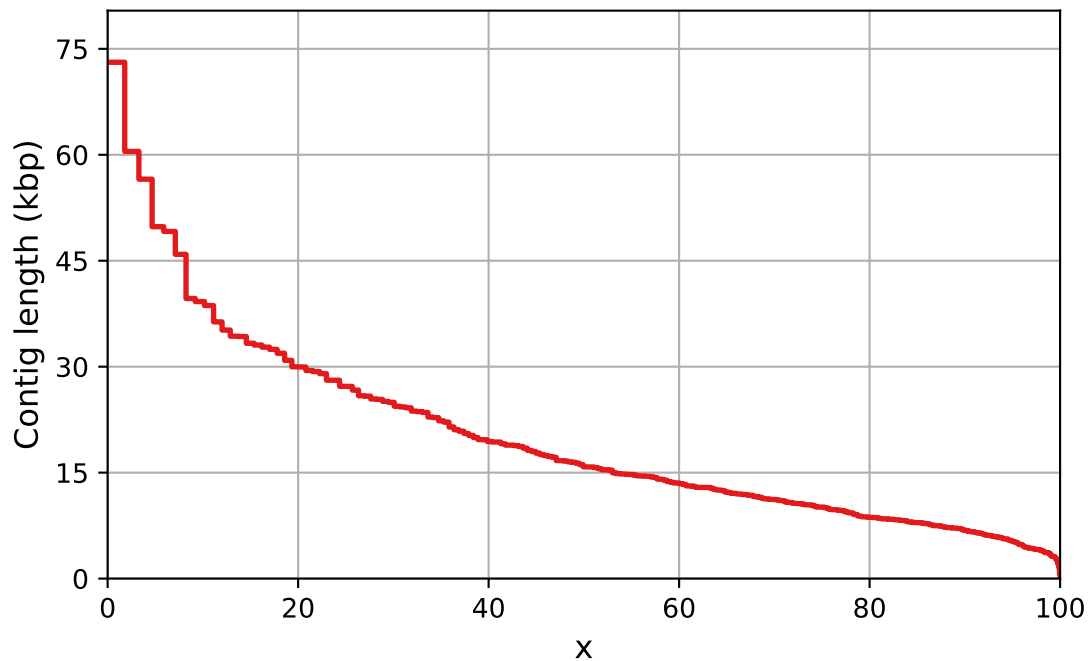


Report

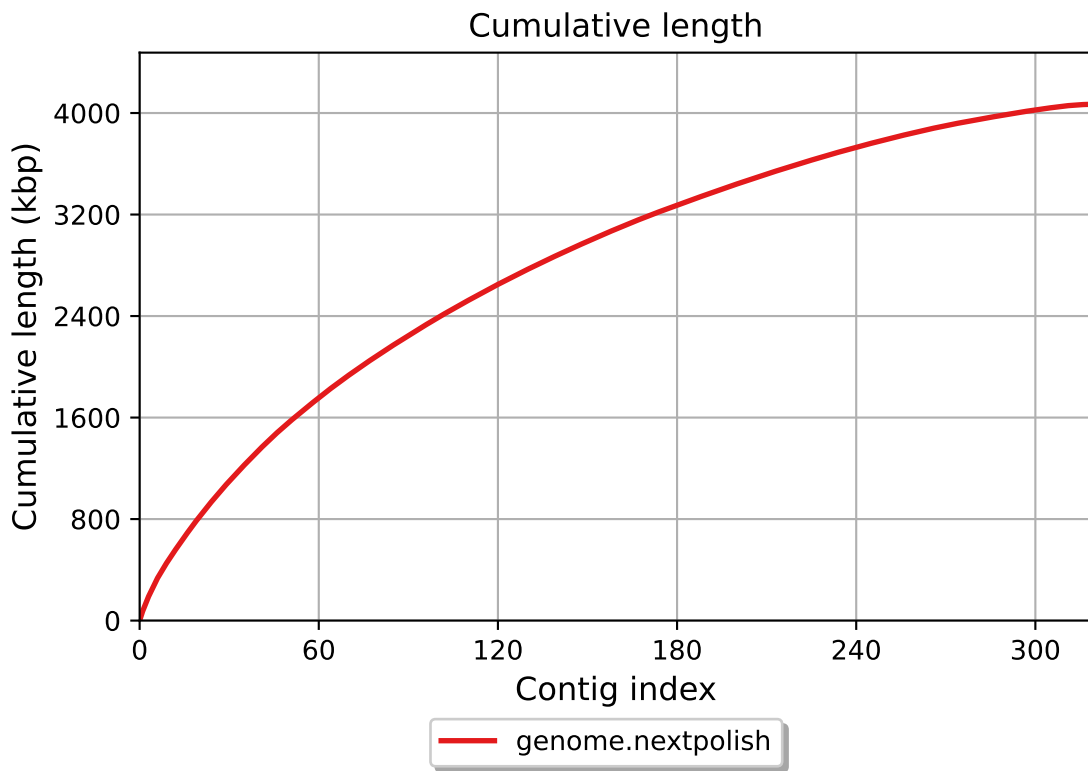
	genome.nextpolish
# contigs (>= 0 bp)	319
# contigs (>= 1000 bp)	315
# contigs (>= 5000 bp)	268
# contigs (>= 10000 bp)	158
# contigs (>= 25000 bp)	34
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	4068992
Total length (>= 1000 bp)	4065878
Total length (>= 5000 bp)	3890818
Total length (>= 10000 bp)	3070242
Total length (>= 25000 bp)	1199638
Total length (>= 50000 bp)	190121
# contigs	319
Largest contig	73098
Total length	4068992
GC (%)	45.70
N50	15825
N90	6863
auN	20539.2
L50	77
L90	230
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 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).

