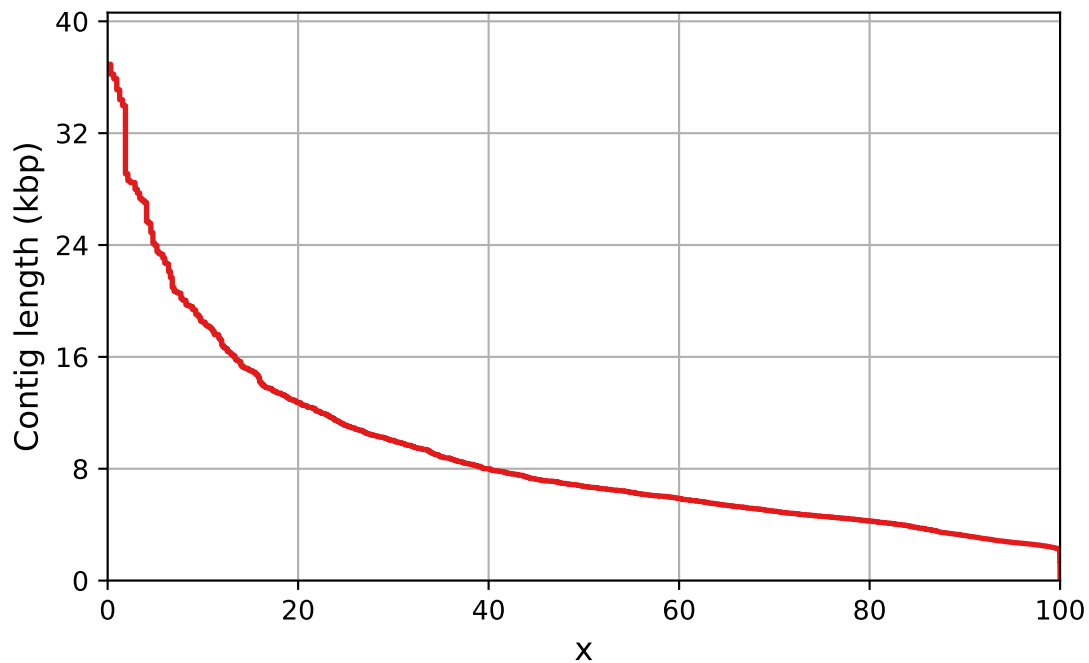


Report

	genome.nextpolish
# contigs (>= 0 bp)	1868
# contigs (>= 1000 bp)	1867
# contigs (>= 5000 bp)	886
# contigs (>= 10000 bp)	225
# contigs (>= 25000 bp)	17
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	11364203
Total length (>= 1000 bp)	11363648
Total length (>= 5000 bp)	7892353
Total length (>= 10000 bp)	3420206
Total length (>= 25000 bp)	515600
Total length (>= 50000 bp)	0
# contigs	1868
Largest contig	36928
Total length	11364203
GC (%)	41.63
N50	6739
N90	3227
auN	9110.3
L50	508
L90	1450
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 3 part

