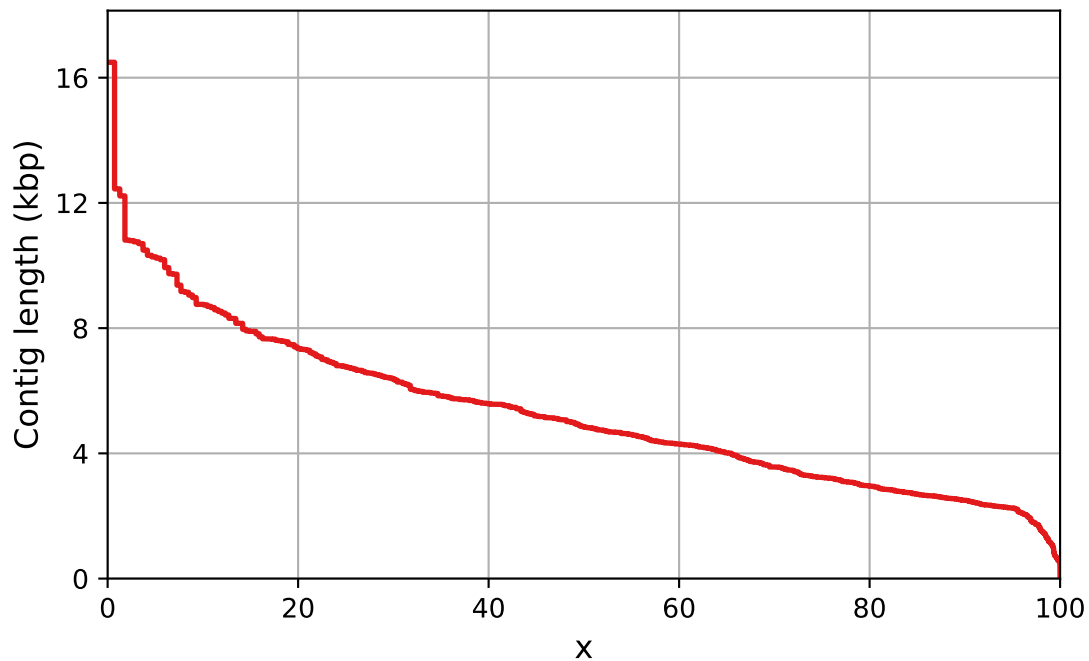


Report

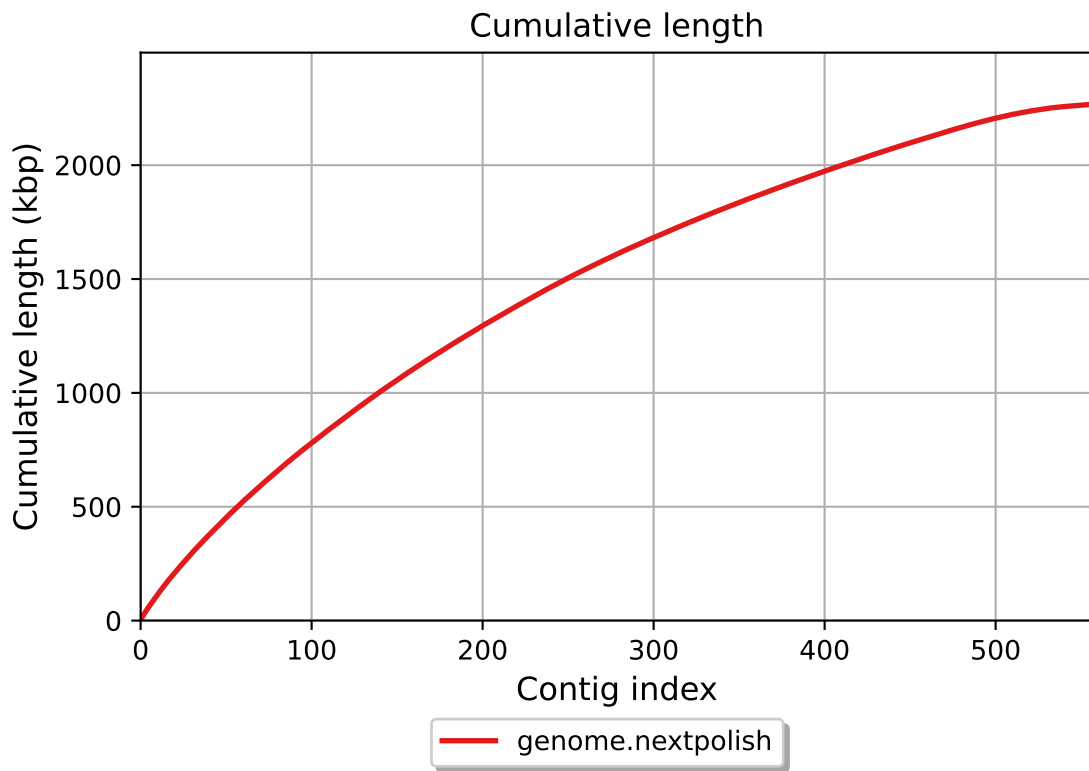
	genome.nextpolish
# contigs (>= 0 bp)	558
# contigs (>= 1000 bp)	532
# contigs (>= 5000 bp)	159
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2267946
Total length (>= 1000 bp)	2251077
Total length (>= 5000 bp)	1102328
Total length (>= 10000 bp)	135744
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	557
Largest contig	16492
Total length	2267450
GC (%)	40.26
N50	4844
N90	2498
auN	5309.9
L50	166
L90	427
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

