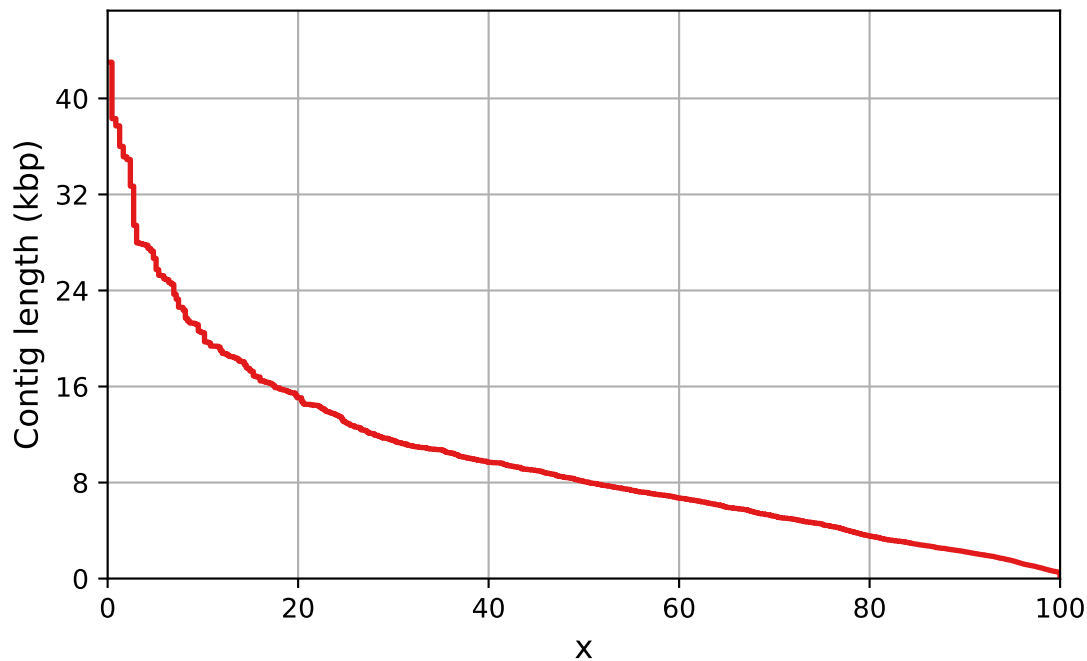


Report

	genome.nextpolish
# contigs (>= 0 bp)	1981
# contigs (>= 1000 bp)	1637
# contigs (>= 5000 bp)	671
# contigs (>= 10000 bp)	233
# contigs (>= 25000 bp)	18
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	9435567
Total length (>= 1000 bp)	9193315
Total length (>= 5000 bp)	6744115
Total length (>= 10000 bp)	3594133
Total length (>= 25000 bp)	556373
Total length (>= 50000 bp)	0
# contigs	1975
Largest contig	43010
Total length	9433581
GC (%)	51.10
N50	8108
N90	2237
auN	10027.2
L50	357
L90	1193
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 1 part

