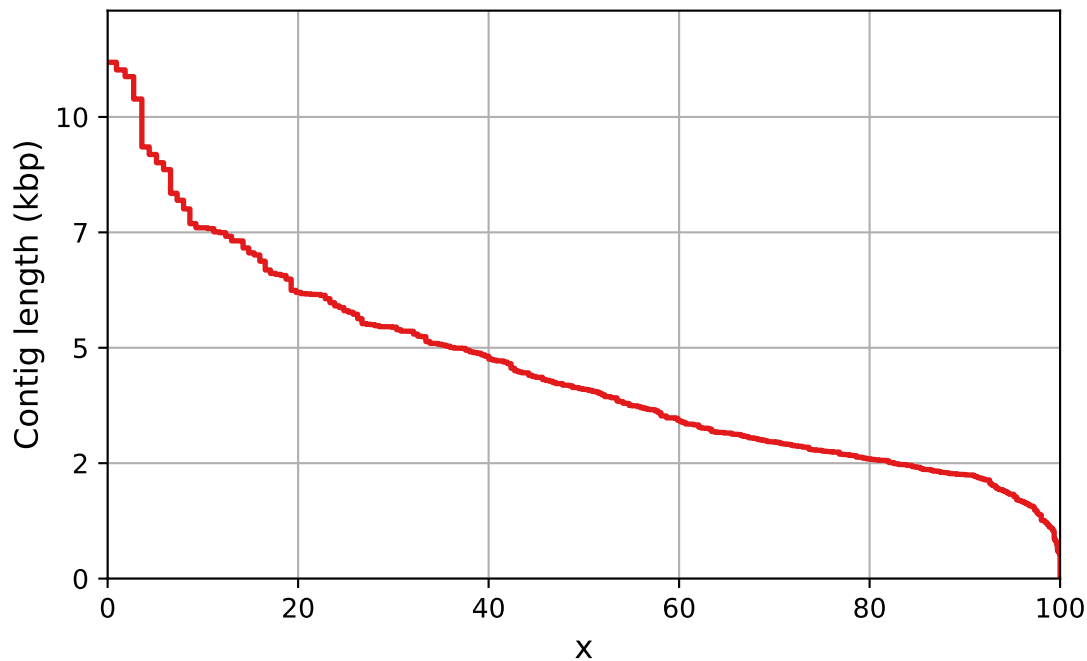


Report

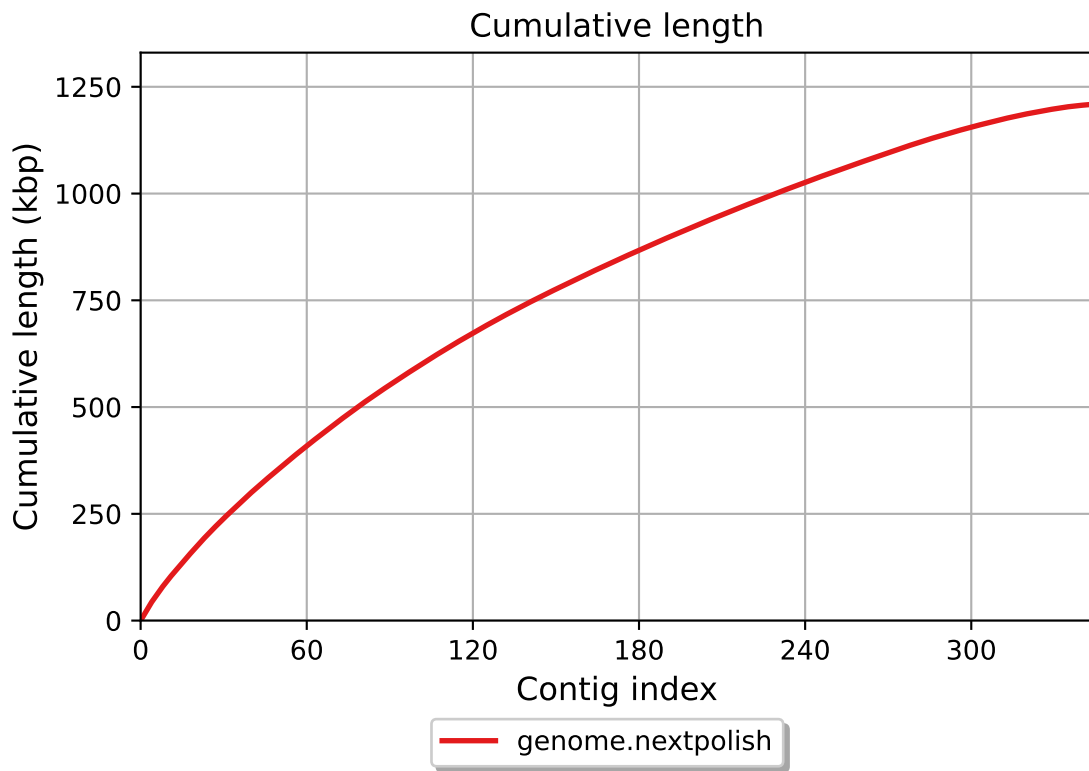
	genome.nextpolish
# contigs (>= 0 bp)	345
# contigs (>= 1000 bp)	333
# contigs (>= 5000 bp)	66
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1209572
Total length (>= 1000 bp)	1201538
Total length (>= 5000 bp)	439529
Total length (>= 10000 bp)	43465
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	344
Largest contig	11184
Total length	1209073
GC (%)	40.03
N50	4102
N90	2254
auN	4556.3
L50	103
L90	267
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

