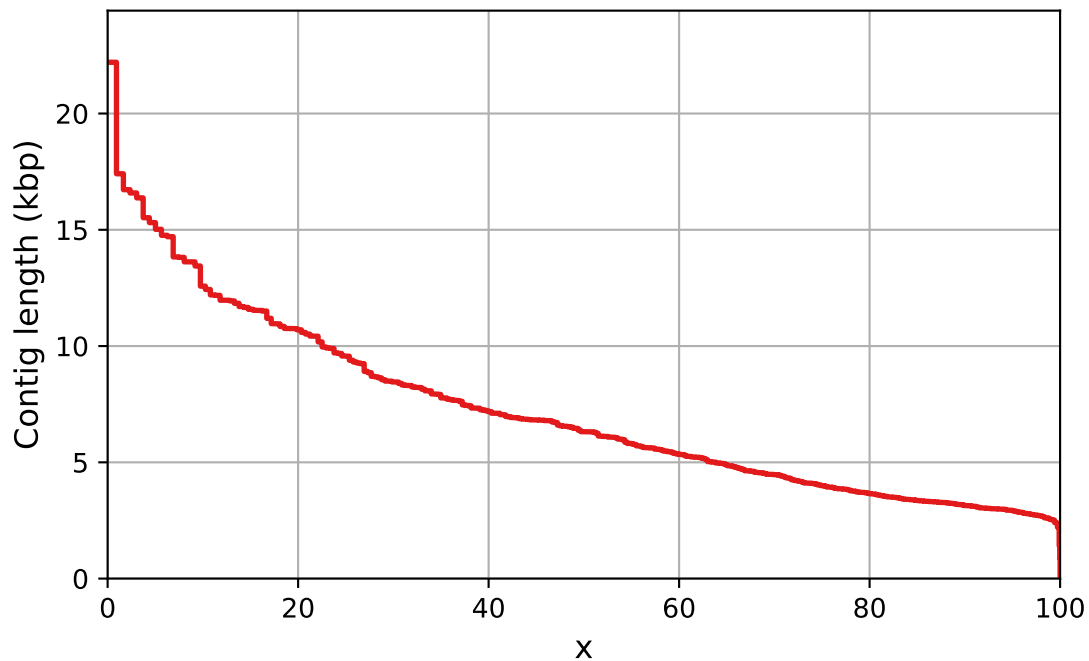


Report

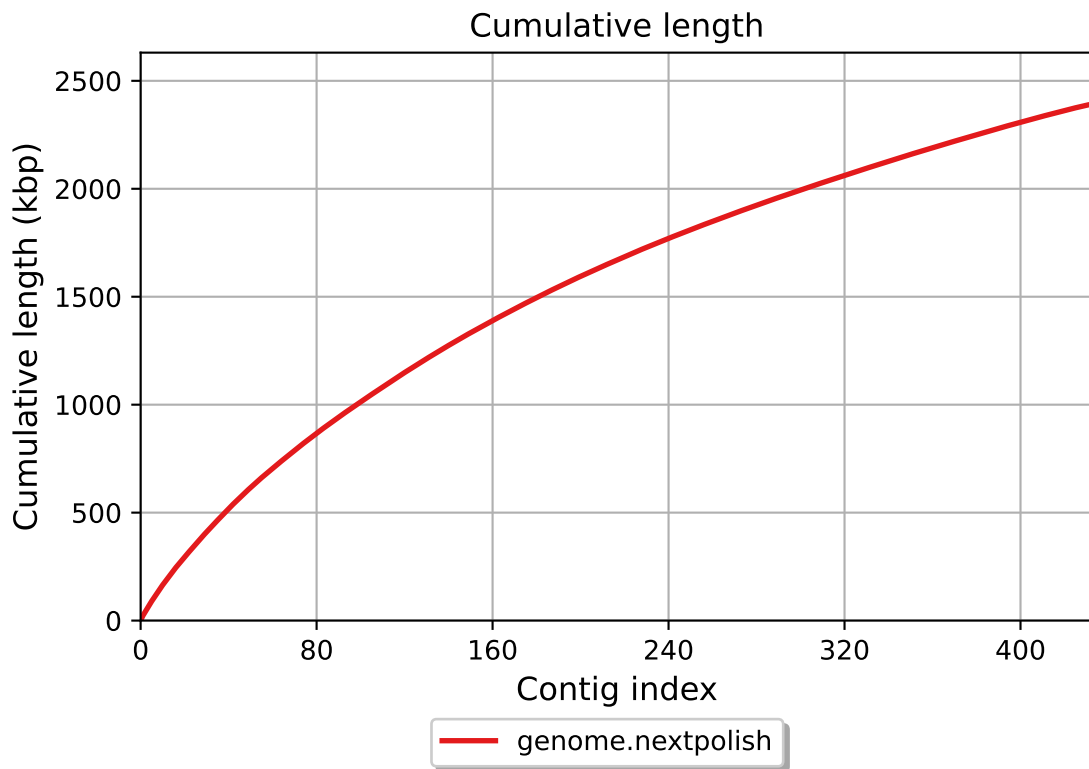
	genome.nextpolish
# contigs (>= 0 bp)	433
# contigs (>= 1000 bp)	432
# contigs (>= 5000 bp)	186
# contigs (>= 10000 bp)	42
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2391424
Total length (>= 1000 bp)	2390542
Total length (>= 5000 bp)	1526429
Total length (>= 10000 bp)	538612
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	433
Largest contig	22204
Total length	2391424
GC (%)	33.55
N50	6316
N90	3142
auN	7245.2
L50	128
L90	348
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 2 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).

