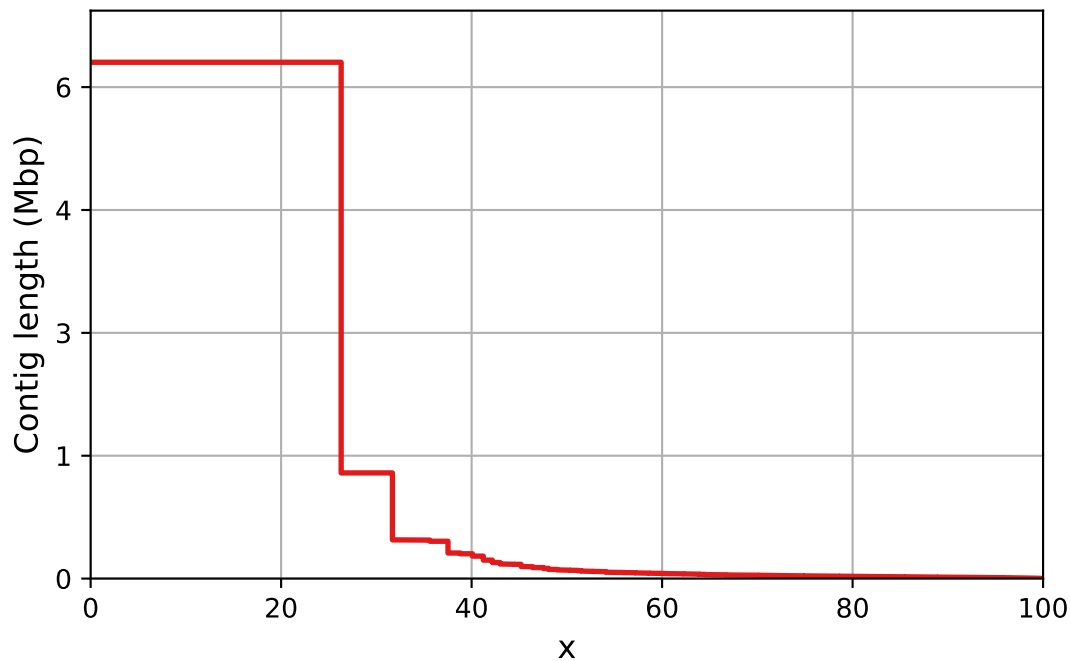


Report

	genome.nextpolish
# contigs (>= 0 bp)	548
# contigs (>= 1000 bp)	547
# contigs (>= 5000 bp)	493
# contigs (>= 10000 bp)	401
# contigs (>= 25000 bp)	195
# contigs (>= 50000 bp)	72
Total length (>= 0 bp)	23965028
Total length (>= 1000 bp)	23964187
Total length (>= 5000 bp)	23789643
Total length (>= 10000 bp)	23109683
Total length (>= 25000 bp)	19716461
Total length (>= 50000 bp)	15465605
# contigs	548
Largest contig	6303516
Total length	23965028
GC (%)	56.35
N50	102739
N90	17063
auN	1799014.5
L50	23
L90	285
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	52 + 3 part

