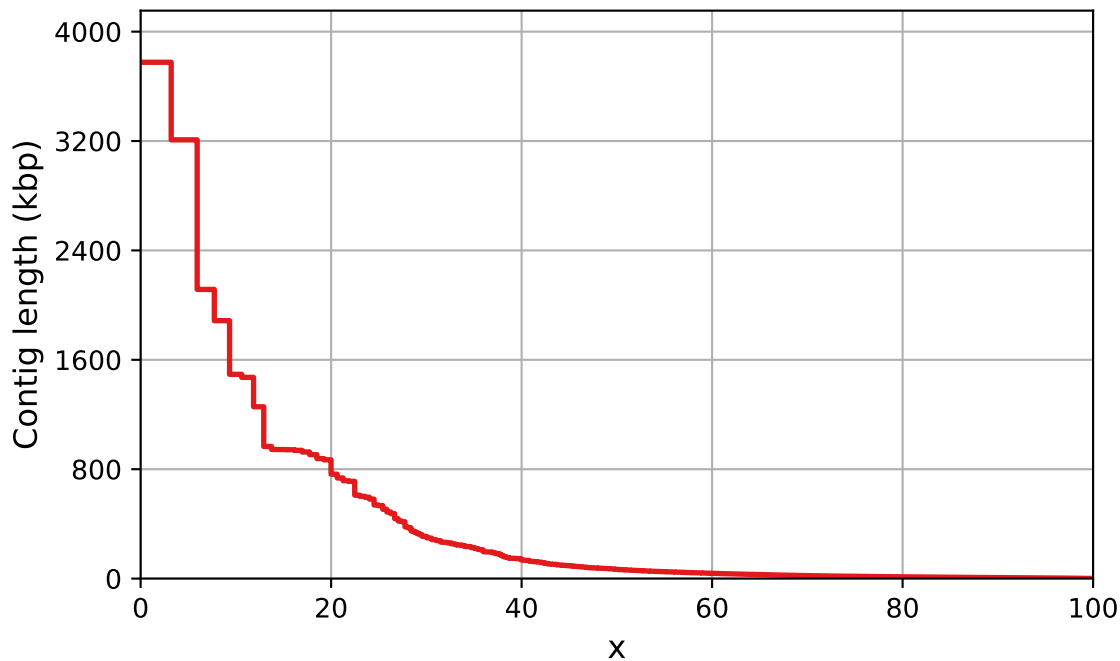


Report

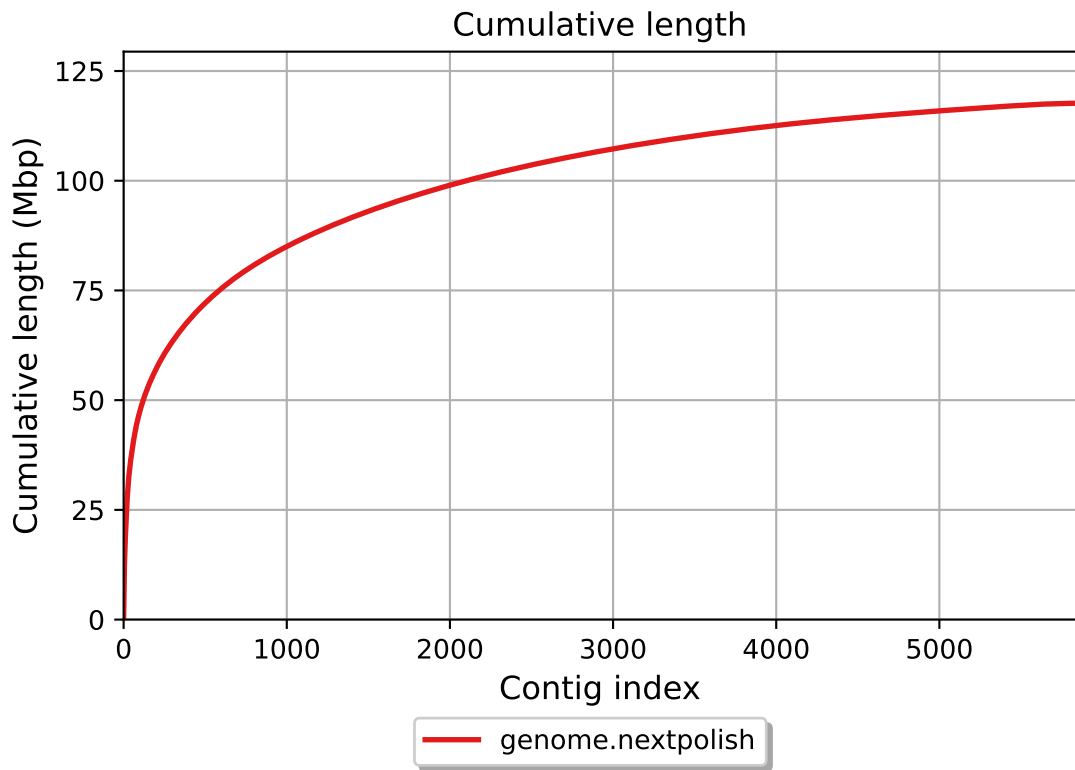
	genome.nextpolish
# contigs (>= 0 bp)	5840
# contigs (>= 1000 bp)	5731
# contigs (>= 5000 bp)	3628
# contigs (>= 10000 bp)	2076
# contigs (>= 25000 bp)	752
# contigs (>= 50000 bp)	328
Total length (>= 0 bp)	117650784
Total length (>= 1000 bp)	117575364
Total length (>= 5000 bp)	110876247
Total length (>= 10000 bp)	99760871
Total length (>= 25000 bp)	79653932
Total length (>= 50000 bp)	64881004
# contigs	5838
Largest contig	3776008
Total length	117649813
GC (%)	45.75
N50	67747
N90	7169
auN	489512.2
L50	223
L90	2799
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	275 + 21 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).

