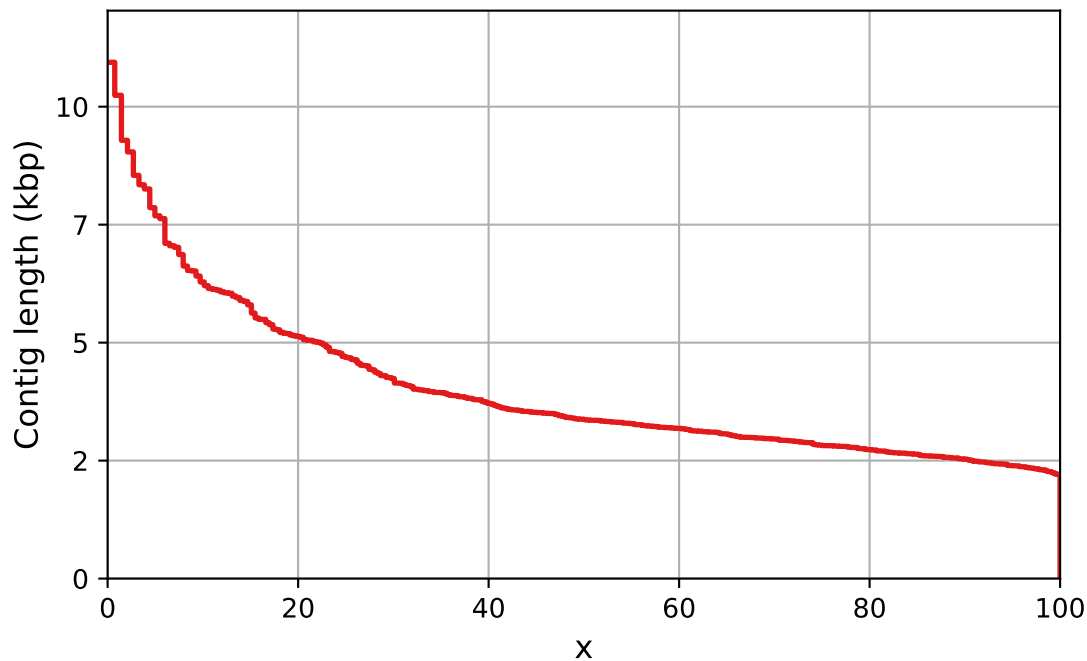


Report

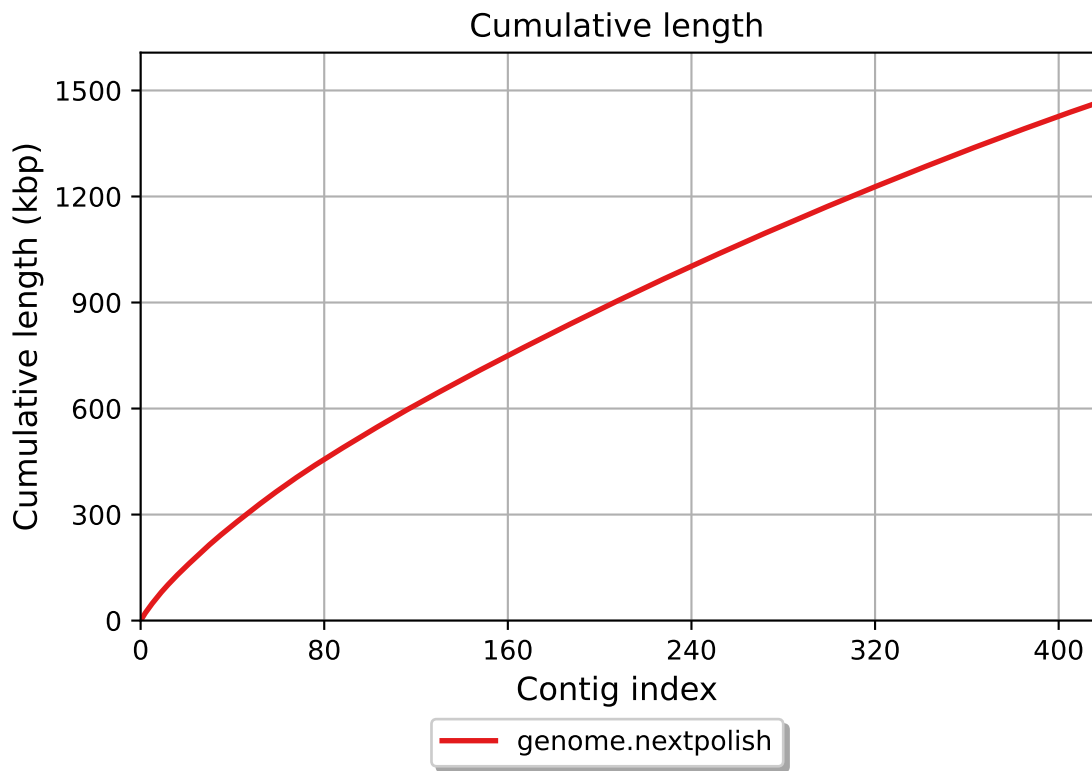
	genome.nextpolish
# contigs (>= 0 bp)	415
# contigs (>= 1000 bp)	415
# contigs (>= 5000 bp)	51
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1460942
Total length (>= 1000 bp)	1460942
Total length (>= 5000 bp)	325367
Total length (>= 10000 bp)	21181
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	415
Largest contig	10940
Total length	1460942
GC (%)	36.32
N50	3371
N90	2531
auN	4008.7
L50	155
L90	354
# 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).

