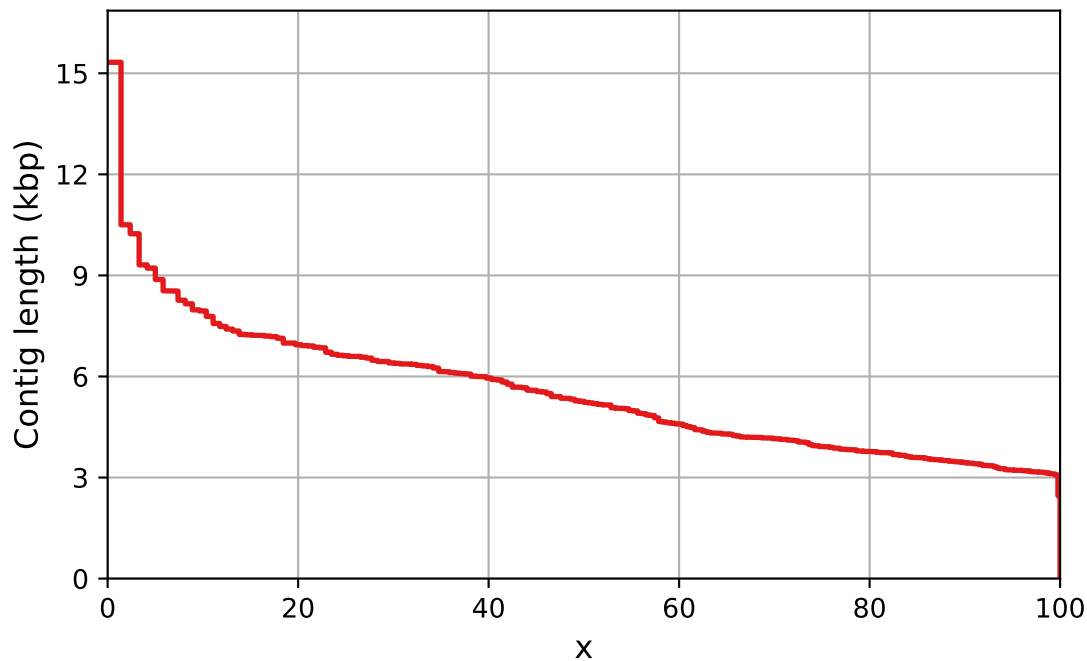


Report

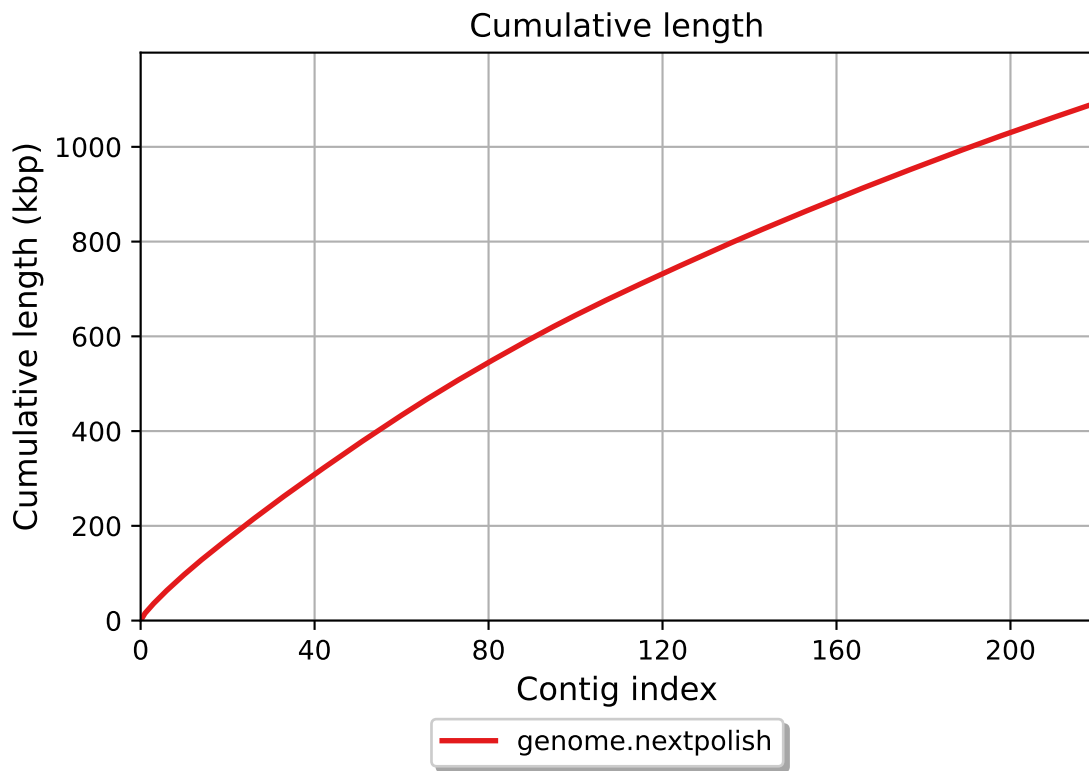
	genome.nextpolish
# contigs (>= 0 bp)	219
# contigs (>= 1000 bp)	219
# contigs (>= 5000 bp)	90
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1089876
Total length (>= 1000 bp)	1089876
Total length (>= 5000 bp)	596321
Total length (>= 10000 bp)	36067
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	219
Largest contig	15327
Total length	1089876
GC (%)	50.86
N50	5263
N90	3441
auN	5566.8
L50	80
L90	186
# 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).

