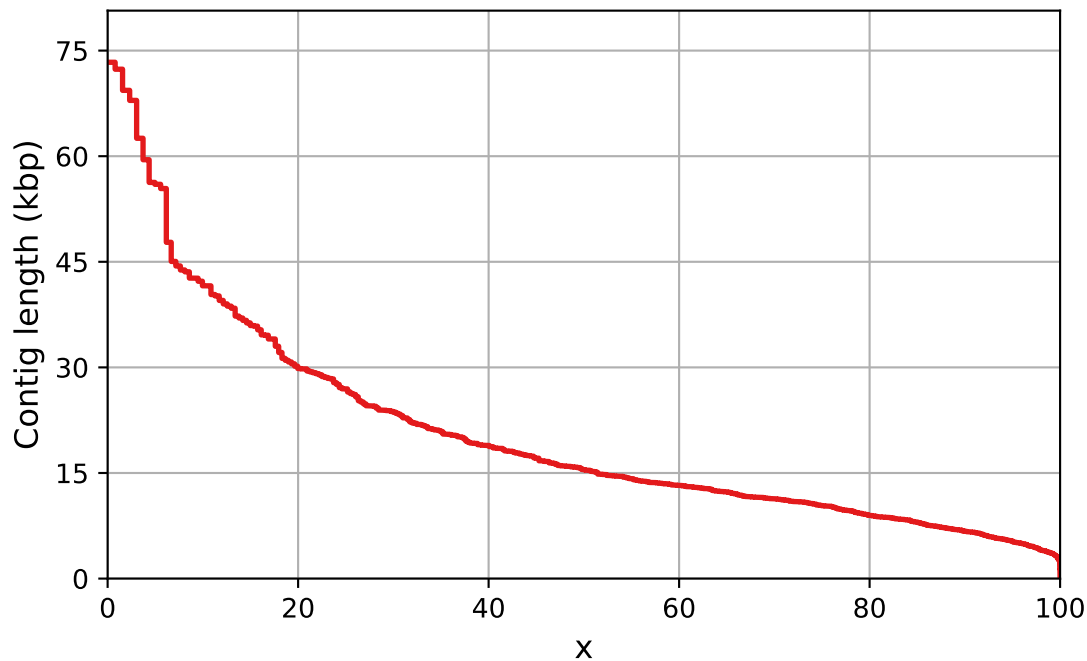


Report

	genome.nextpolish
# contigs (>= 0 bp)	719
# contigs (>= 1000 bp)	718
# contigs (>= 5000 bp)	621
# contigs (>= 10000 bp)	376
# contigs (>= 25000 bp)	66
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	9292946
Total length (>= 1000 bp)	9292337
Total length (>= 5000 bp)	8912594
Total length (>= 10000 bp)	7118744
Total length (>= 25000 bp)	2495929
Total length (>= 50000 bp)	572704
# contigs	719
Largest contig	73341
Total length	9292946
GC (%)	52.79
N50	15466
N90	6723
auN	20728.5
L50	178
L90	528
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 0 part

