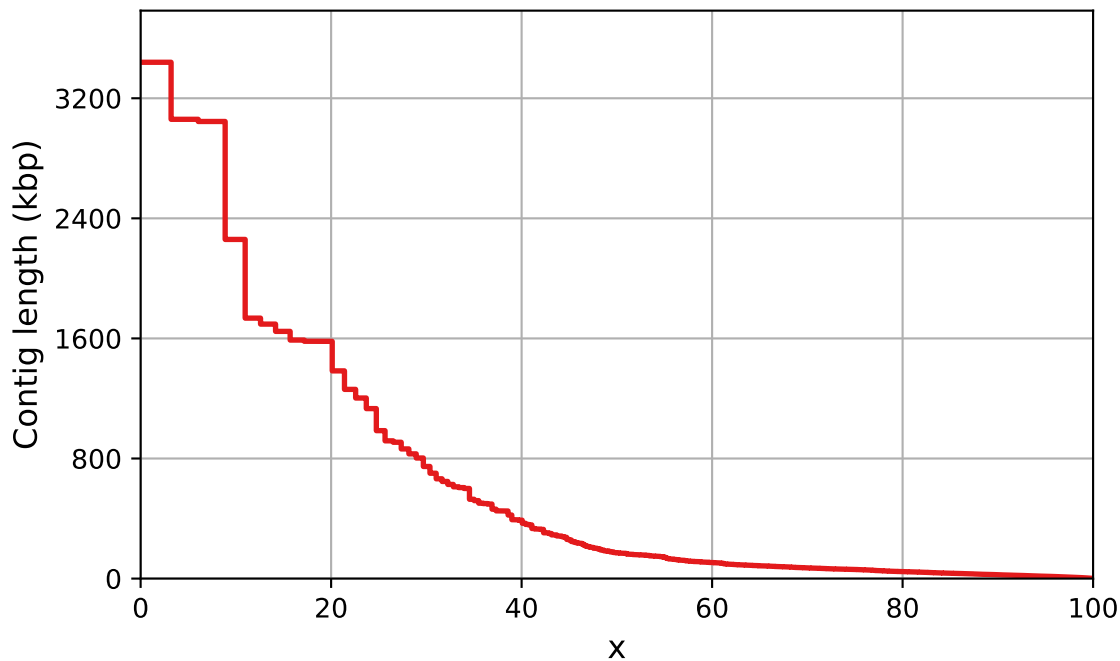


Report

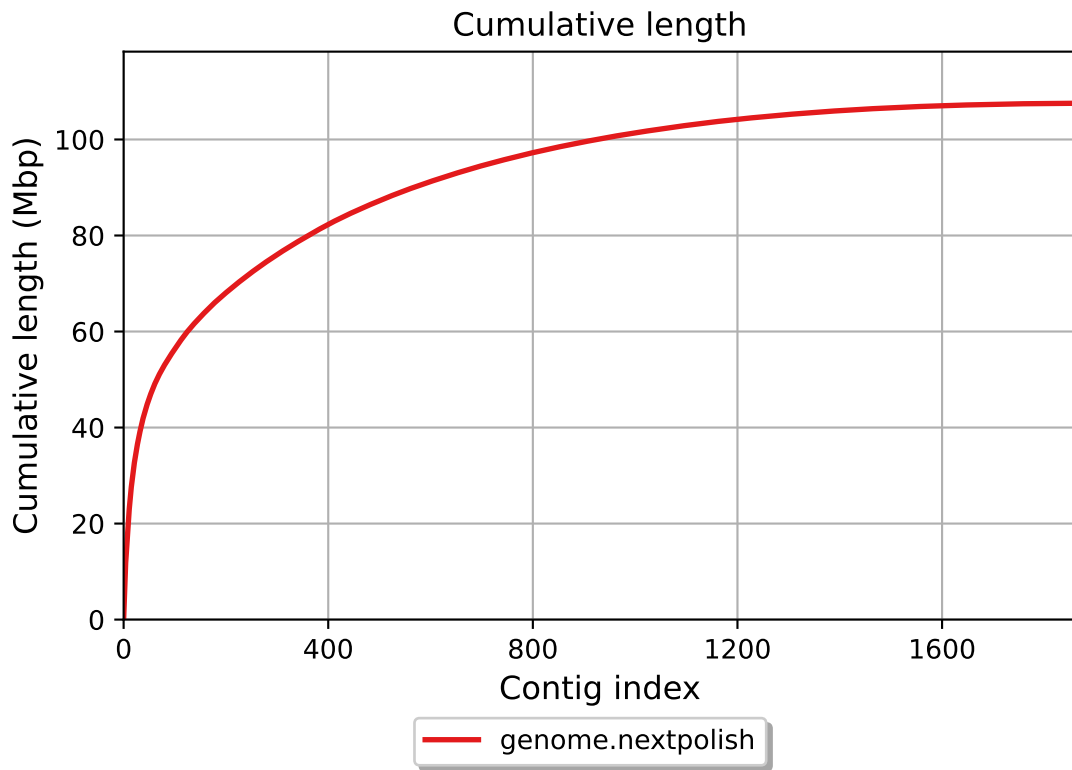
	genome.nextpolish
# contigs (>= 0 bp)	1865
# contigs (>= 1000 bp)	1808
# contigs (>= 5000 bp)	1510
# contigs (>= 10000 bp)	1255
# contigs (>= 25000 bp)	794
# contigs (>= 50000 bp)	438
Total length (>= 0 bp)	107543140
Total length (>= 1000 bp)	107502437
Total length (>= 5000 bp)	106656101
Total length (>= 10000 bp)	104786562
Total length (>= 25000 bp)	97102730
Total length (>= 50000 bp)	84333816
# contigs	1862
Largest contig	3439803
Total length	107542137
GC (%)	47.09
N50	171694
N90	25677
auN	699352.0
L50	85
L90	782
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	273 + 10 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).

