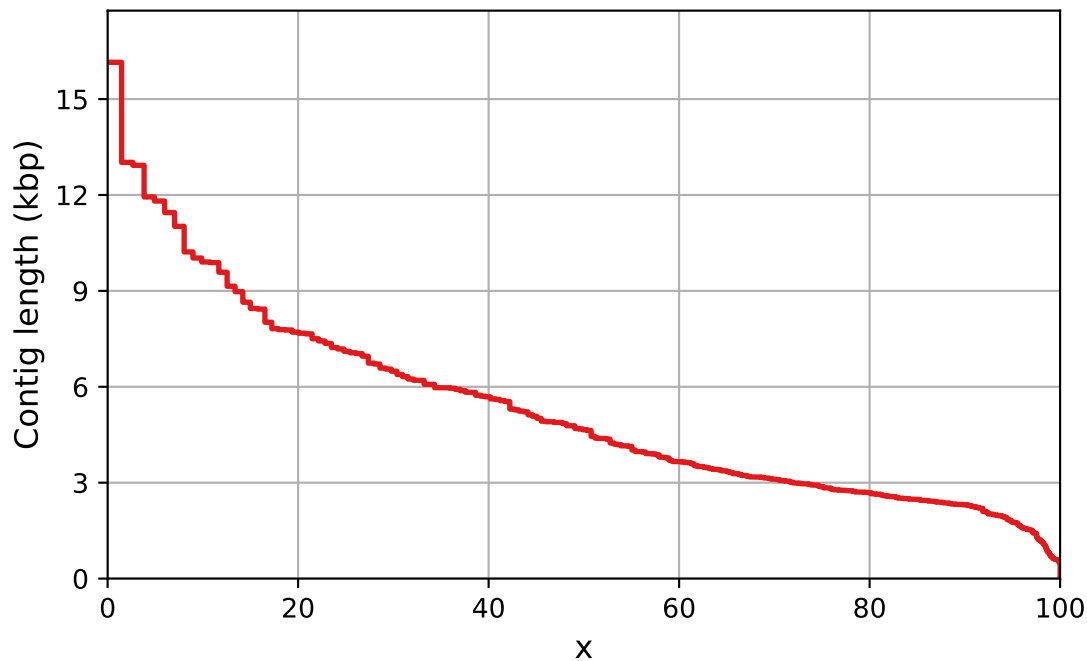


Report

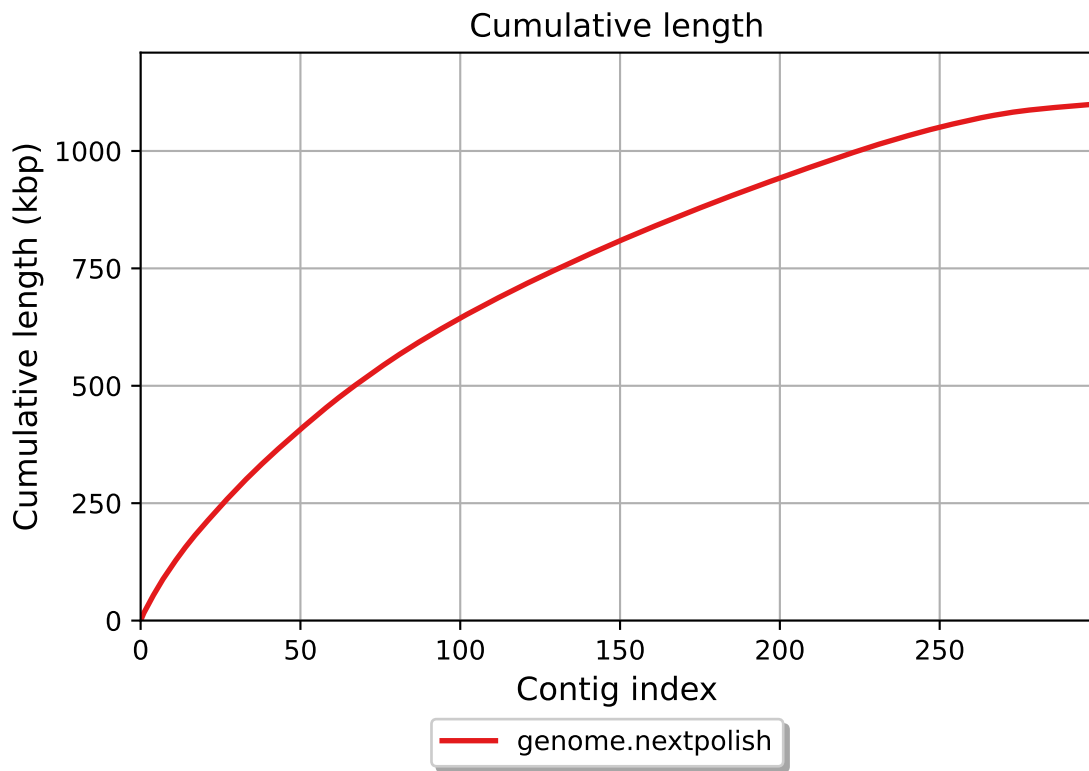
	genome.nextpolish
# contigs (>= 0 bp)	299
# contigs (>= 1000 bp)	273
# contigs (>= 5000 bp)	67
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1099795
Total length (>= 1000 bp)	1082720
Total length (>= 5000 bp)	500497
Total length (>= 10000 bp)	108566
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	298
Largest contig	16151
Total length	1099457
GC (%)	48.78
N50	4654
N90	2310
auN	5370.2
L50	78
L90	220
# 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).

