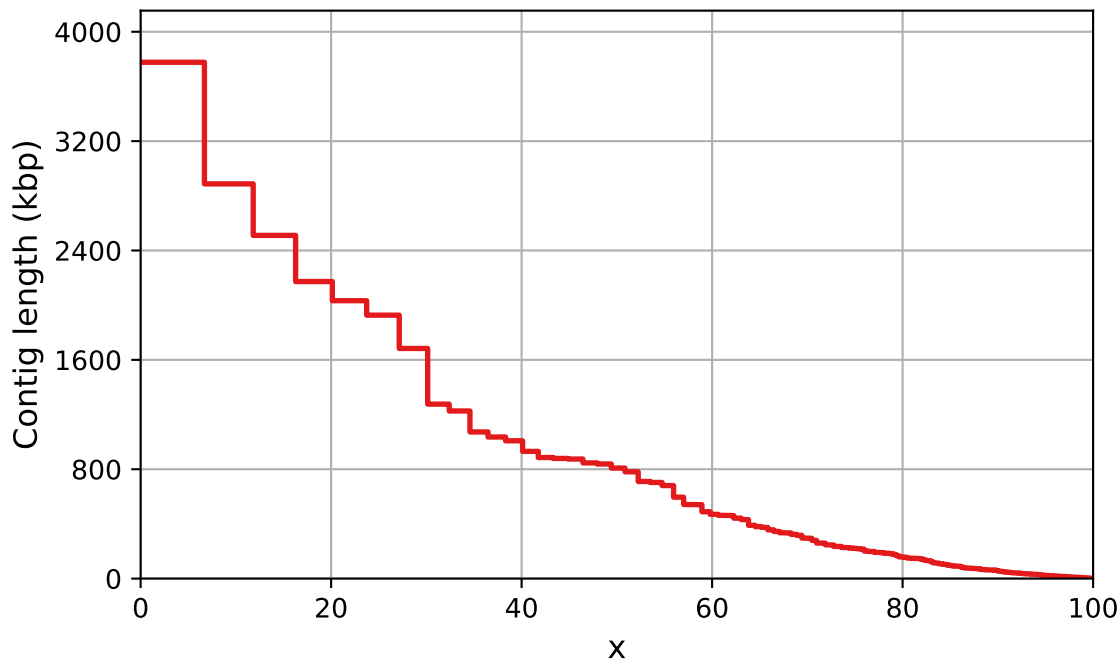


Report

	genome.nextpolish
# contigs (>= 0 bp)	542
# contigs (>= 1000 bp)	515
# contigs (>= 5000 bp)	414
# contigs (>= 10000 bp)	324
# contigs (>= 25000 bp)	199
# contigs (>= 50000 bp)	135
Total length (>= 0 bp)	56378423
Total length (>= 1000 bp)	56361069
Total length (>= 5000 bp)	56074421
Total length (>= 10000 bp)	55408857
Total length (>= 25000 bp)	53392030
Total length (>= 50000 bp)	51066672
# contigs	541
Largest contig	3776868
Total length	56377931
GC (%)	42.93
N50	808504
N90	57529
auN	1124392.9
L50	19
L90	129
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	137 + 0 part