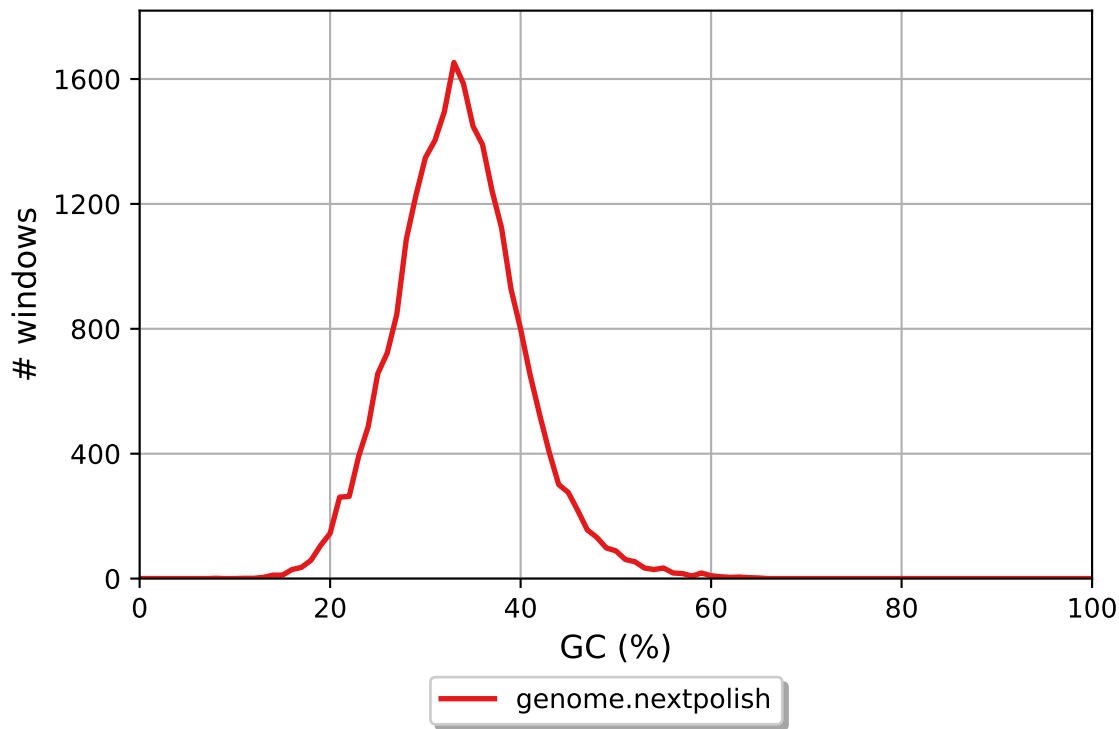
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

