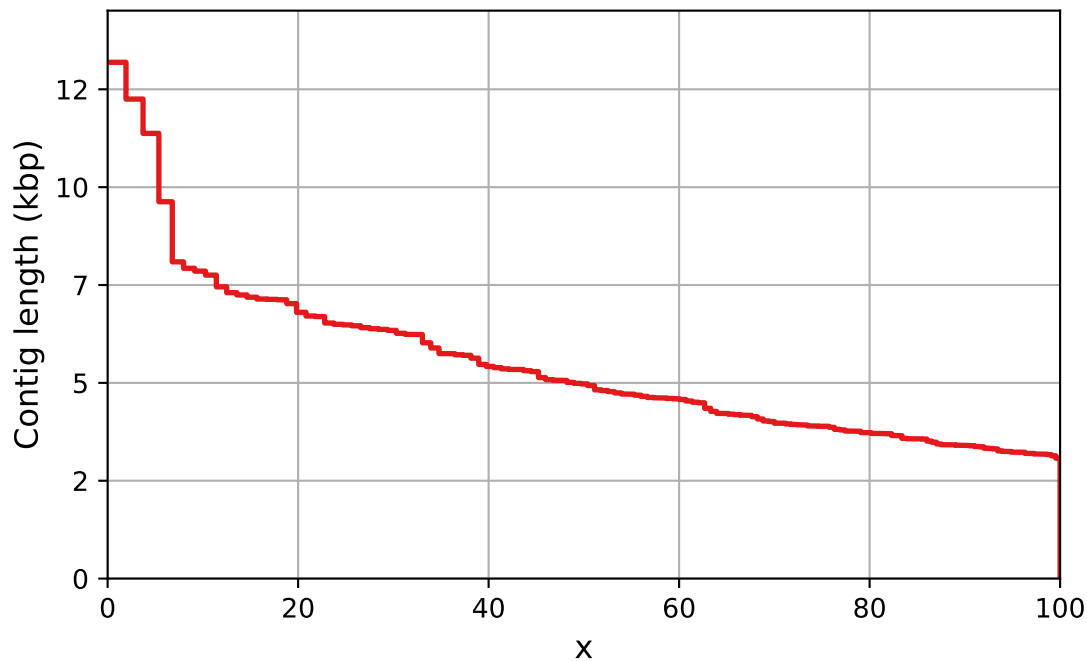


Report

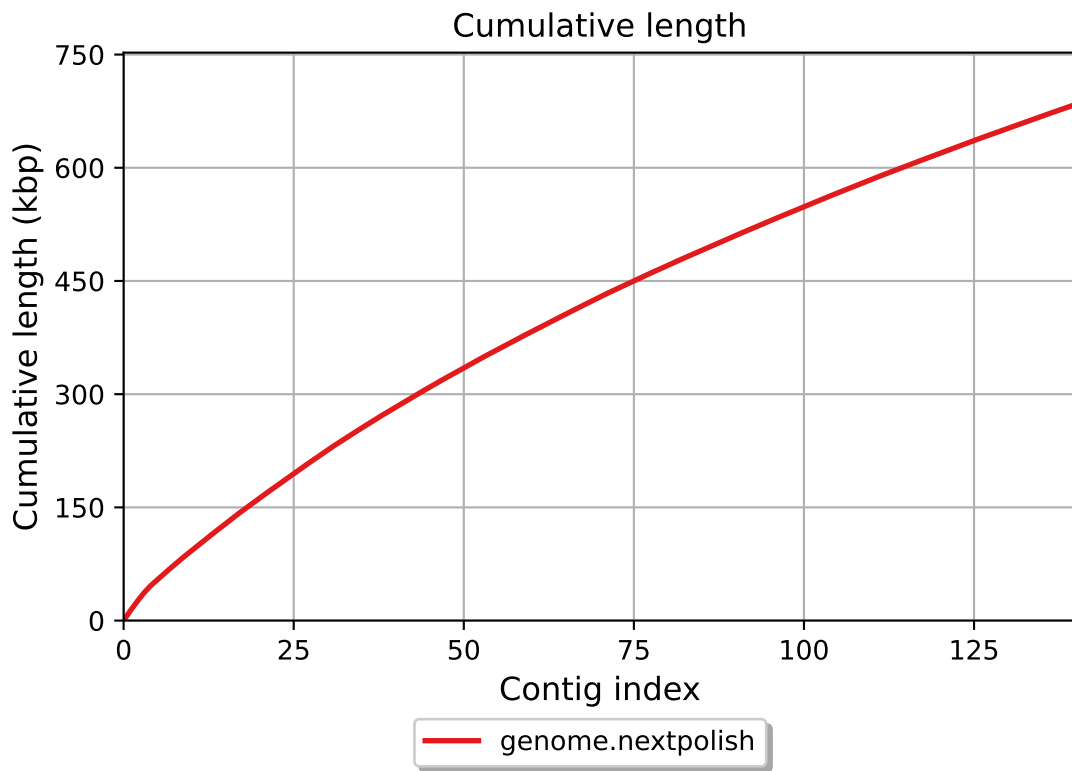
	genome.nextpolish
# contigs (>= 0 bp)	140
# contigs (>= 1000 bp)	140
# contigs (>= 5000 bp)	50
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	684131
Total length (>= 1000 bp)	684131
Total length (>= 5000 bp)	334849
Total length (>= 10000 bp)	36814
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	140
Largest contig	13190
Total length	684131
GC (%)	50.52
N50	4974
N90	3405
auN	5510.6
L50	52
L90	119
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