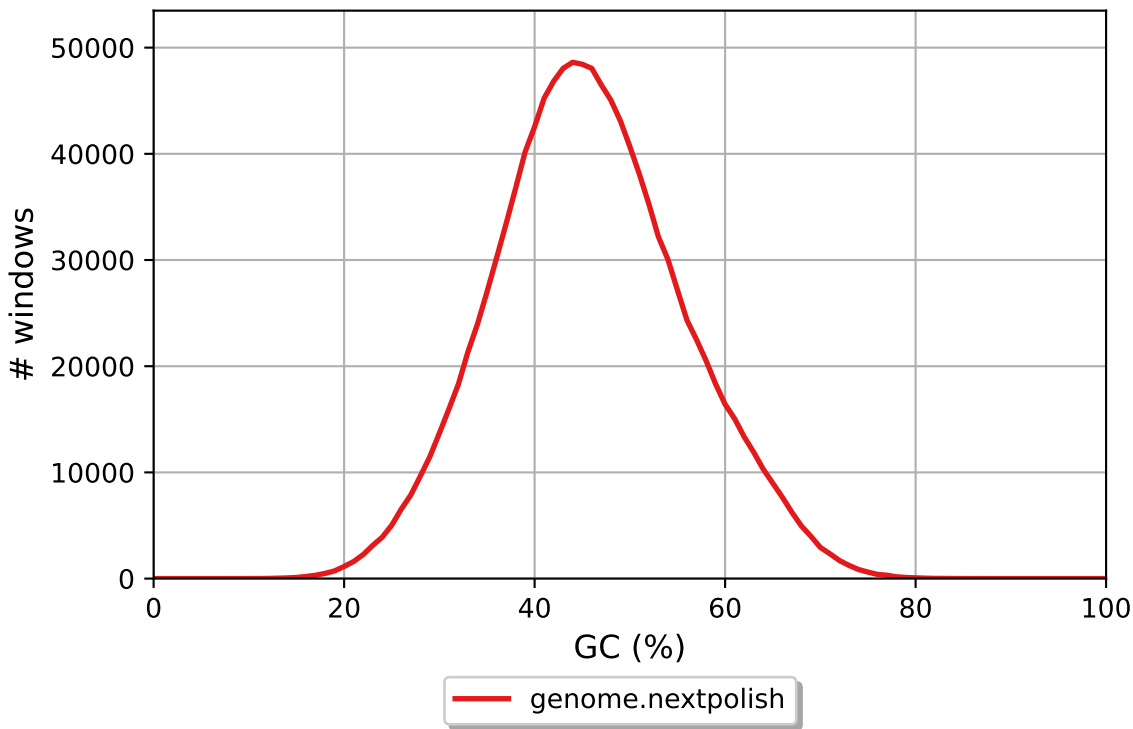
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

