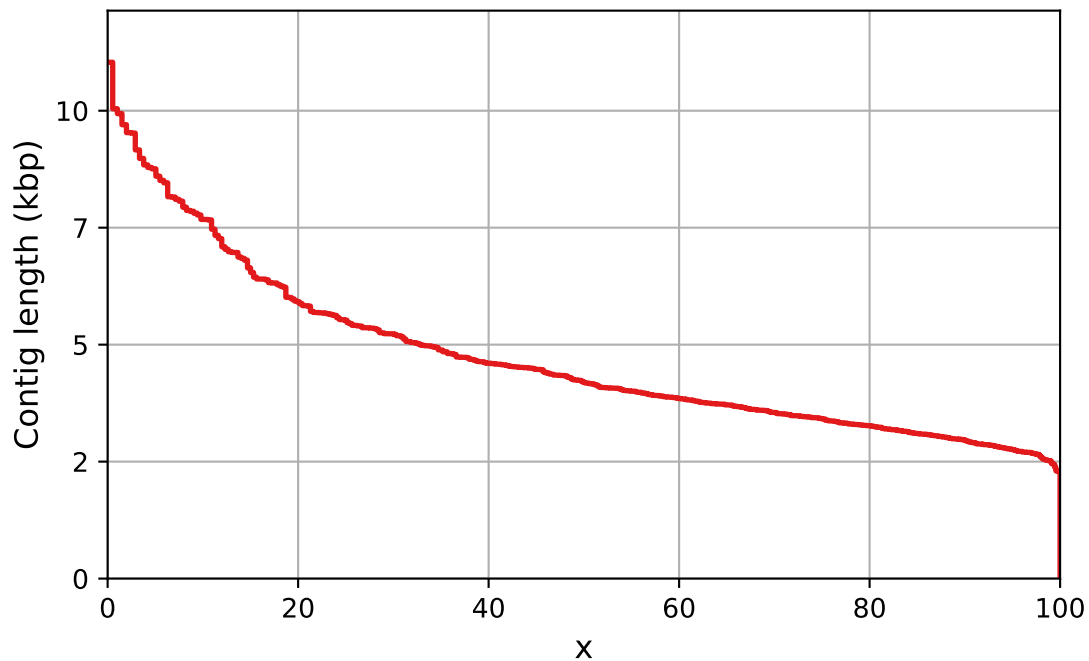


Report

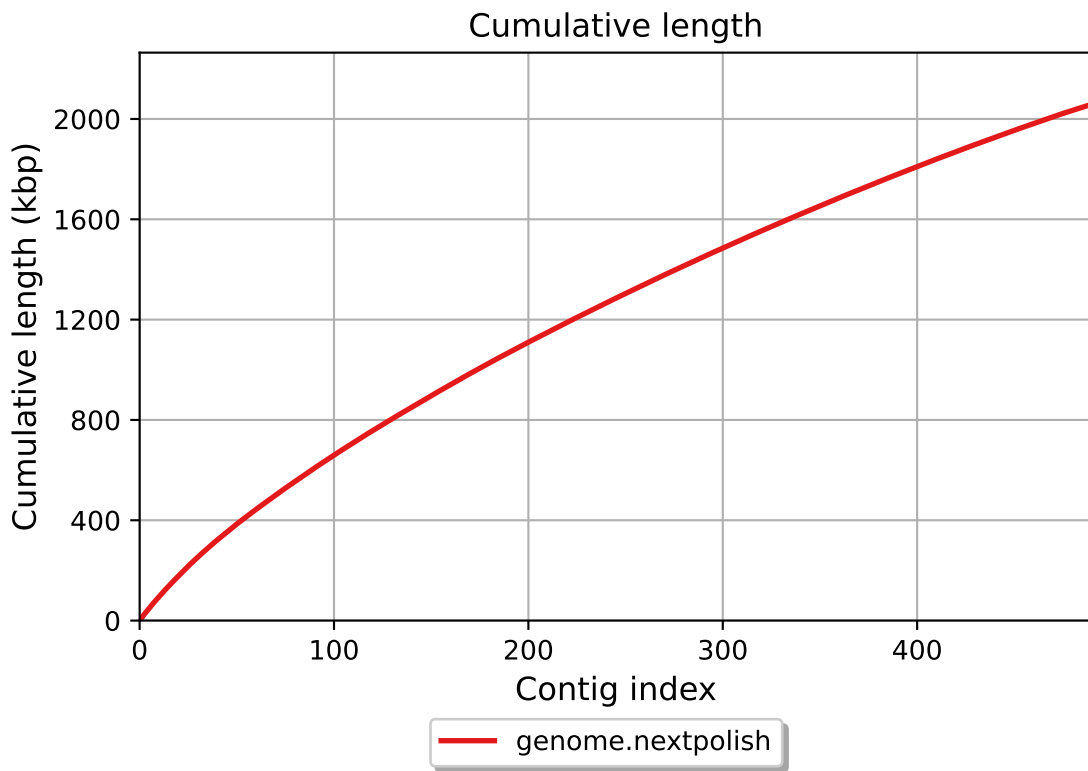
	genome.nextpolish
# contigs (>= 0 bp)	490
# contigs (>= 1000 bp)	490
# contigs (>= 5000 bp)	103
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2058599
Total length (>= 1000 bp)	2058599
Total length (>= 5000 bp)	674853
Total length (>= 10000 bp)	21077
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	490
Largest contig	11035
Total length	2058599
GC (%)	55.00
N50	4194
N90	2958
auN	4731.9
L50	181
L90	415
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

