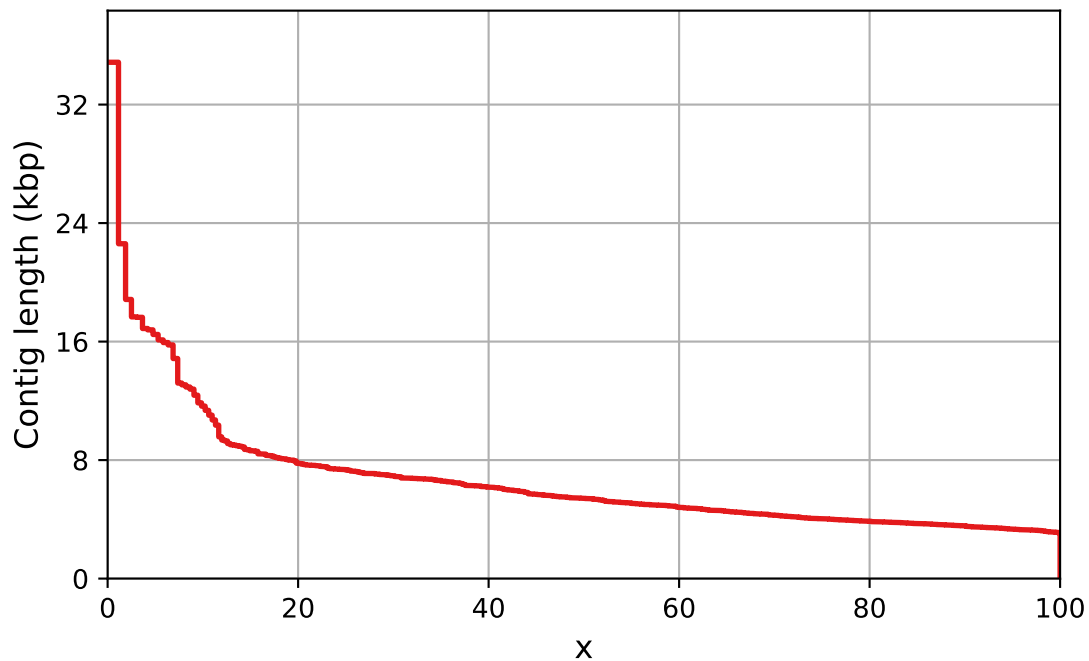


Report

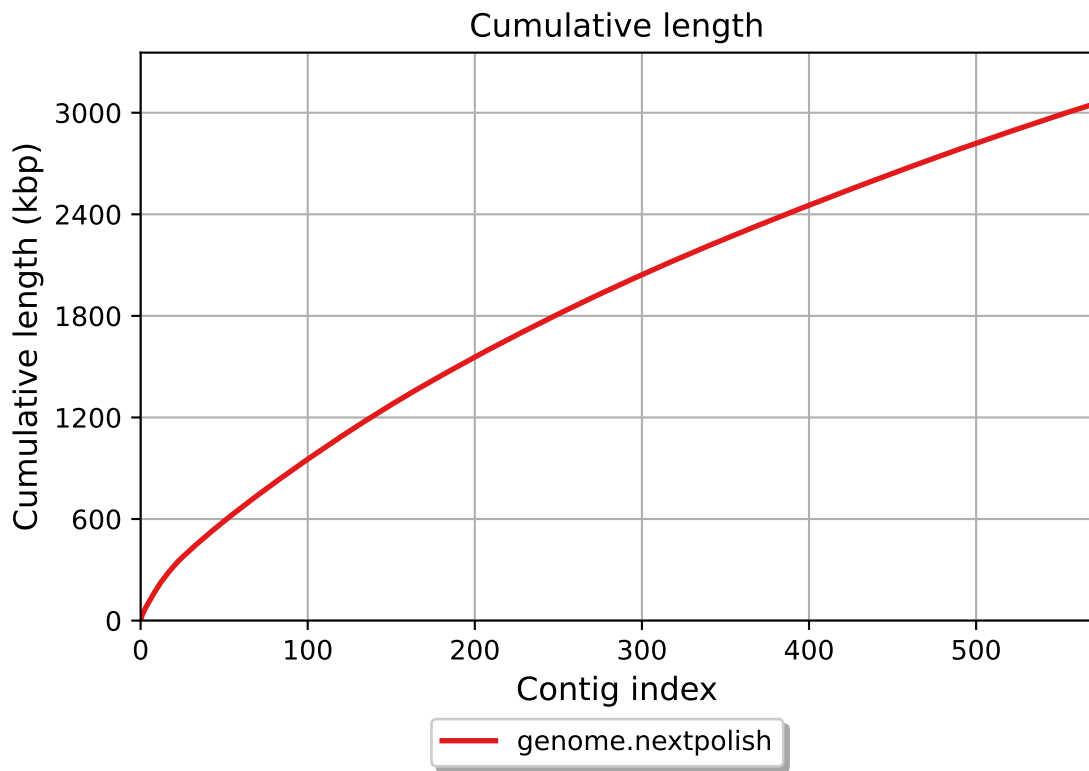
	genome.nextpolish
# contigs (>= 0 bp)	570
# contigs (>= 1000 bp)	570
# contigs (>= 5000 bp)	234
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3050082
Total length (>= 1000 bp)	3050082
Total length (>= 5000 bp)	1731948
Total length (>= 10000 bp)	355785
Total length (>= 25000 bp)	34856
Total length (>= 50000 bp)	0
# contigs	570
Largest contig	34856
Total length	3050082
GC (%)	59.19
N50	5412
N90	3565
auN	6755.2
L50	195
L90	479
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 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).

