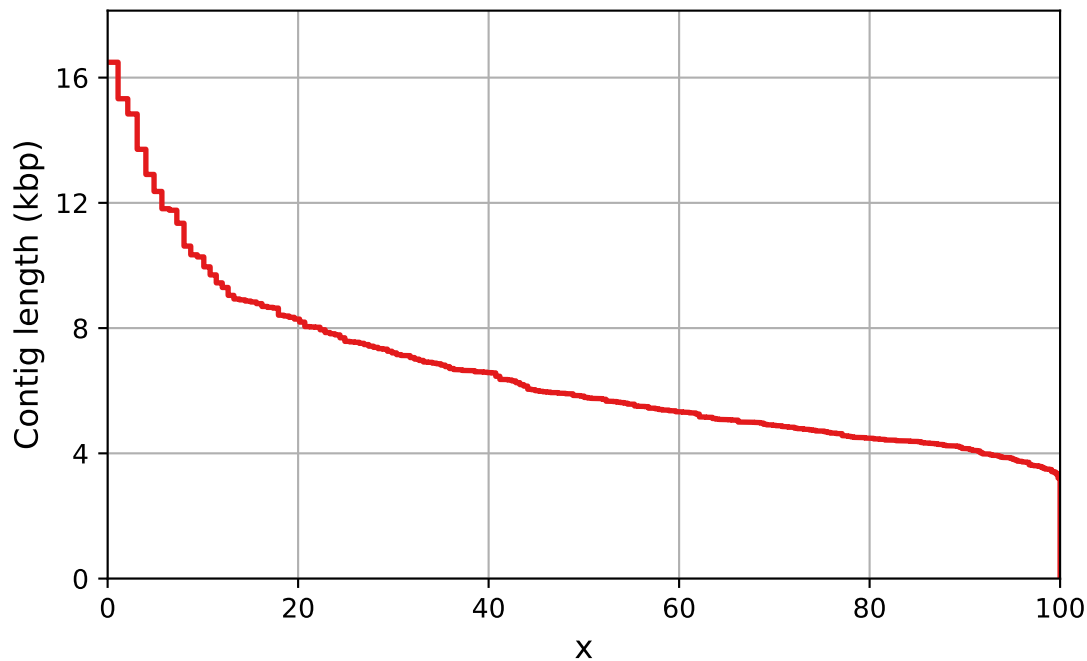


Report

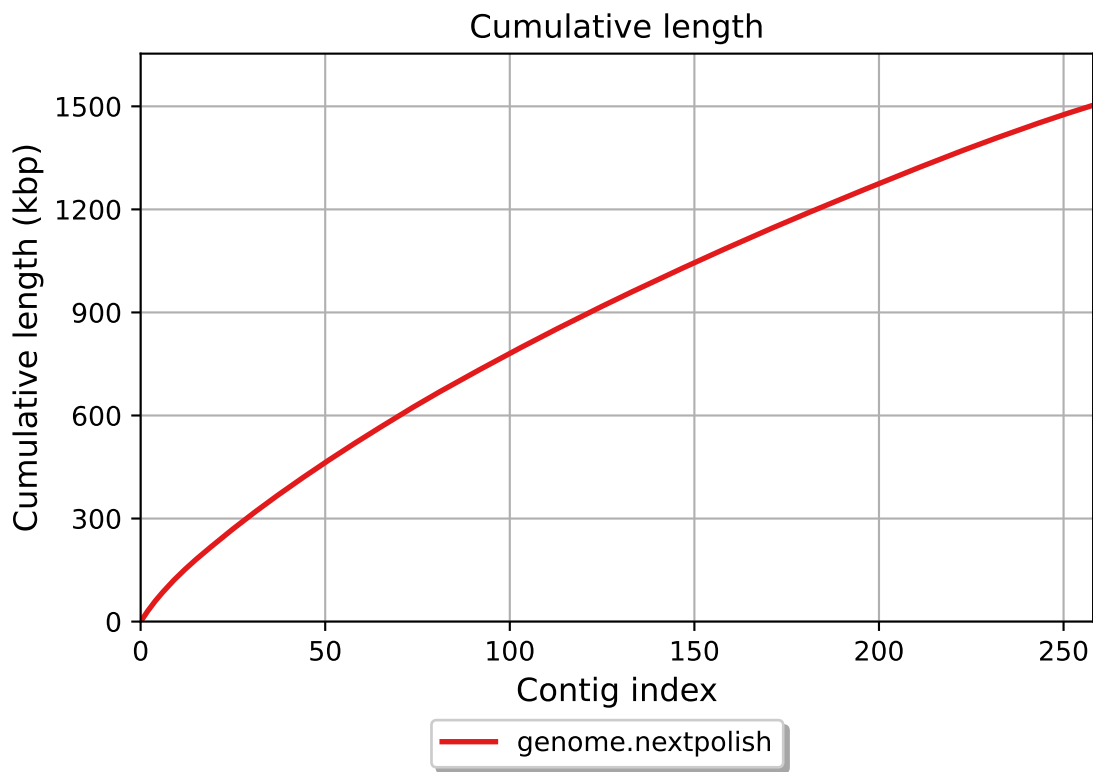
	genome.nextpolish
# contigs (>= 0 bp)	258
# contigs (>= 1000 bp)	258
# contigs (>= 5000 bp)	141
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1503117
Total length (>= 1000 bp)	1503117
Total length (>= 5000 bp)	1000054
Total length (>= 10000 bp)	151794
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	258
Largest contig	16490
Total length	1503117
GC (%)	39.61
N50	5828
N90	4150
auN	6603.8
L50	95
L90	219
# 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).

