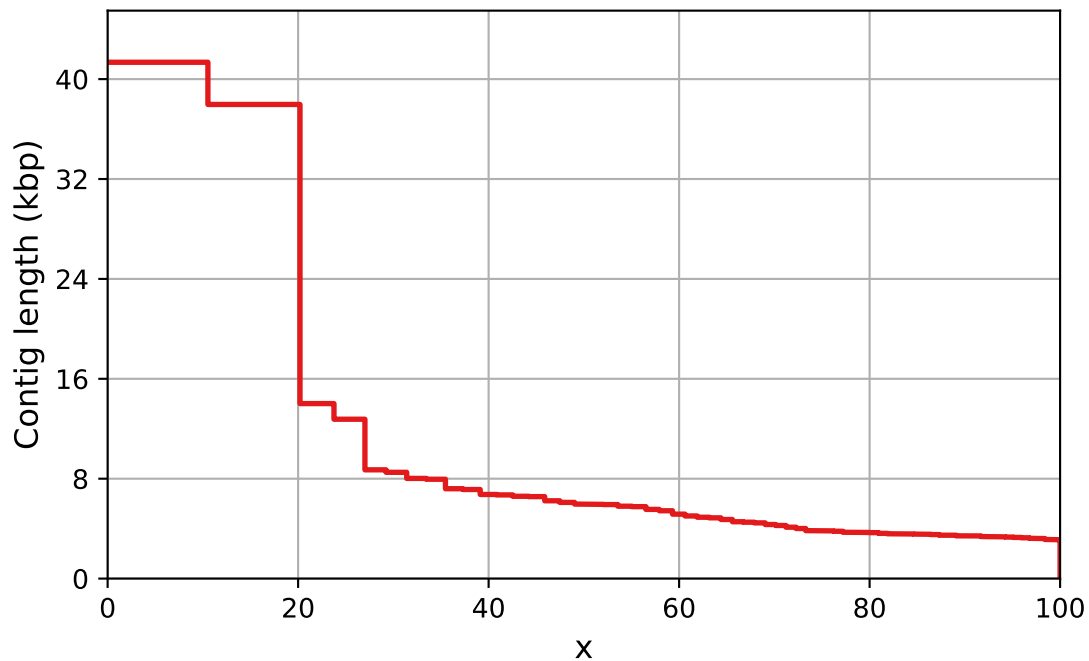


Report

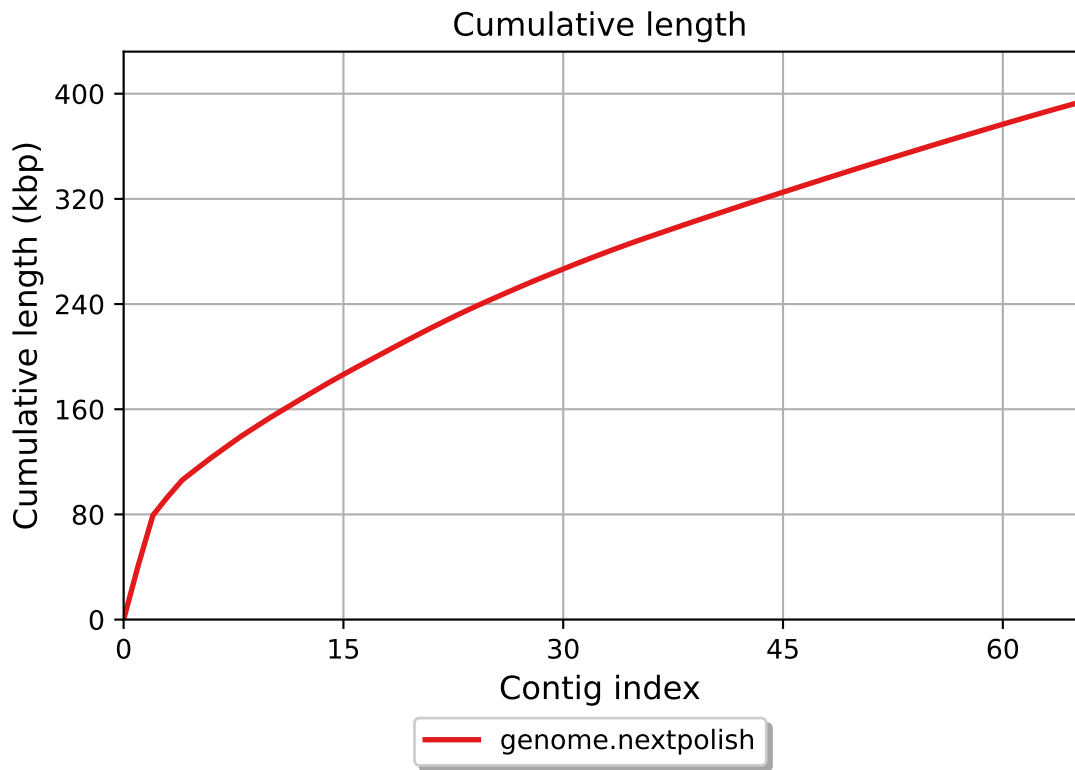
	genome.nextpolish
# contigs (>= 0 bp)	65
# contigs (>= 1000 bp)	65
# contigs (>= 5000 bp)	25
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	392733
Total length (>= 1000 bp)	392733
Total length (>= 5000 bp)	243100
Total length (>= 10000 bp)	106097
Total length (>= 25000 bp)	79320
Total length (>= 50000 bp)	0
# contigs	65
Largest contig	41349
Total length	392733
GC (%)	53.18
N50	5967
N90	3423
auN	12721.8
L50	17
L90	54
# 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).