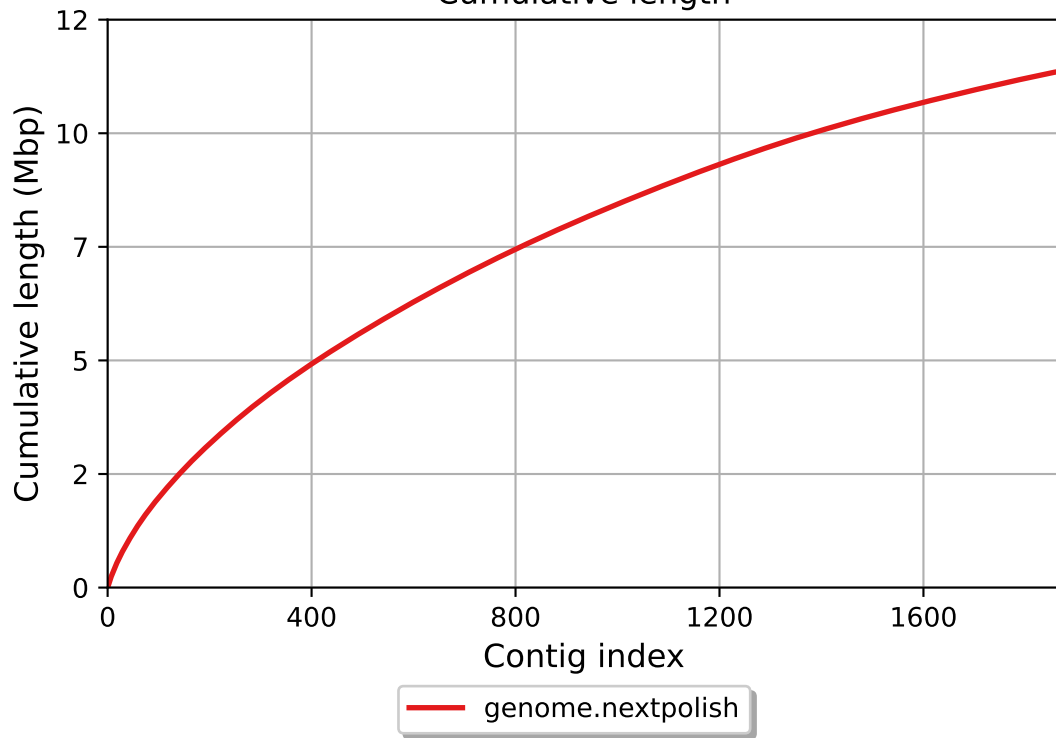
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

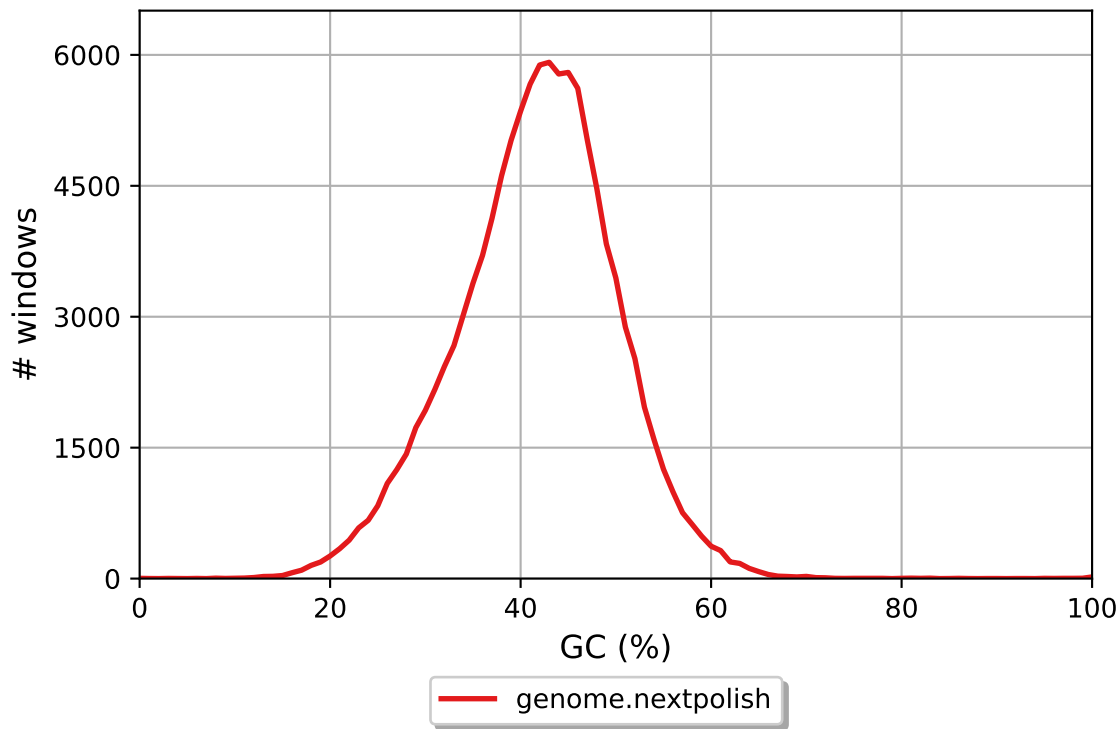


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

