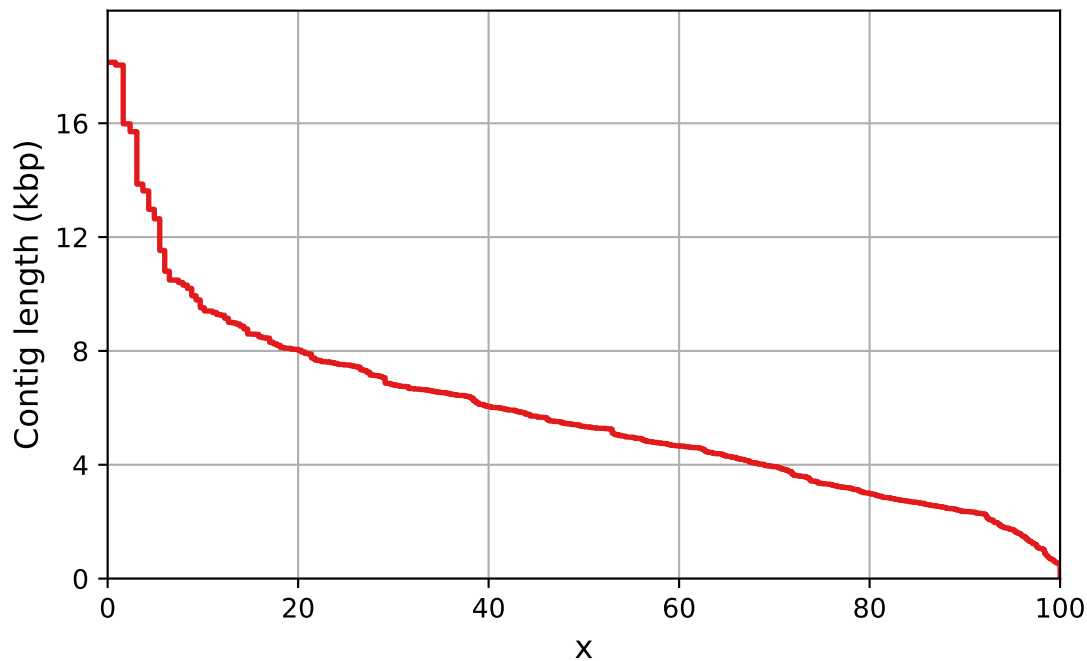


Report

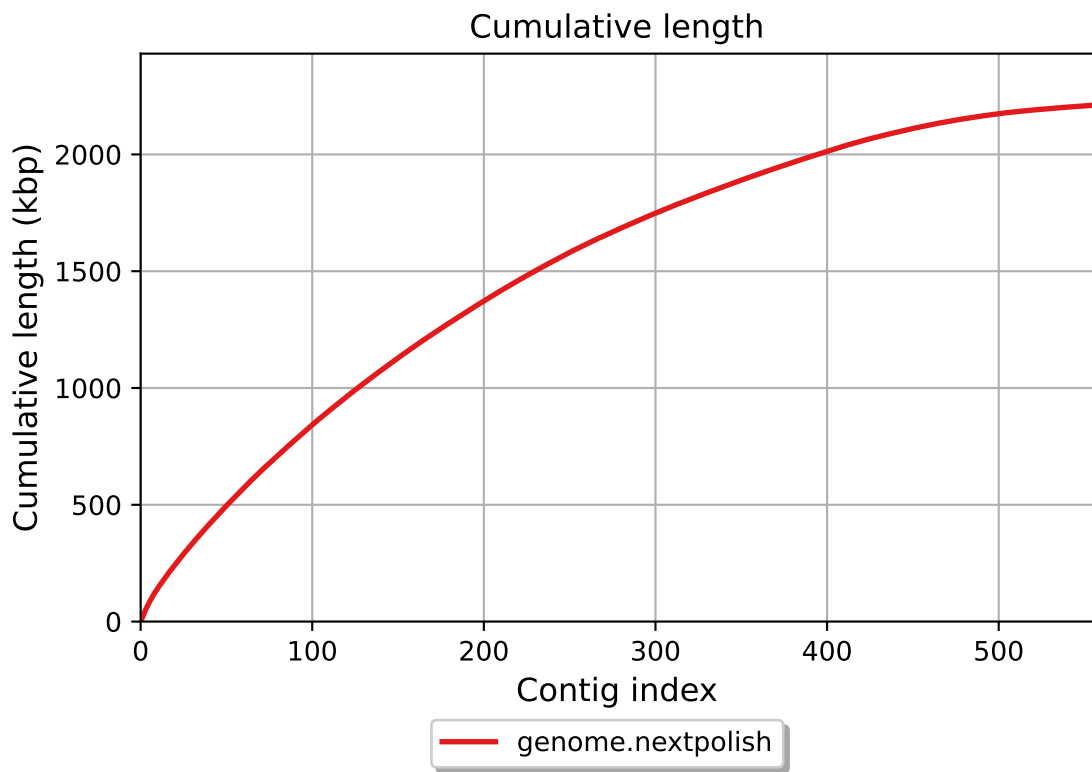
	genome.nextpolish
# contigs (>= 0 bp)	557
# contigs (>= 1000 bp)	499
# contigs (>= 5000 bp)	164
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2211028
Total length (>= 1000 bp)	2173573
Total length (>= 5000 bp)	1201119
Total length (>= 10000 bp)	195191
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	555
Largest contig	18142
Total length	2210374
GC (%)	33.14
N50	5346
N90	2359
auN	5850.8
L50	146
L90	390
# 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).