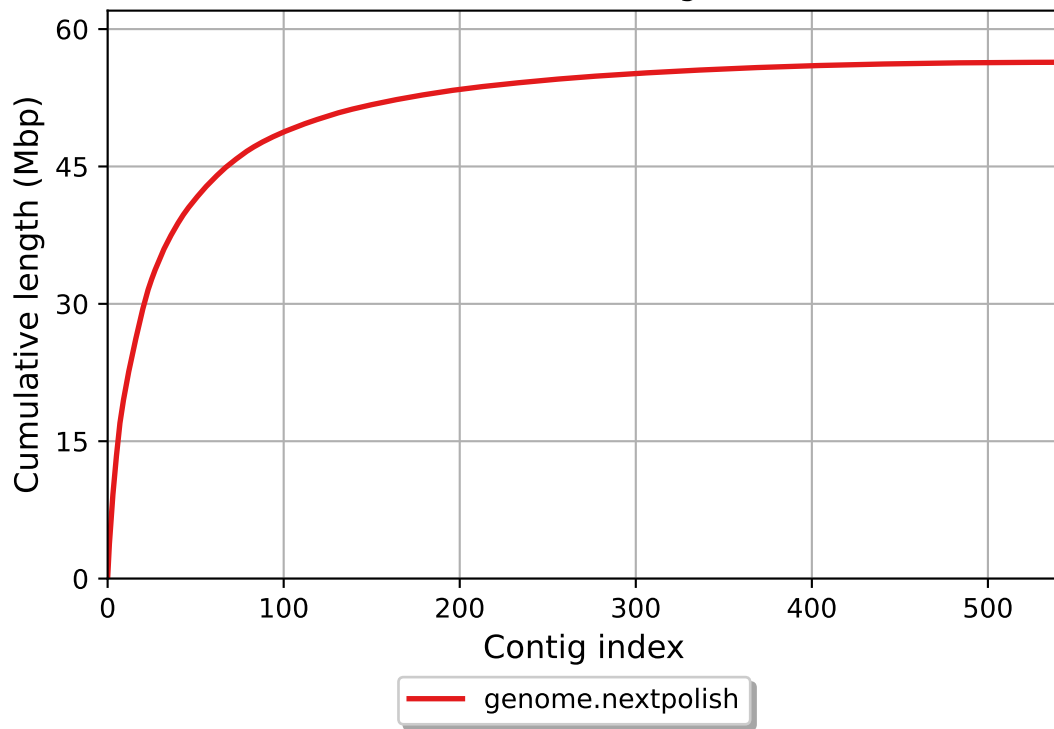
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

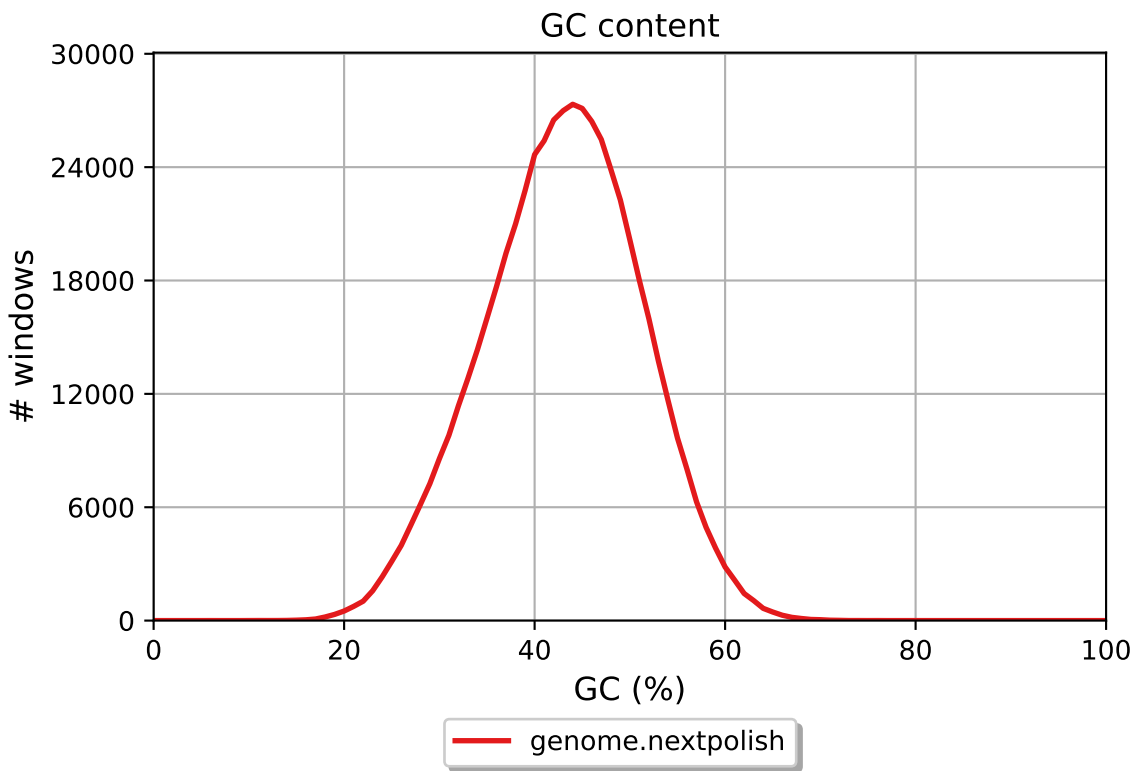
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