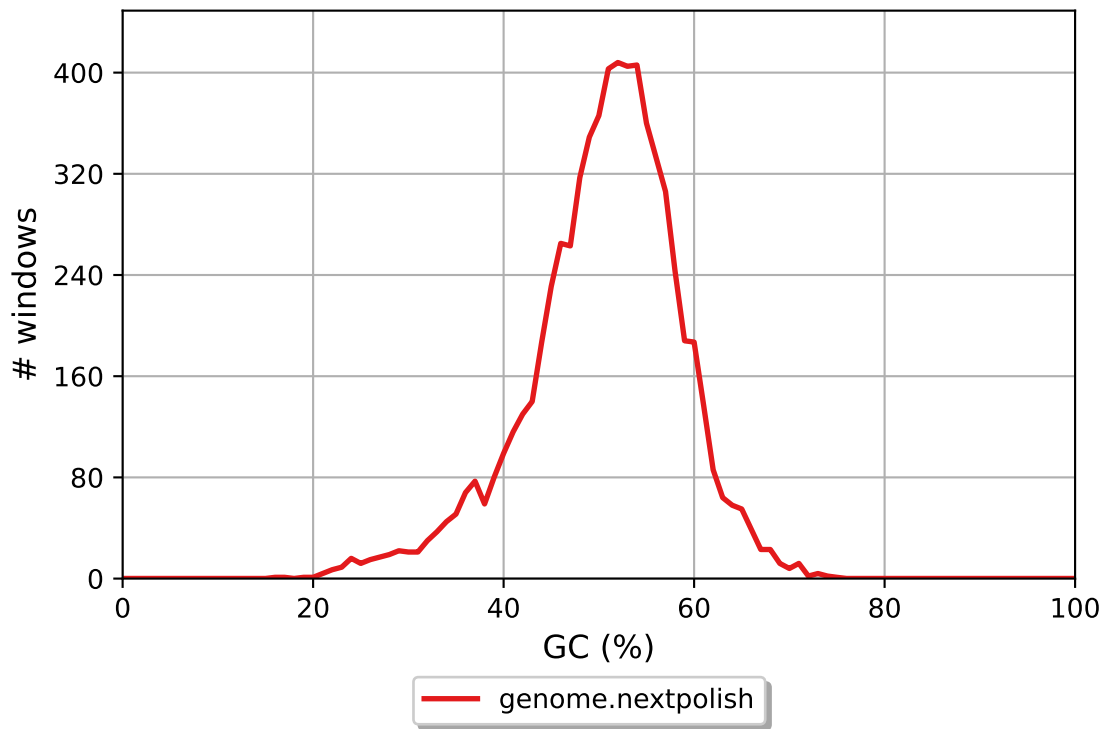
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