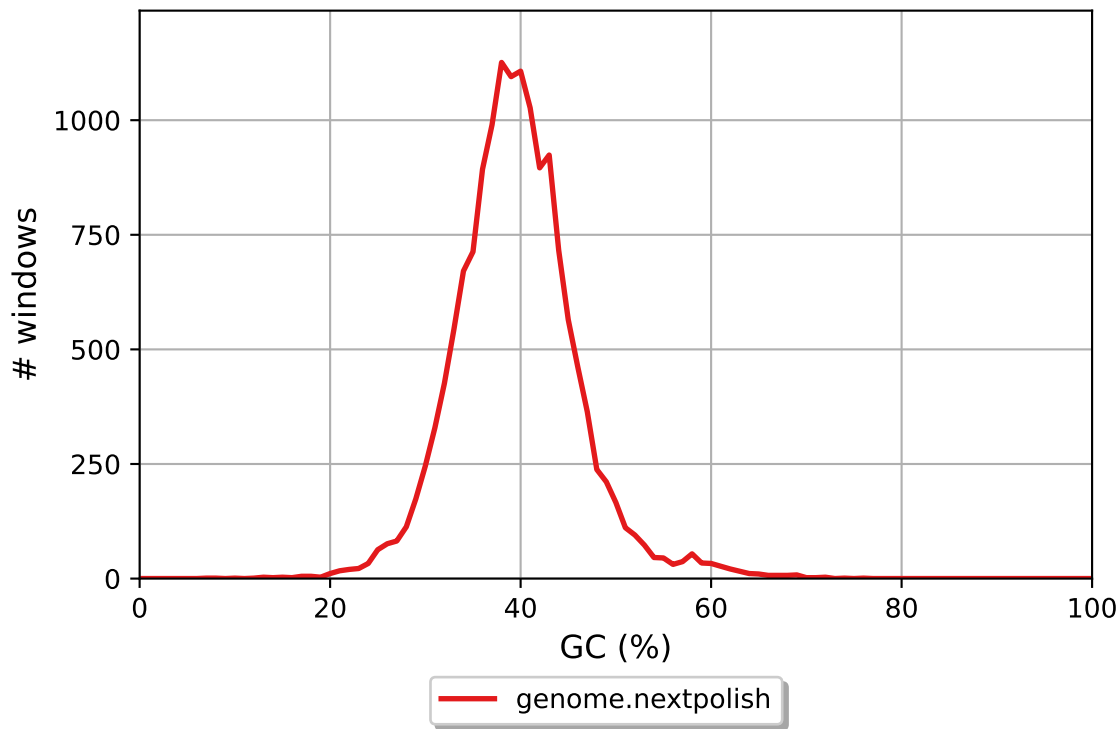
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