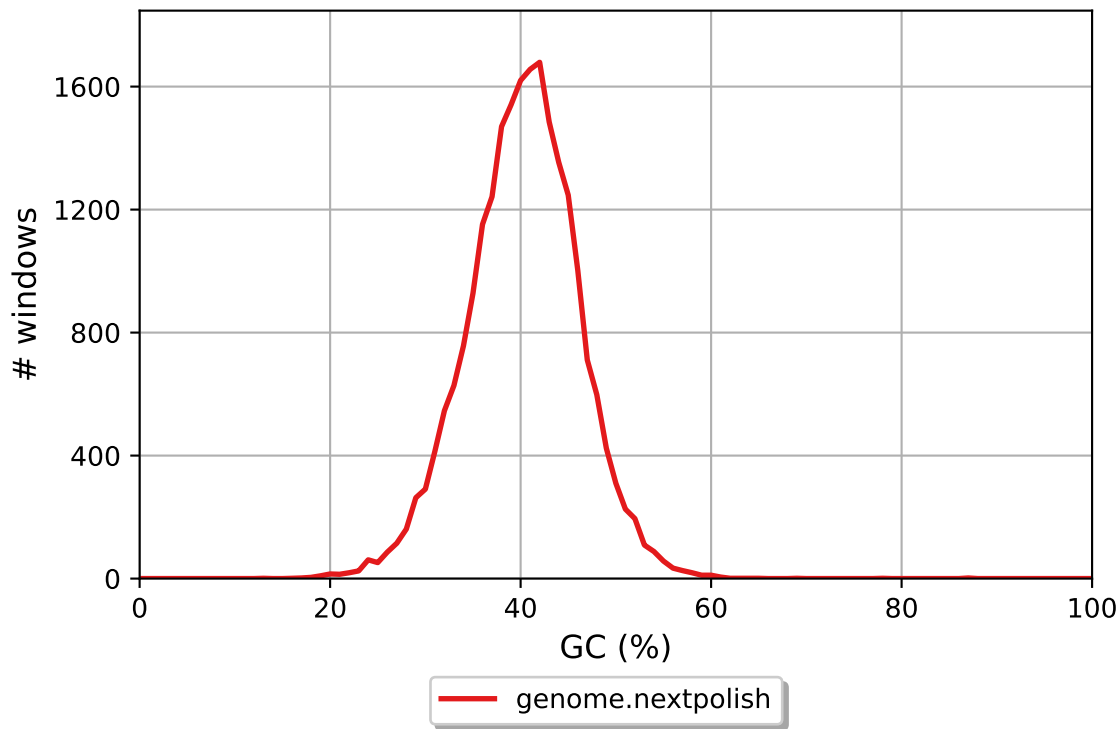
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