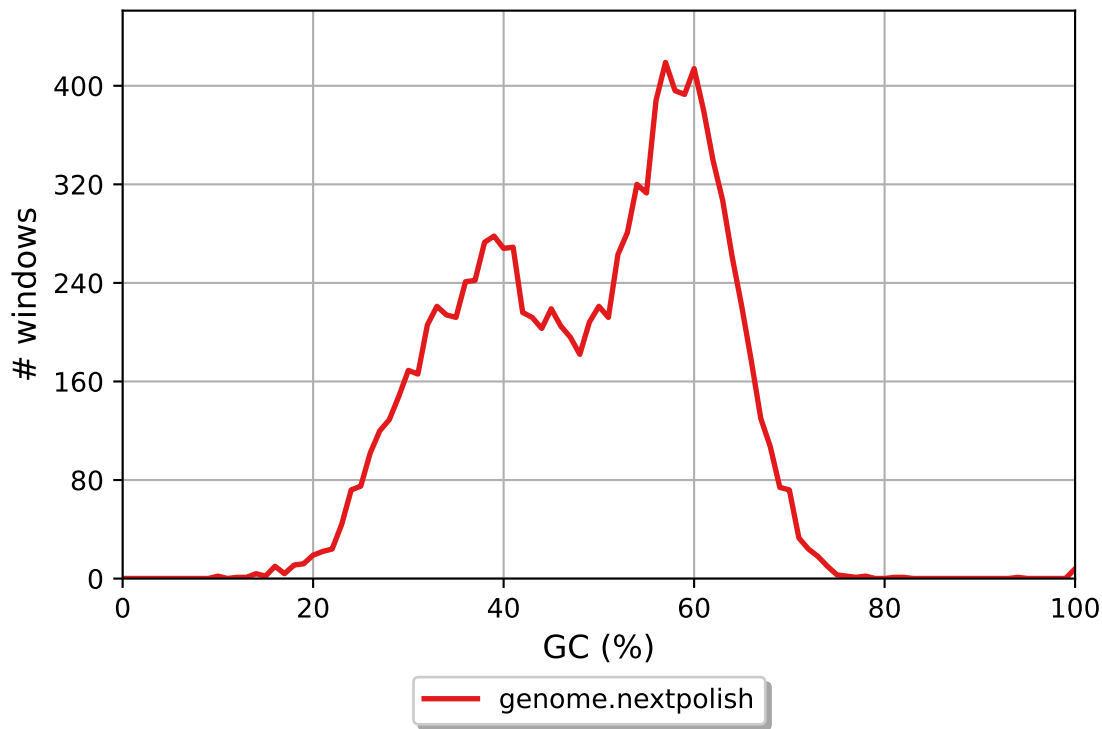
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

