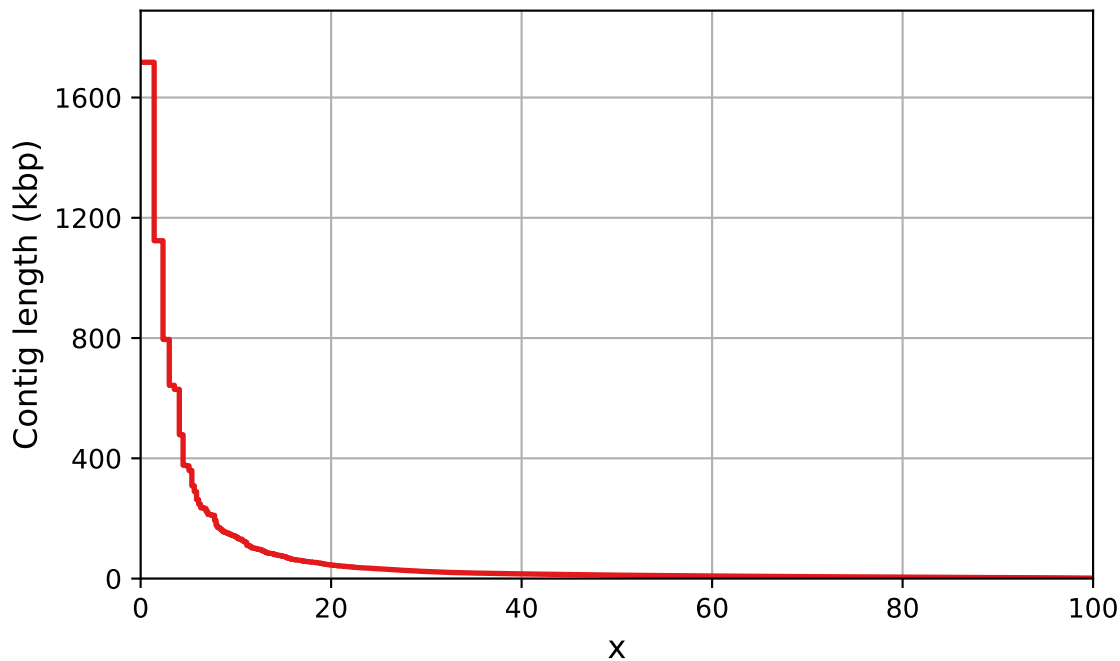


# Report

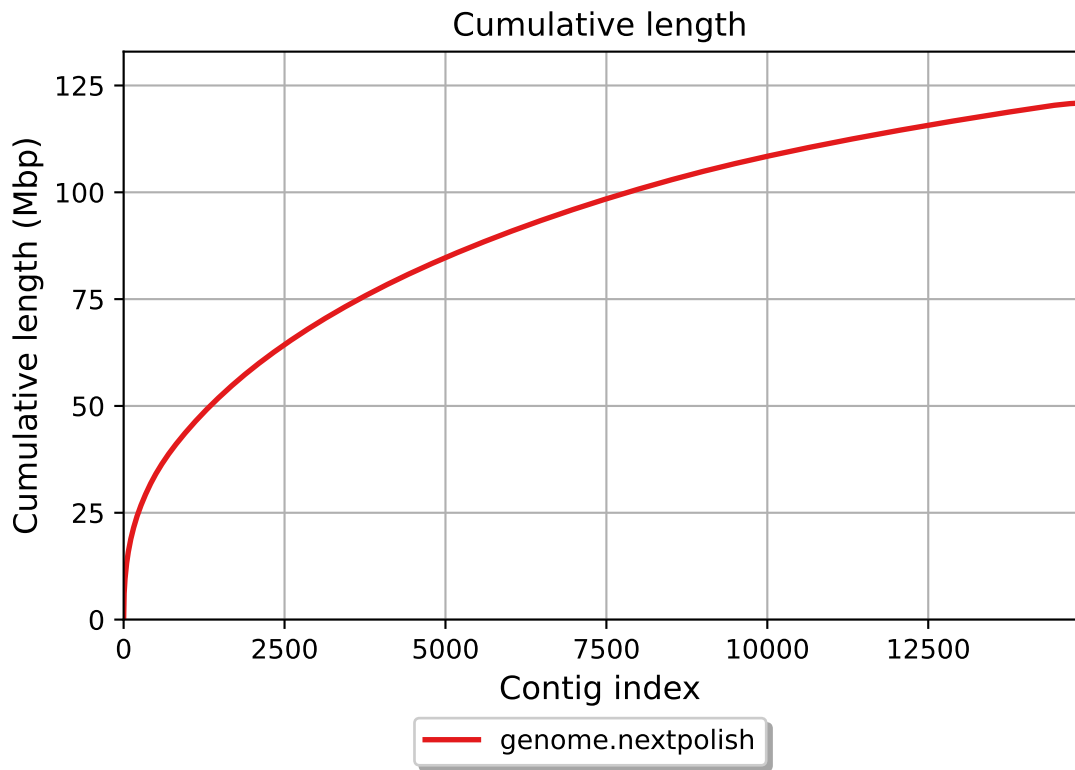
	genome.nextpolish
# contigs (>= 0 bp)	14798
# contigs (>= 1000 bp)	14744
# contigs (>= 5000 bp)	6971
# contigs (>= 10000 bp)	2694
# contigs (>= 25000 bp)	541
# contigs (>= 50000 bp)	184
Total length (>= 0 bp)	120838975
Total length (>= 1000 bp)	120802693
Total length (>= 5000 bp)	95913275
Total length (>= 10000 bp)	66341103
Total length (>= 25000 bp)	35136566
Total length (>= 50000 bp)	23025694
# contigs	14797
Largest contig	1717265
Total length	120838481
GC (%)	49.59
N50	11478
N90	3270
auN	78815.3
L50	2141
L90	10097
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	127 + 18 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).

