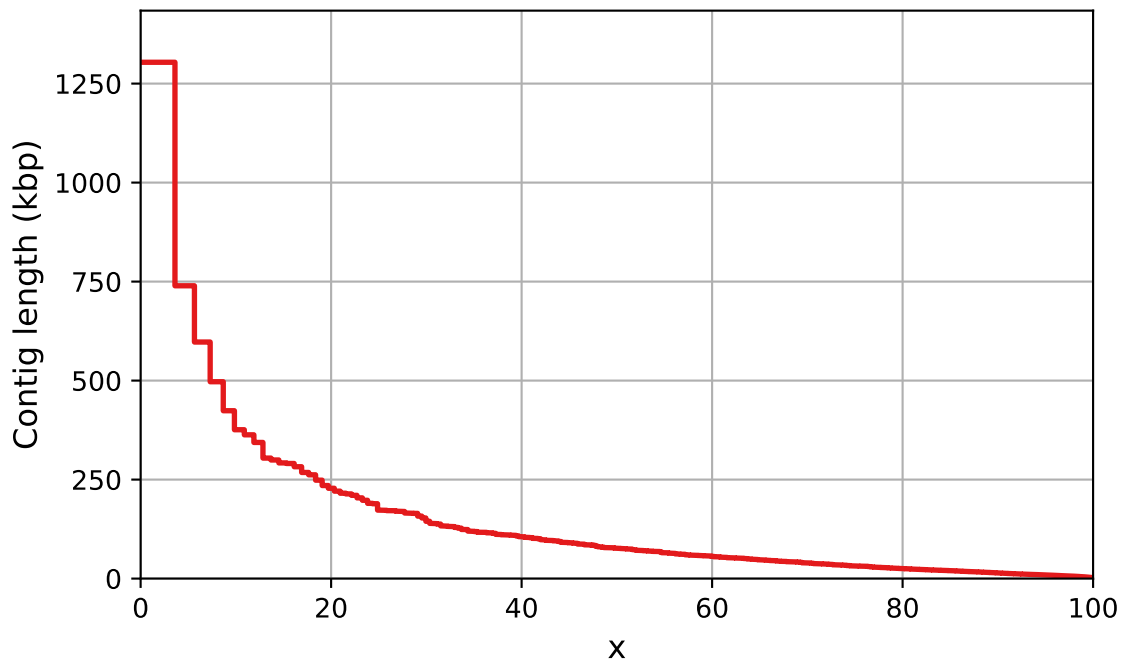


Report

	genome.nextpolish
# contigs (>= 0 bp)	1025
# contigs (>= 1000 bp)	1018
# contigs (>= 5000 bp)	890
# contigs (>= 10000 bp)	671
# contigs (>= 25000 bp)	356
# contigs (>= 50000 bp)	188
Total length (>= 0 bp)	36165243
Total length (>= 1000 bp)	36159826
Total length (>= 5000 bp)	35696580
Total length (>= 10000 bp)	34089158
Total length (>= 25000 bp)	28994445
Total length (>= 50000 bp)	23044760
# contigs	1025
Largest contig	1303851
Total length	36165243
GC (%)	57.01
N50	76781
N90	14237
auN	171724.2
L50	108
L90	541
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	62 + 3 part