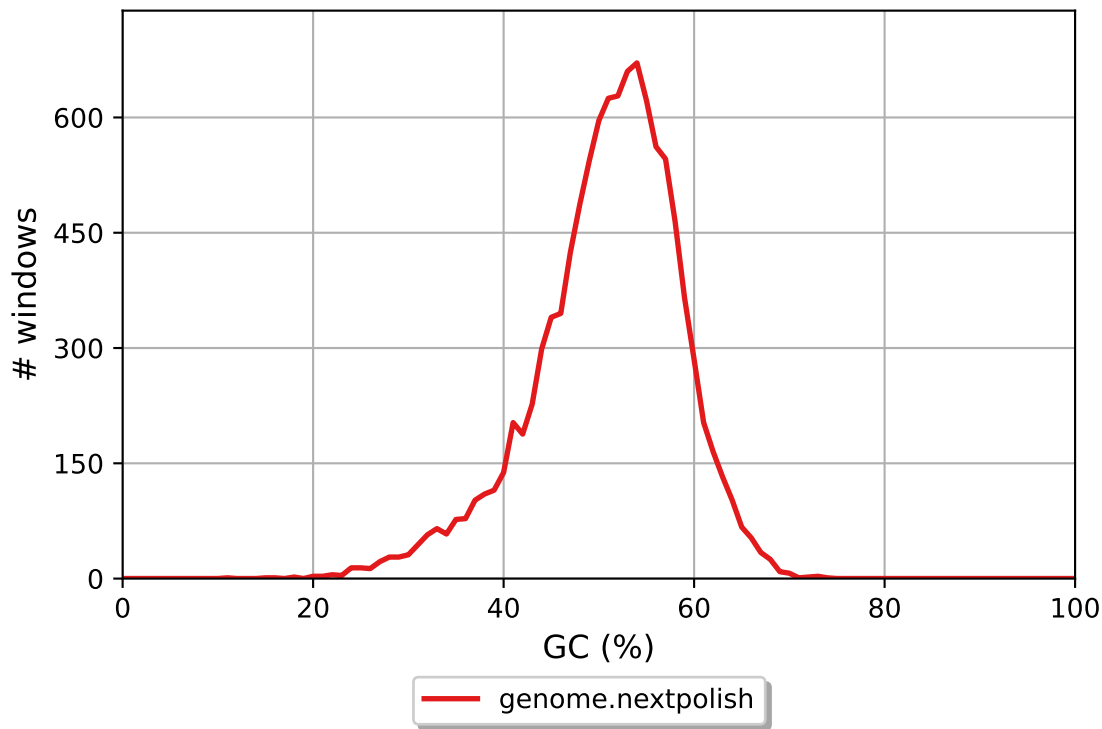
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