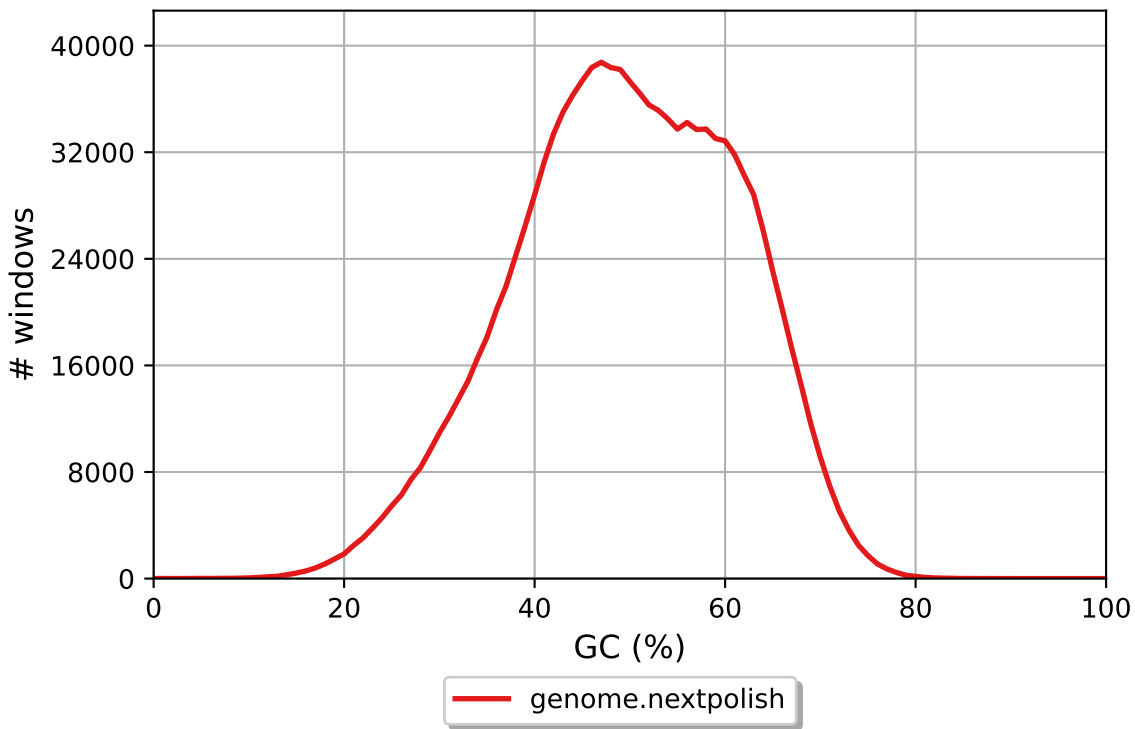
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