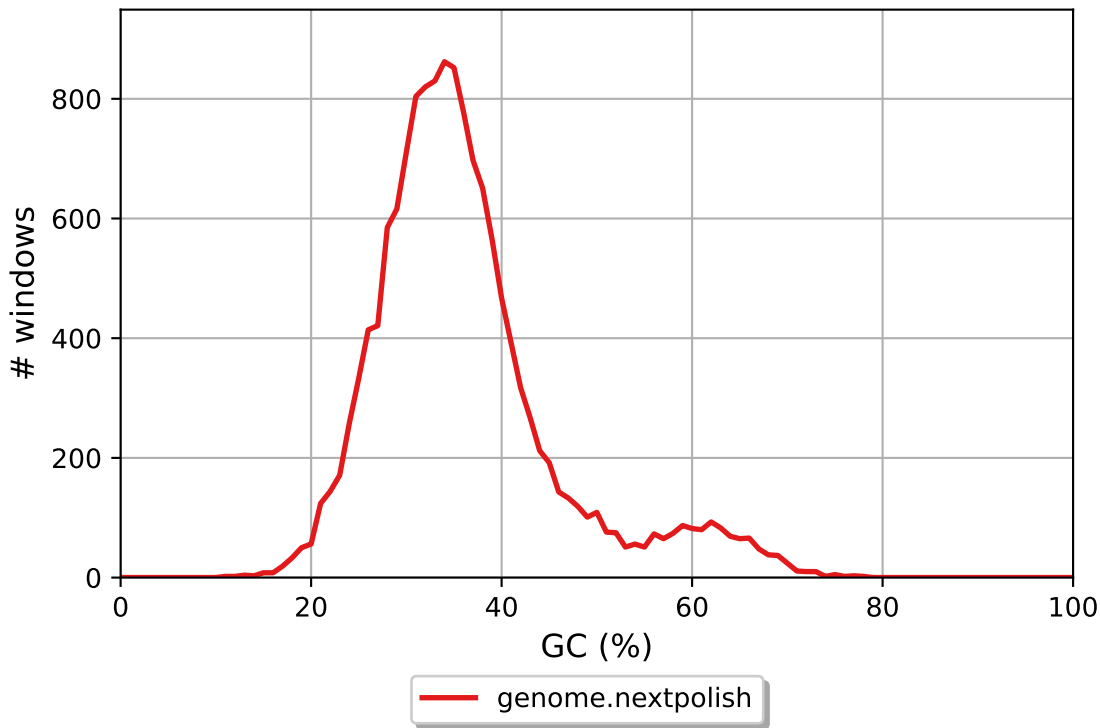
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

