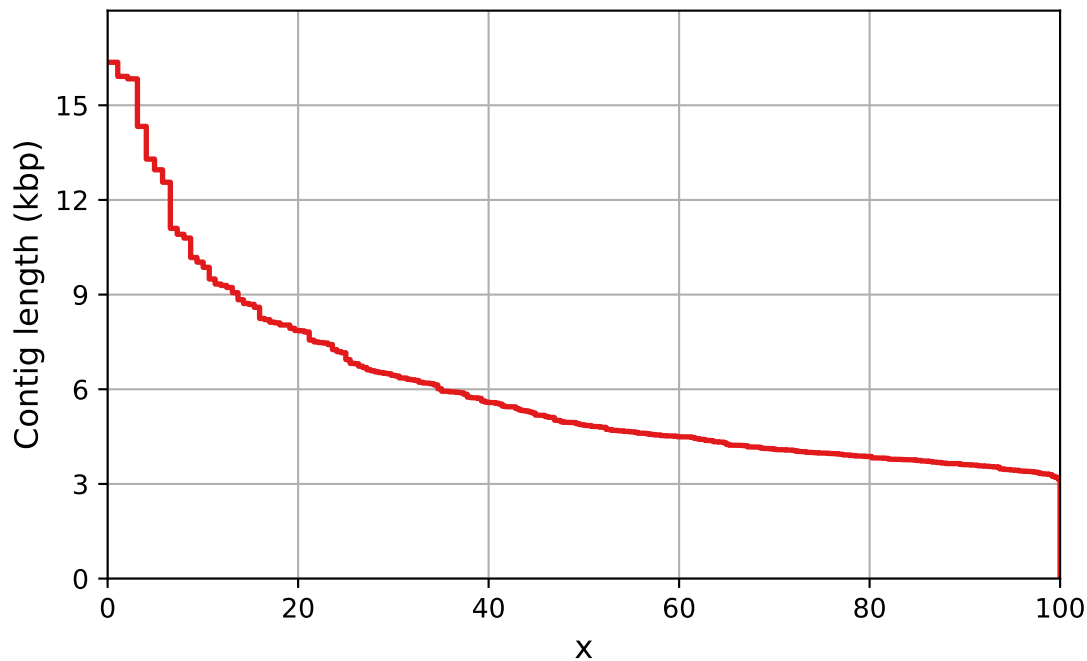


Report

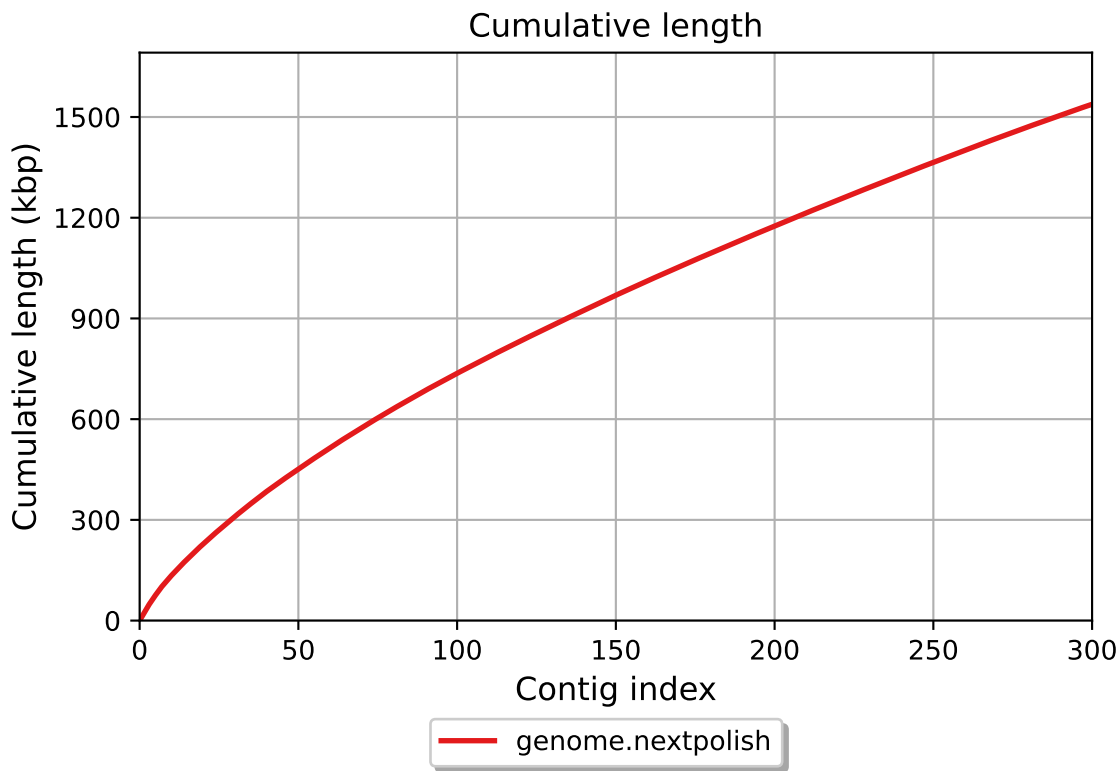
	genome.nextpolish
# contigs (>= 0 bp)	300
# contigs (>= 1000 bp)	300
# contigs (>= 5000 bp)	99
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1537935
Total length (>= 1000 bp)	1537935
Total length (>= 5000 bp)	731603
Total length (>= 10000 bp)	154260
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	300
Largest contig	16362
Total length	1537935
GC (%)	51.09
N50	4870
N90	3612
auN	6039.6
L50	107
L90	256
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