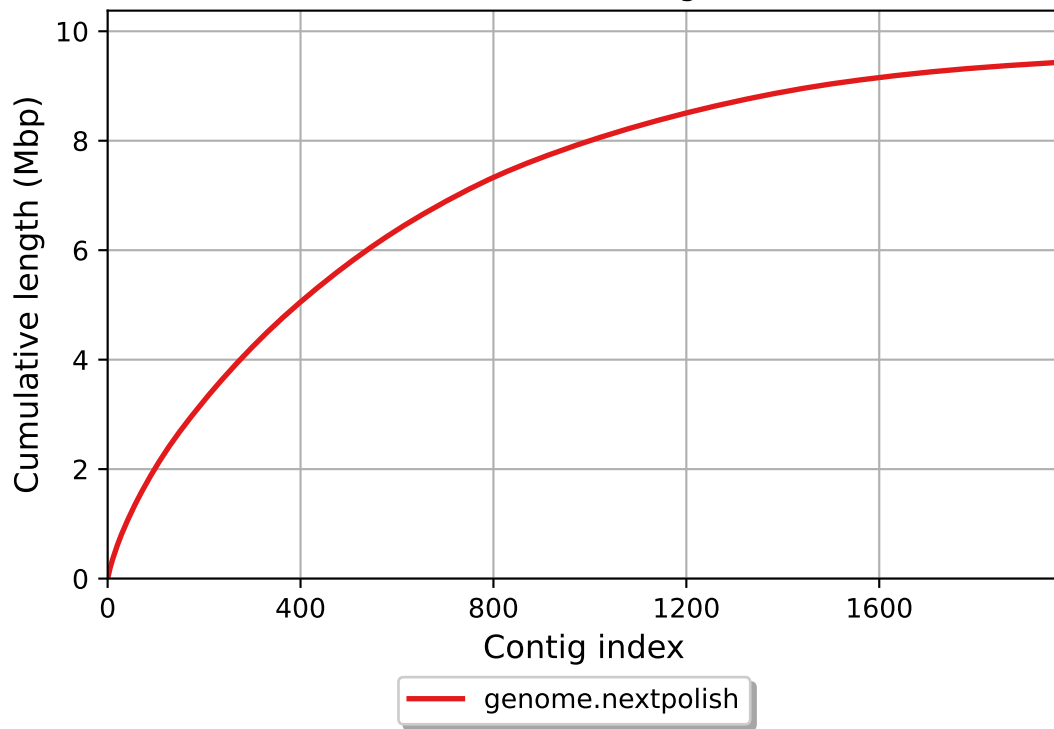
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

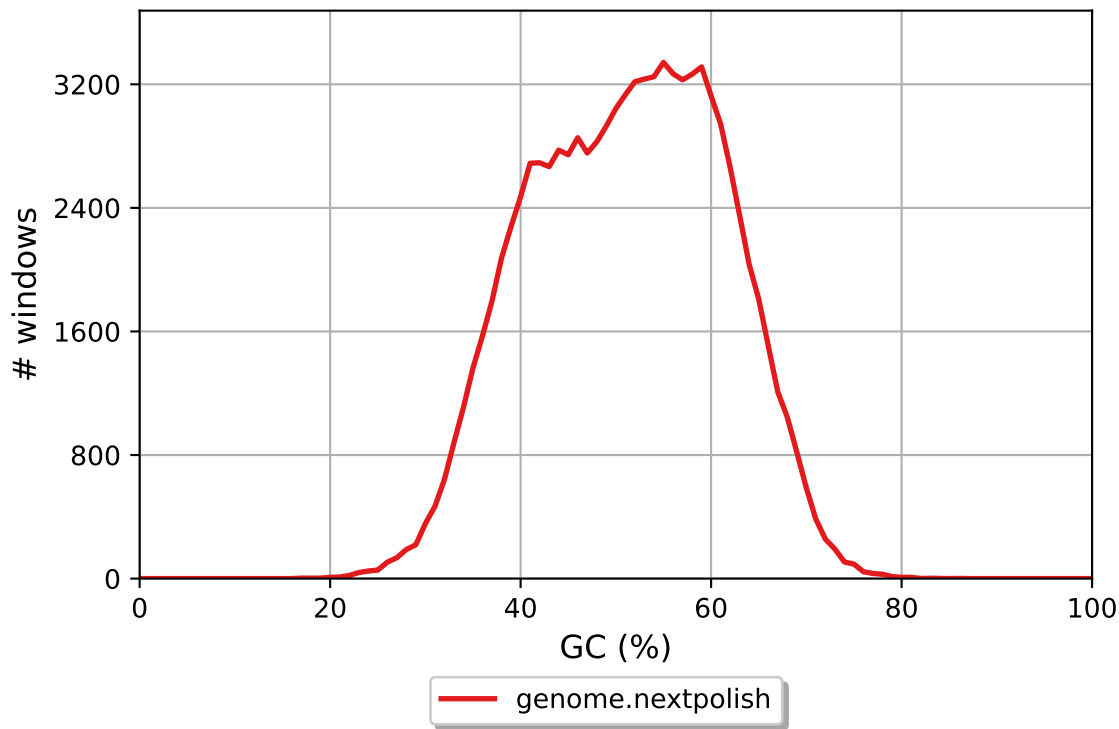


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

