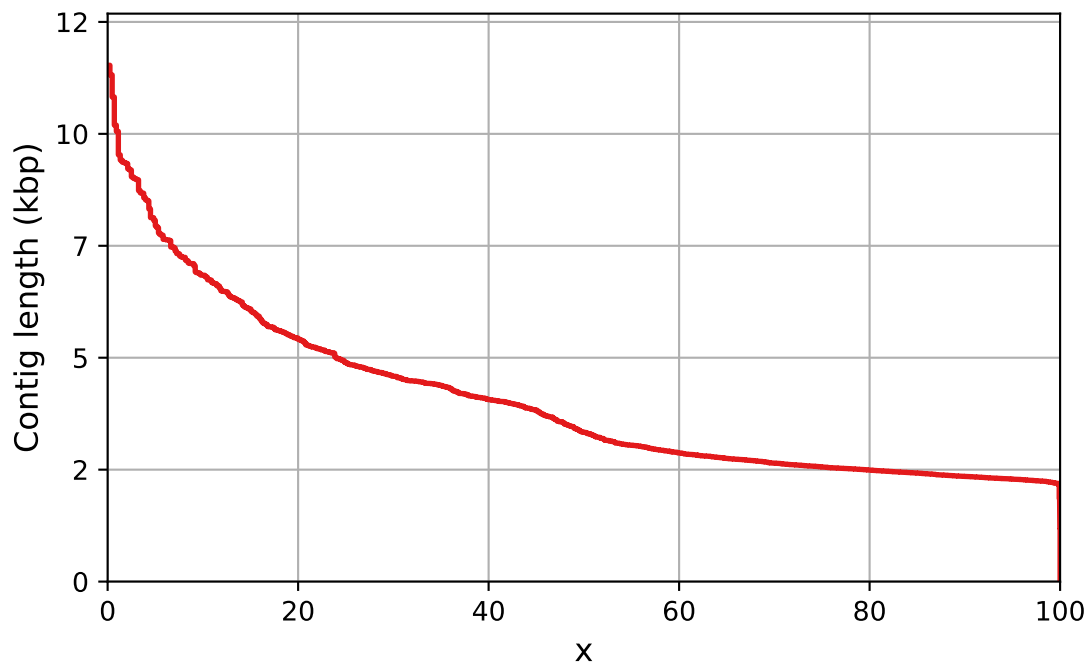


Report

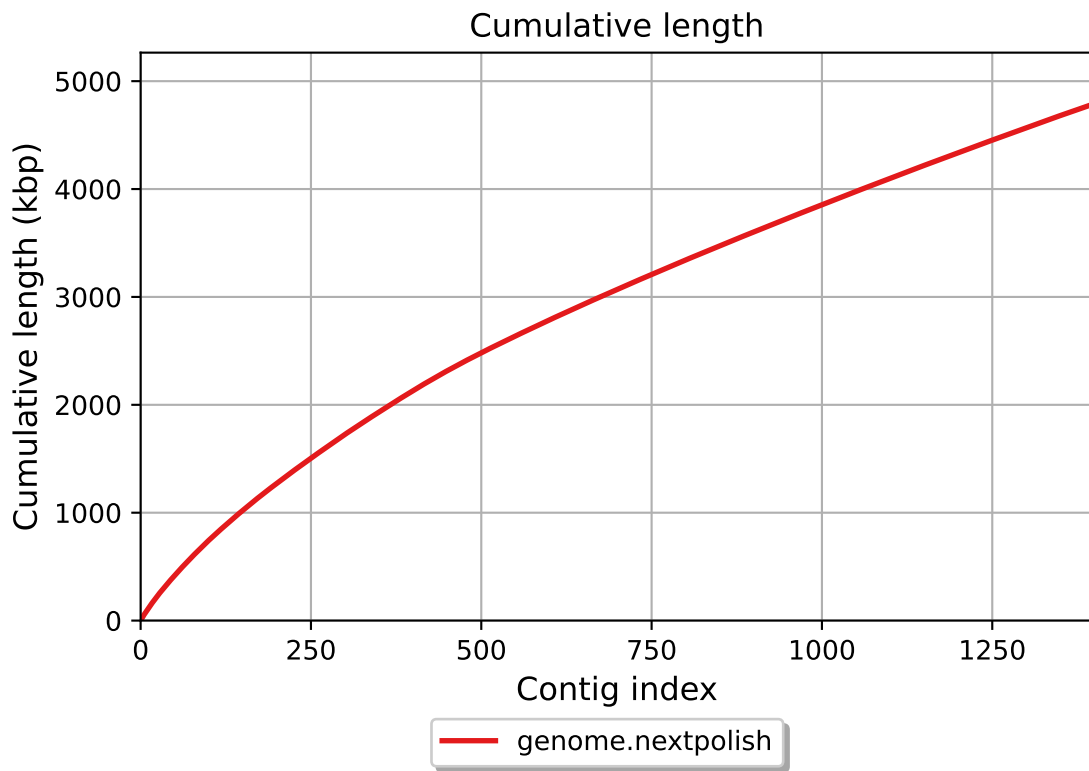
	genome.nextpolish
# contigs (>= 0 bp)	1398
# contigs (>= 1000 bp)	1398
# contigs (>= 5000 bp)	175
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4785812
Total length (>= 1000 bp)	4785812
Total length (>= 5000 bp)	1151067
Total length (>= 10000 bp)	53921
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1398
Largest contig	11531
Total length	4785812
GC (%)	50.63
N50	3334
N90	2349
auN	4053.7
L50	473
L90	1188
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