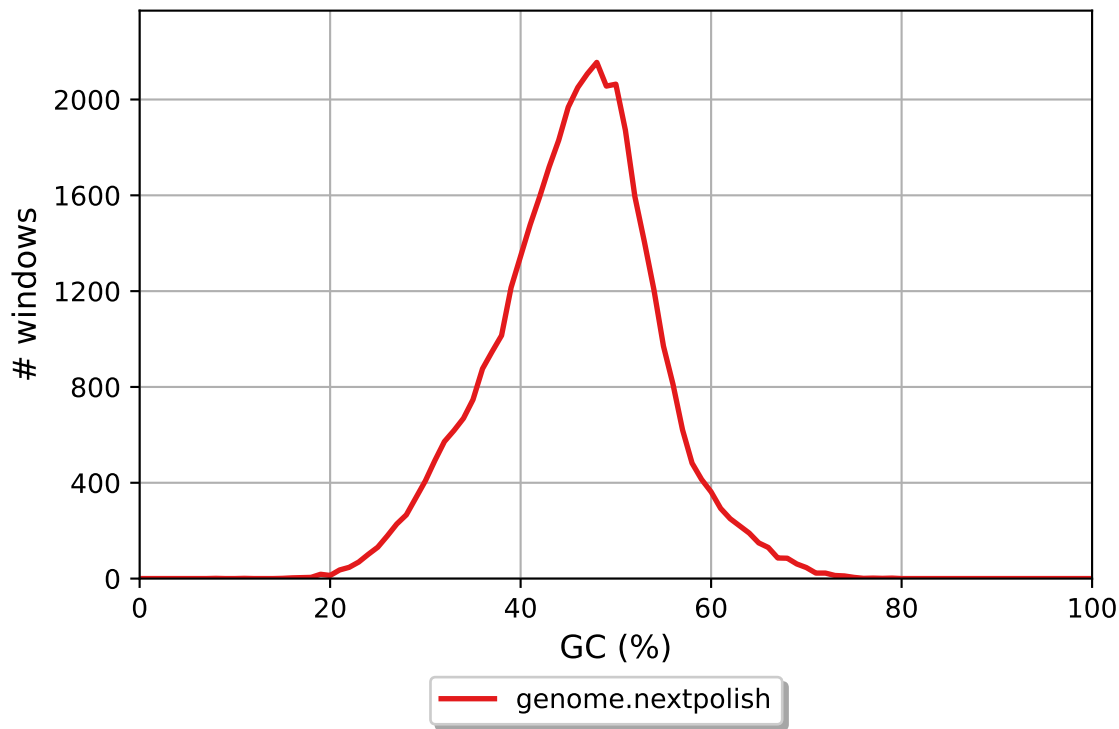
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