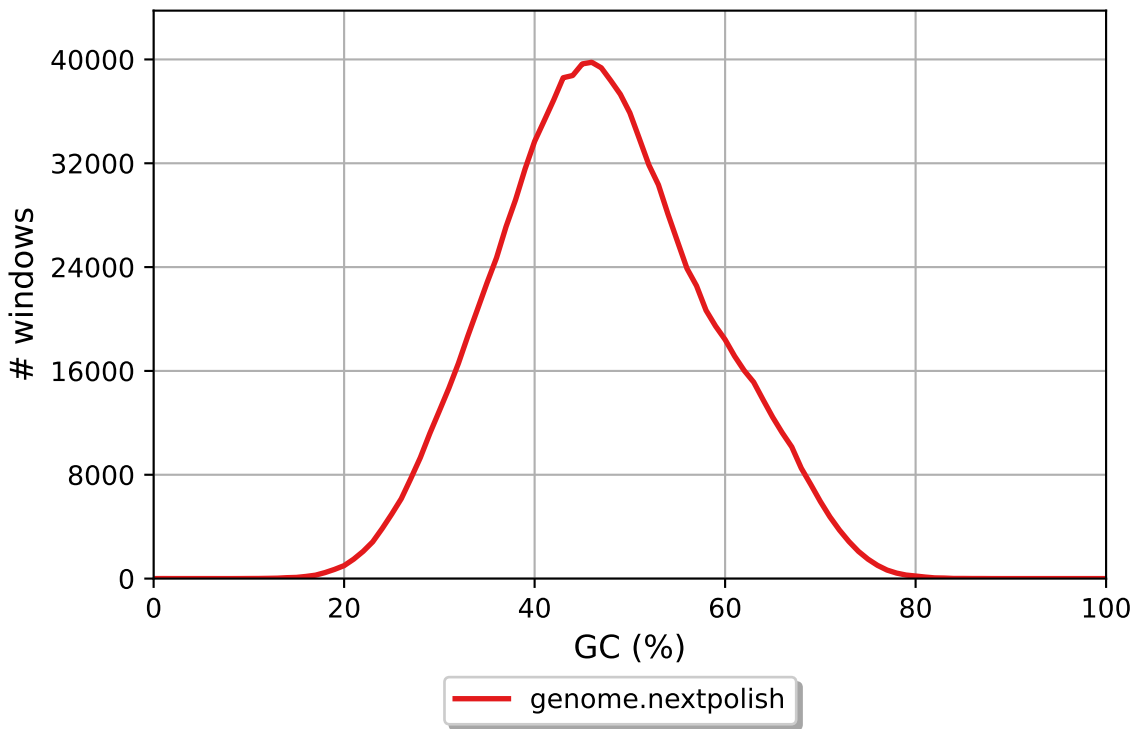
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