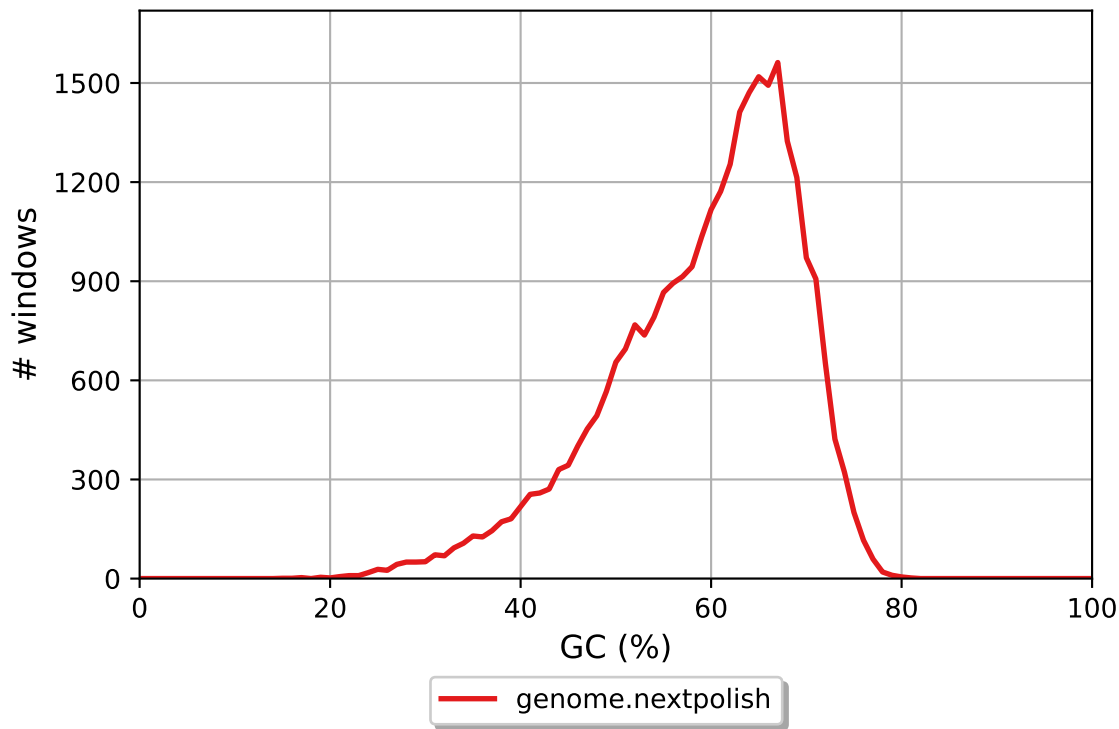
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