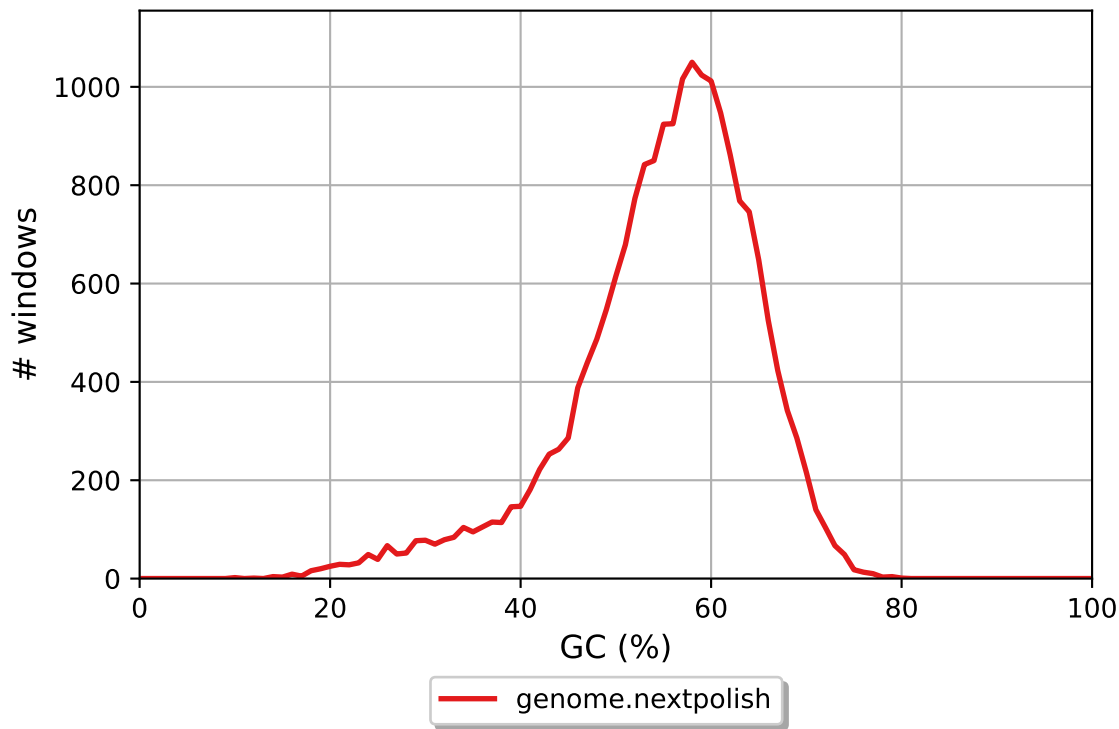
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