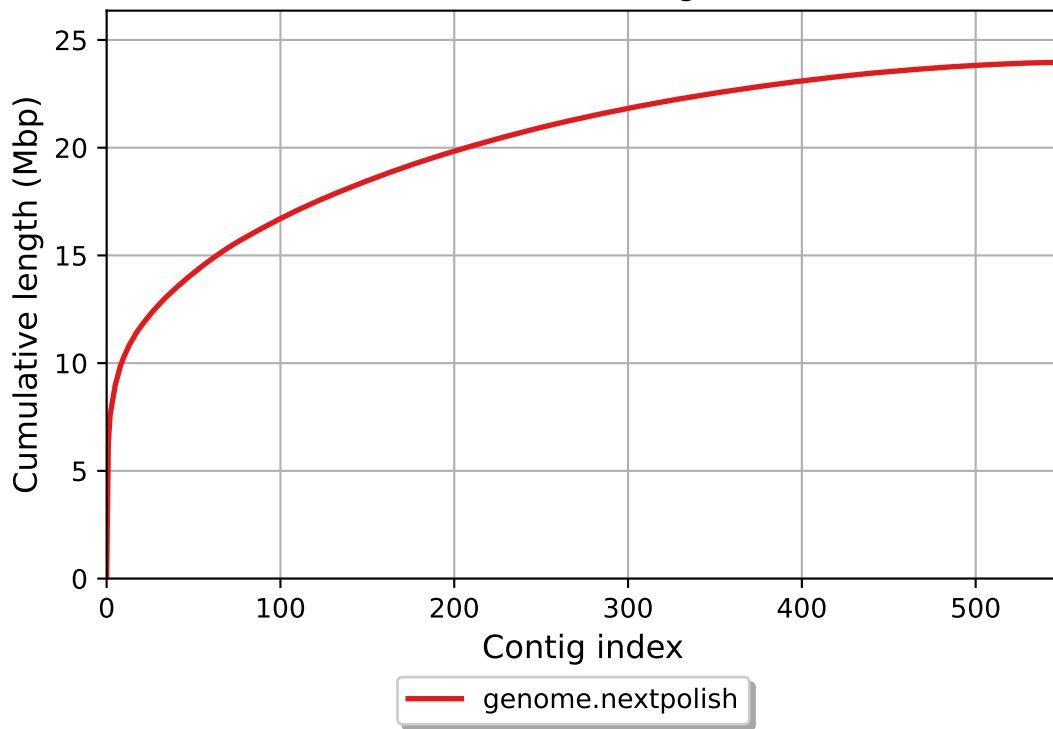
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

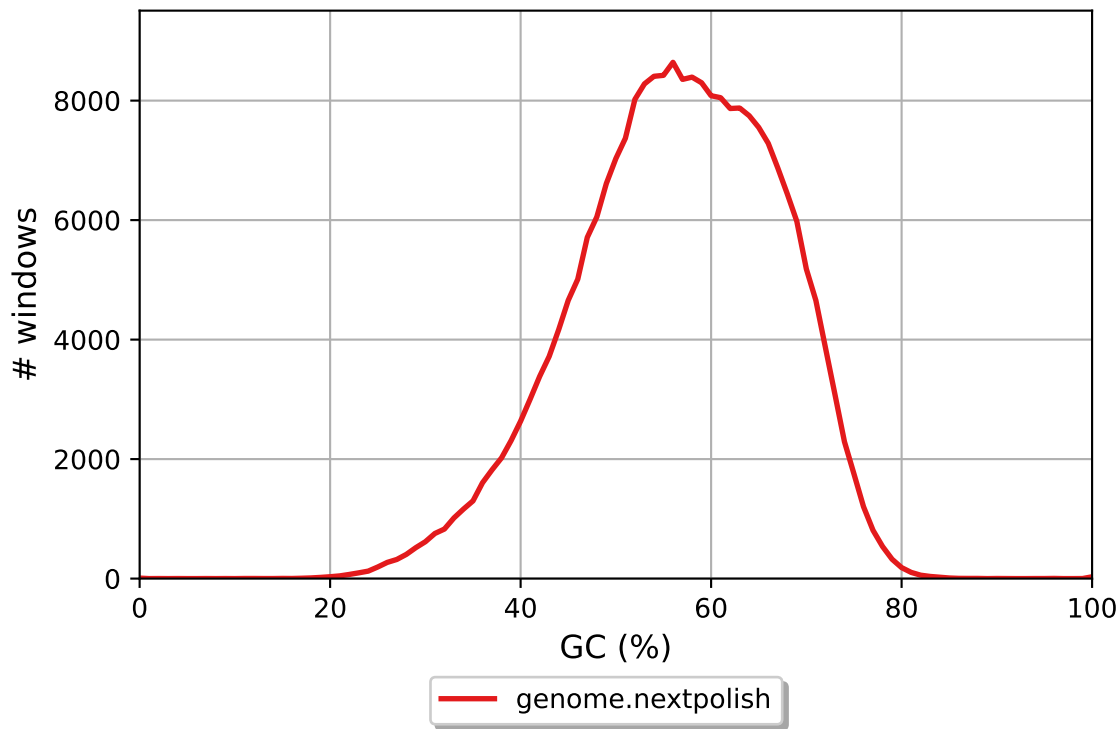


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