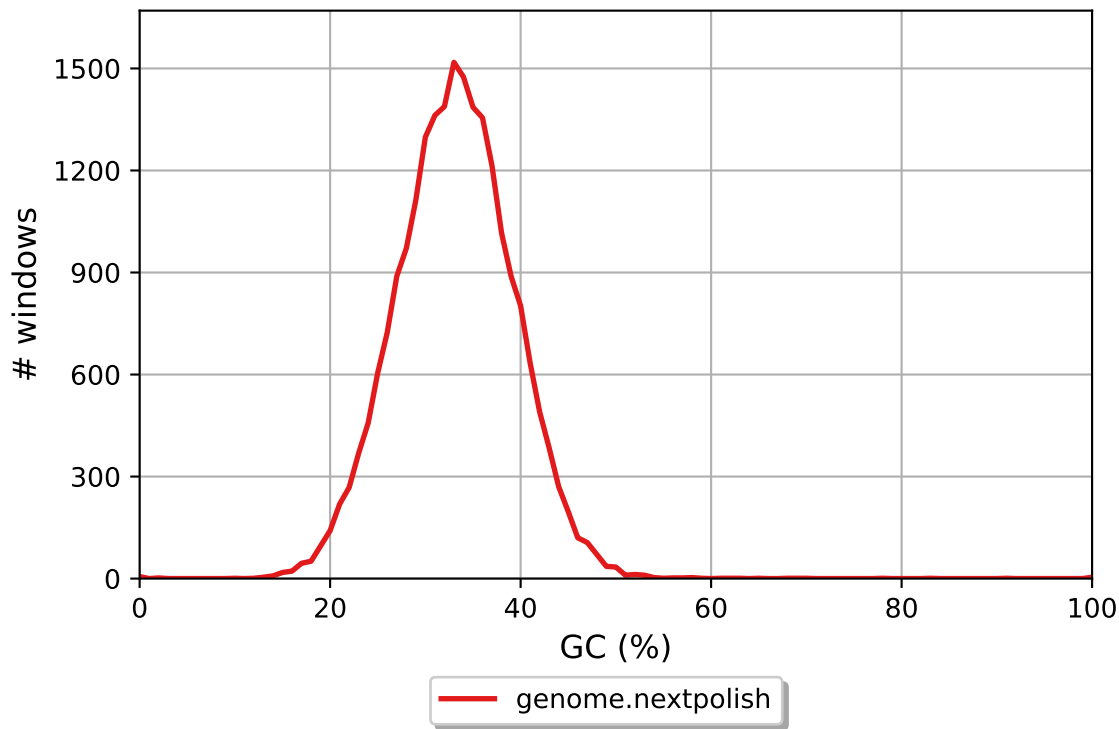
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

