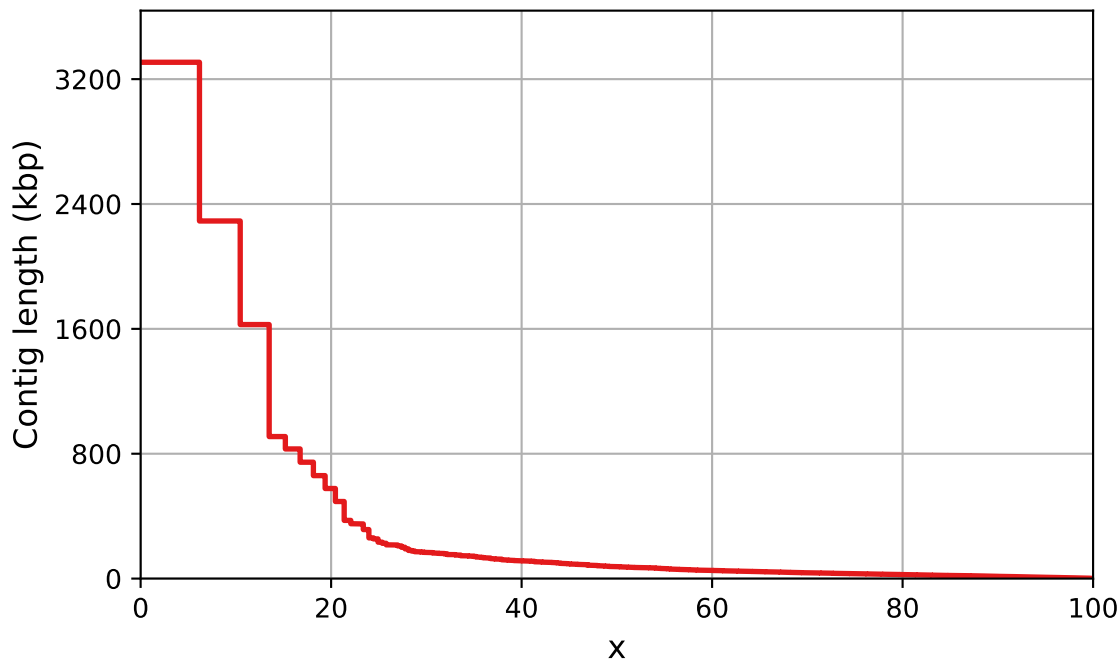


Report

	genome.nextpolish
# contigs (≥ 0 bp)	1628
# contigs (≥ 1000 bp)	1616
# contigs (≥ 5000 bp)	1245
# contigs (≥ 10000 bp)	941
# contigs (≥ 25000 bp)	516
# contigs (≥ 50000 bp)	222
Total length (≥ 0 bp)	53547096
Total length (≥ 1000 bp)	53539166
Total length (≥ 5000 bp)	52362182
Total length (≥ 10000 bp)	50194553
Total length (≥ 25000 bp)	43143926
Total length (≥ 50000 bp)	32743411
# contigs	1628
Largest contig	3308066
Total length	53547096
GC (%)	51.77
N50	76524
N90	14582
auN	471469.9
L50	125
L90	776
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	113 + 11 part