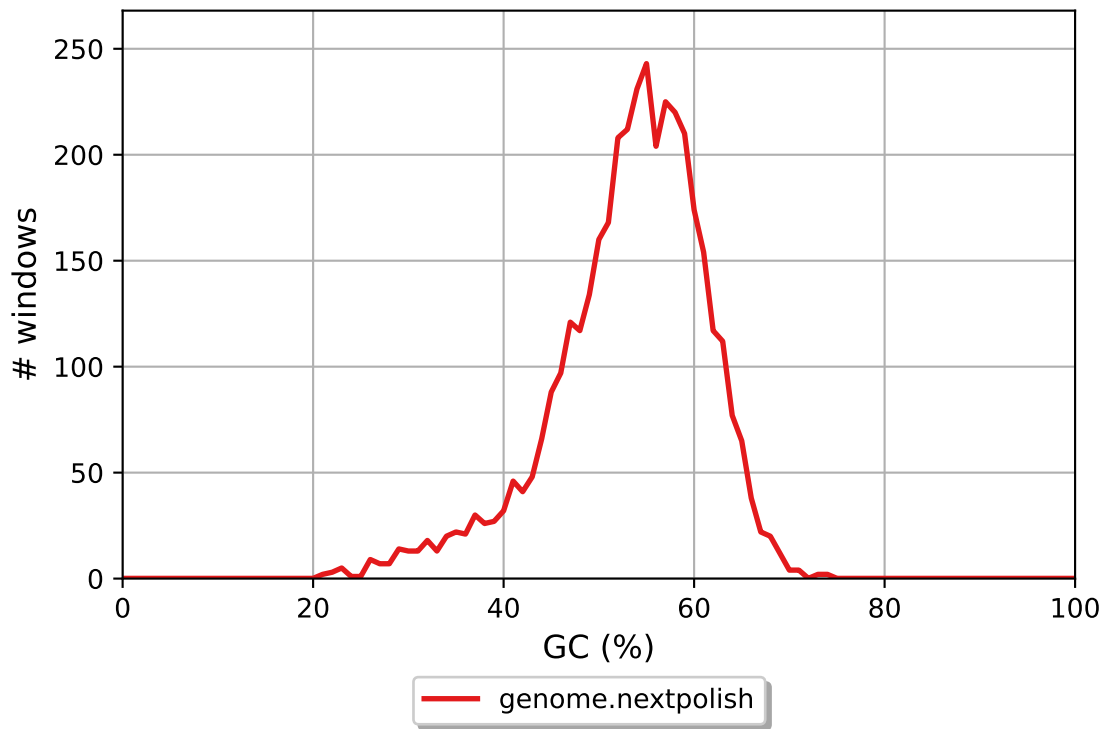
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