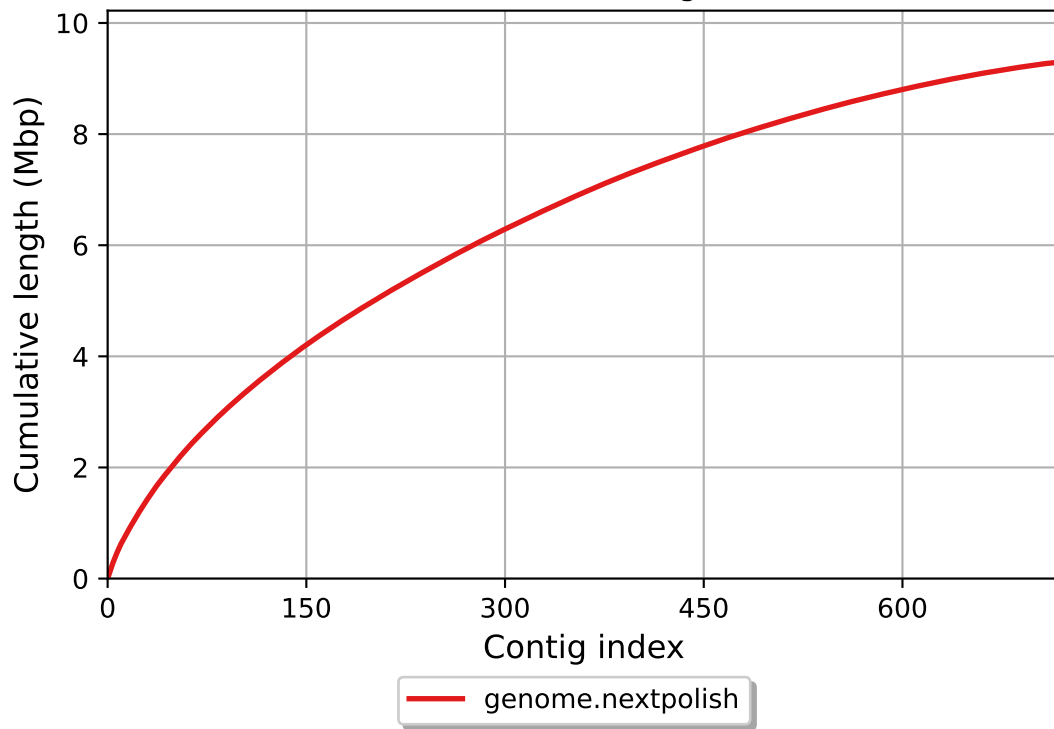
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

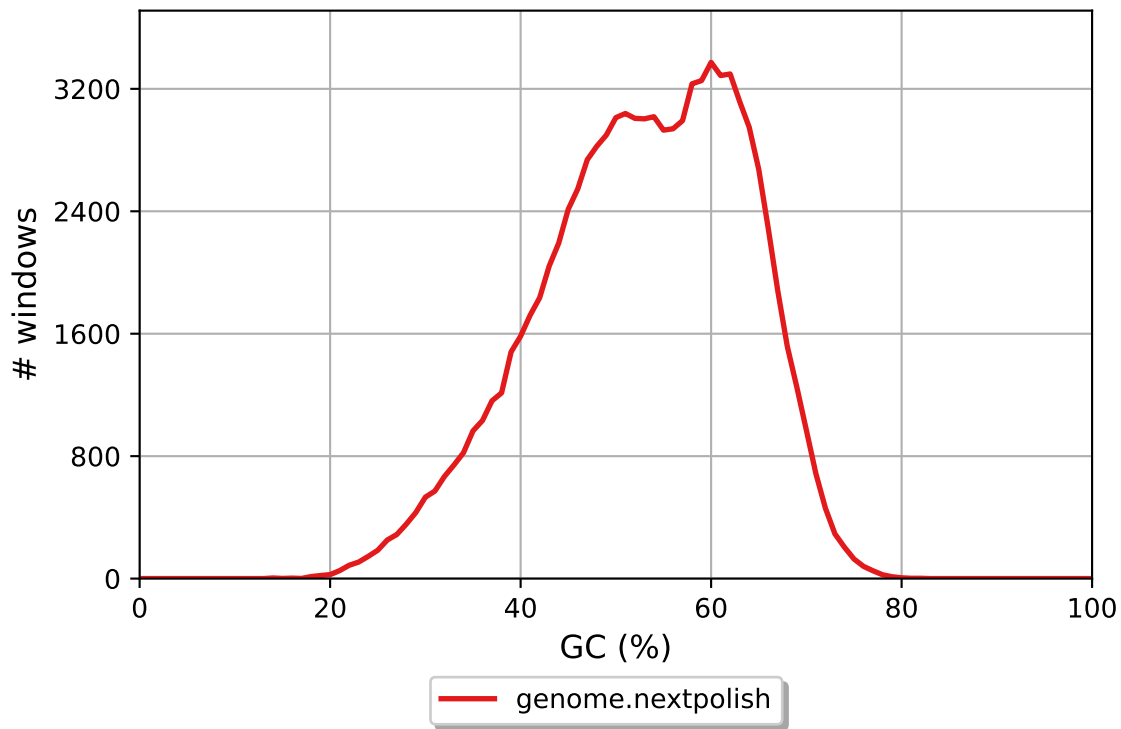


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