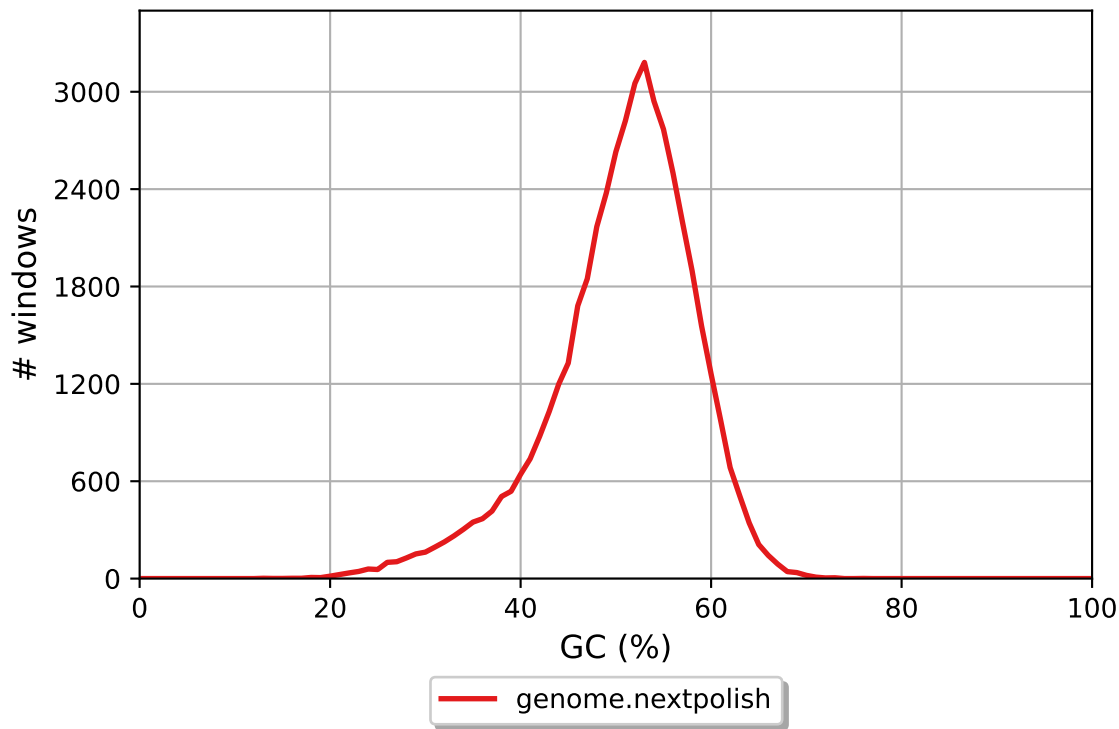
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