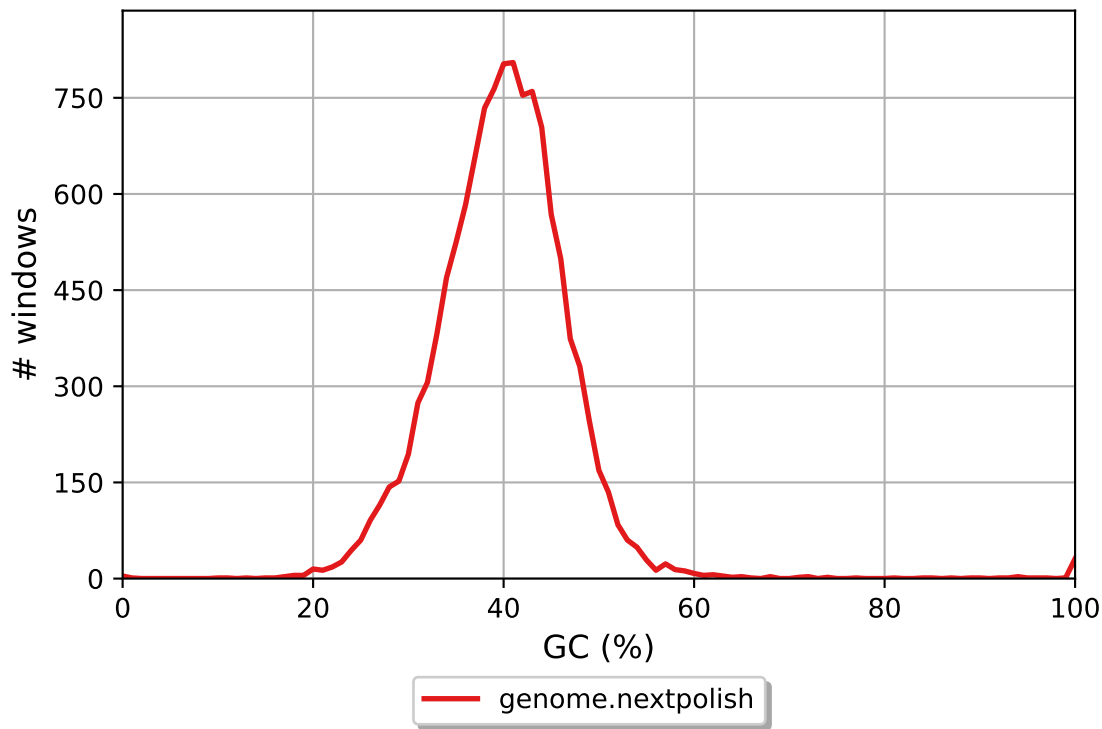
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