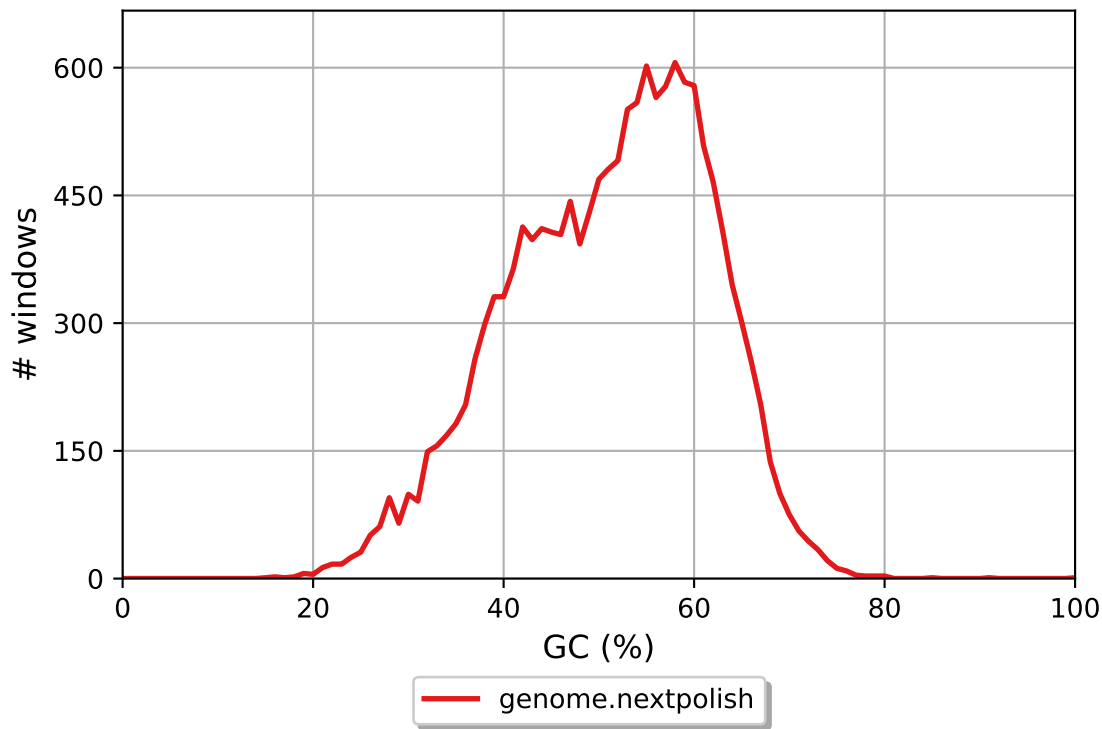
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