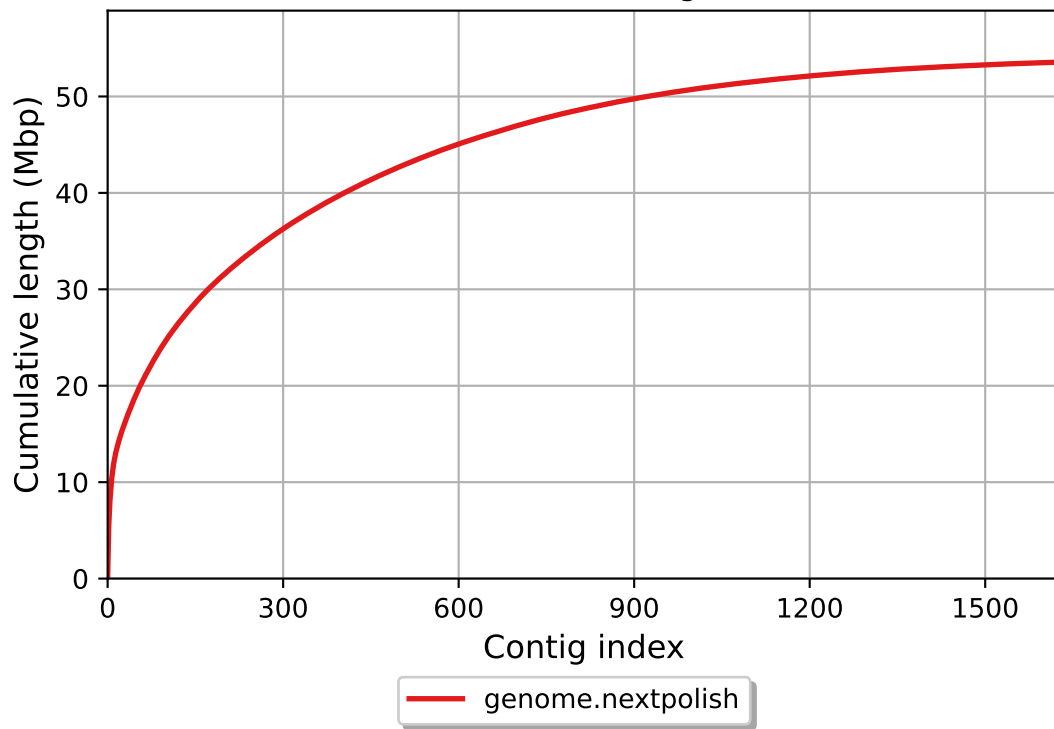
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

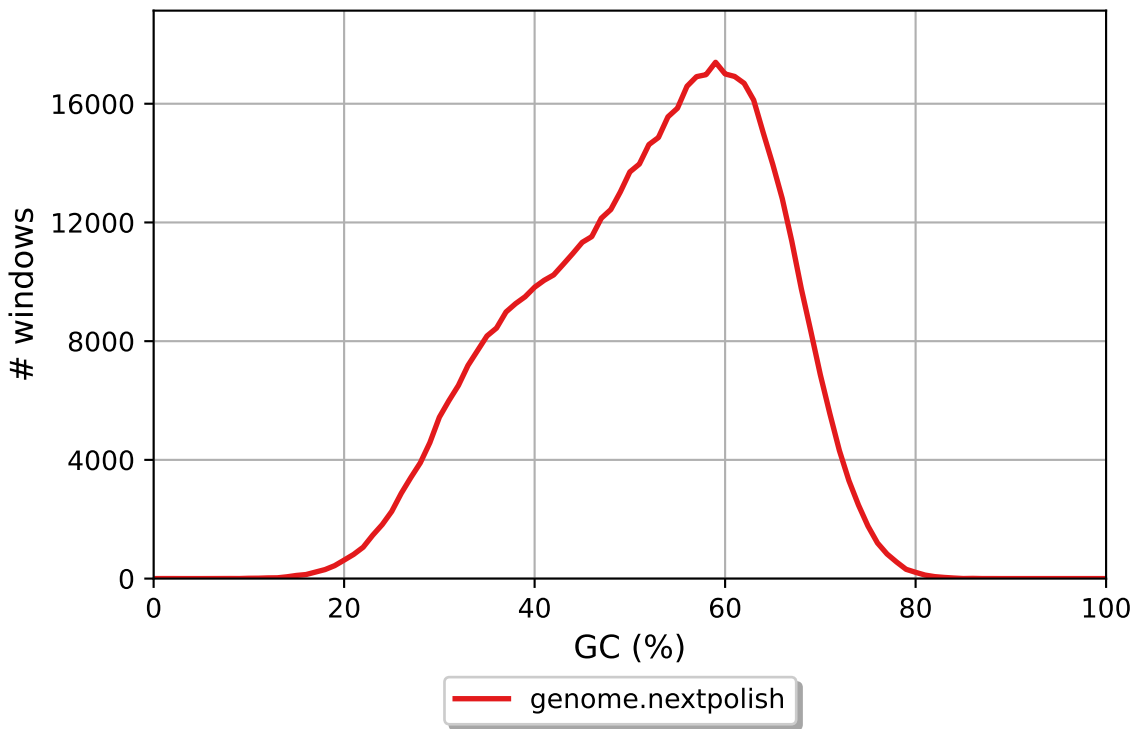


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