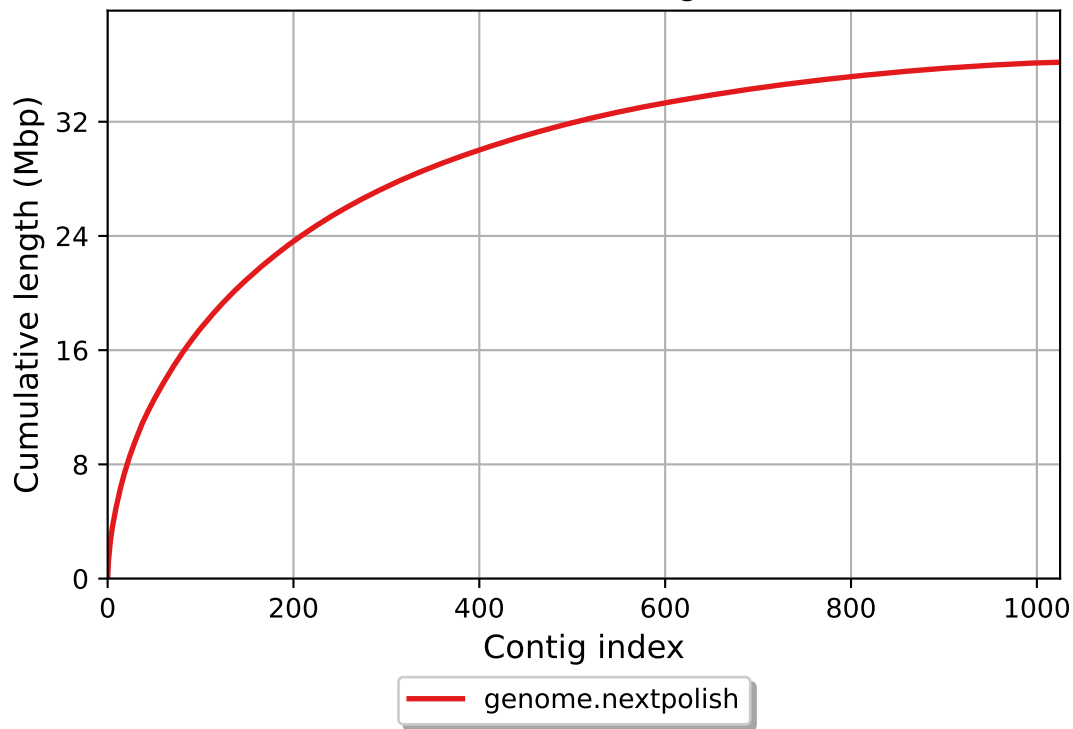
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

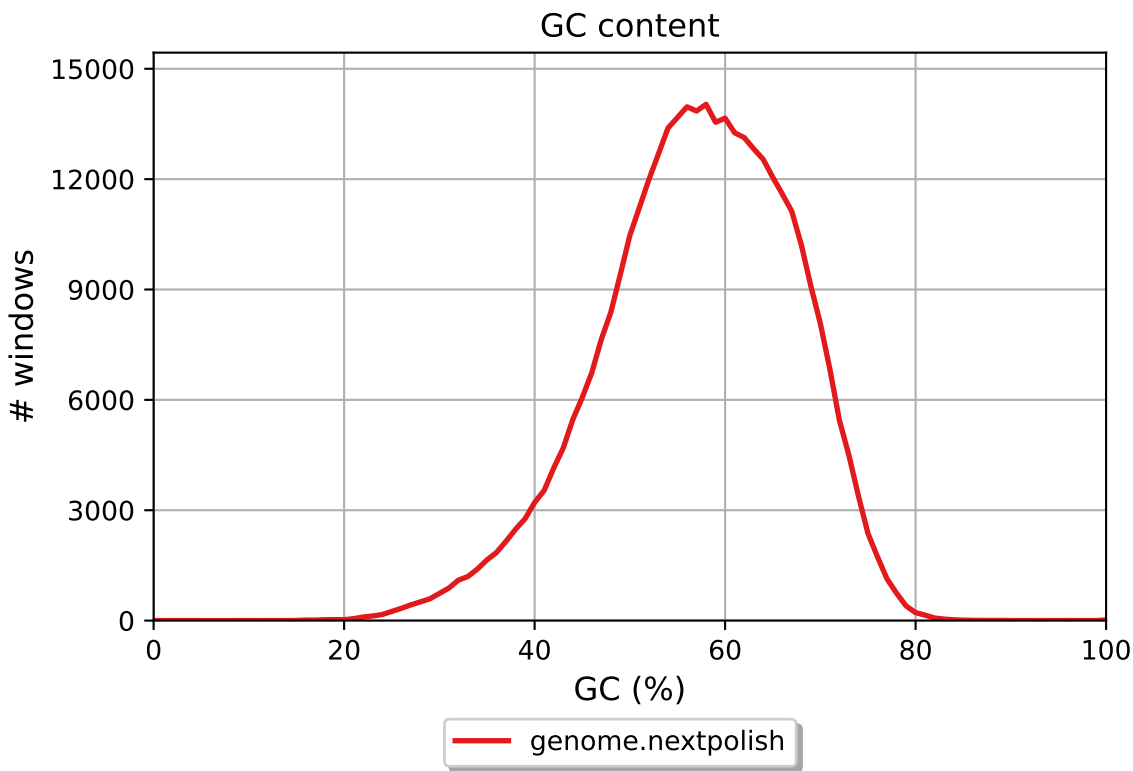
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

