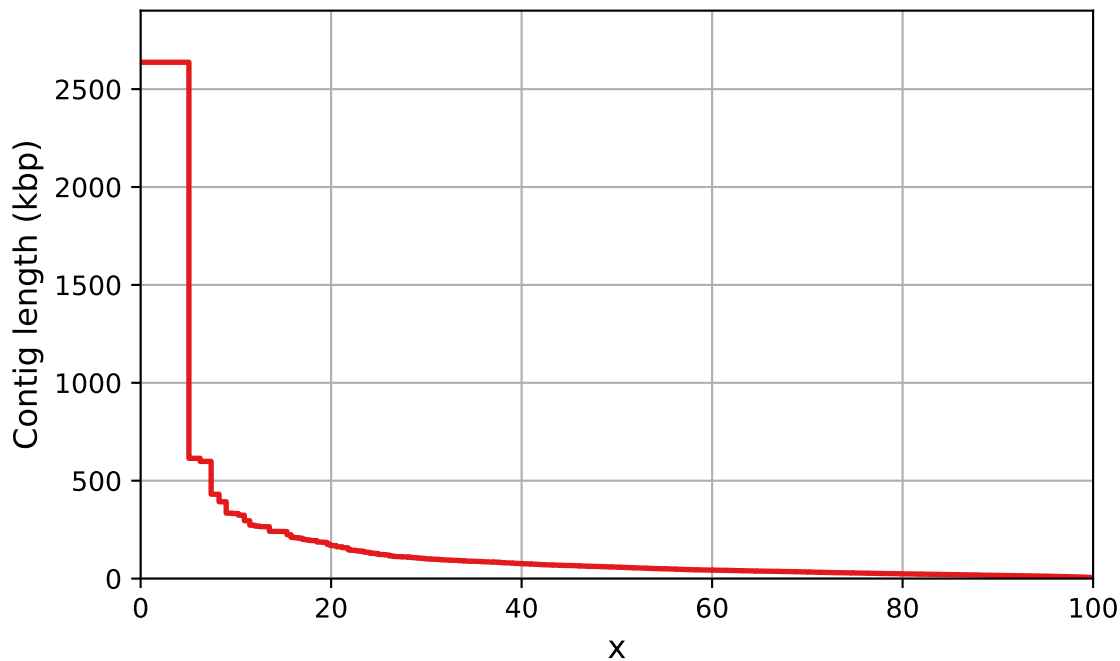


Report

	genome.nextpolish
# contigs (>= 0 bp)	1369
# contigs (>= 1000 bp)	1366
# contigs (>= 5000 bp)	1348
# contigs (>= 10000 bp)	1172
# contigs (>= 25000 bp)	609
# contigs (>= 50000 bp)	257
Total length (>= 0 bp)	52004188
Total length (>= 1000 bp)	52002378
Total length (>= 5000 bp)	51932407
Total length (>= 10000 bp)	50523219
Total length (>= 25000 bp)	41064866
Total length (>= 50000 bp)	28649717
# contigs	1369
Largest contig	2637180
Total length	52004188
GC (%)	53.55
N50	58611
N90	16341
auN	224904.9
L50	208
L90	891
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	146 + 8 part

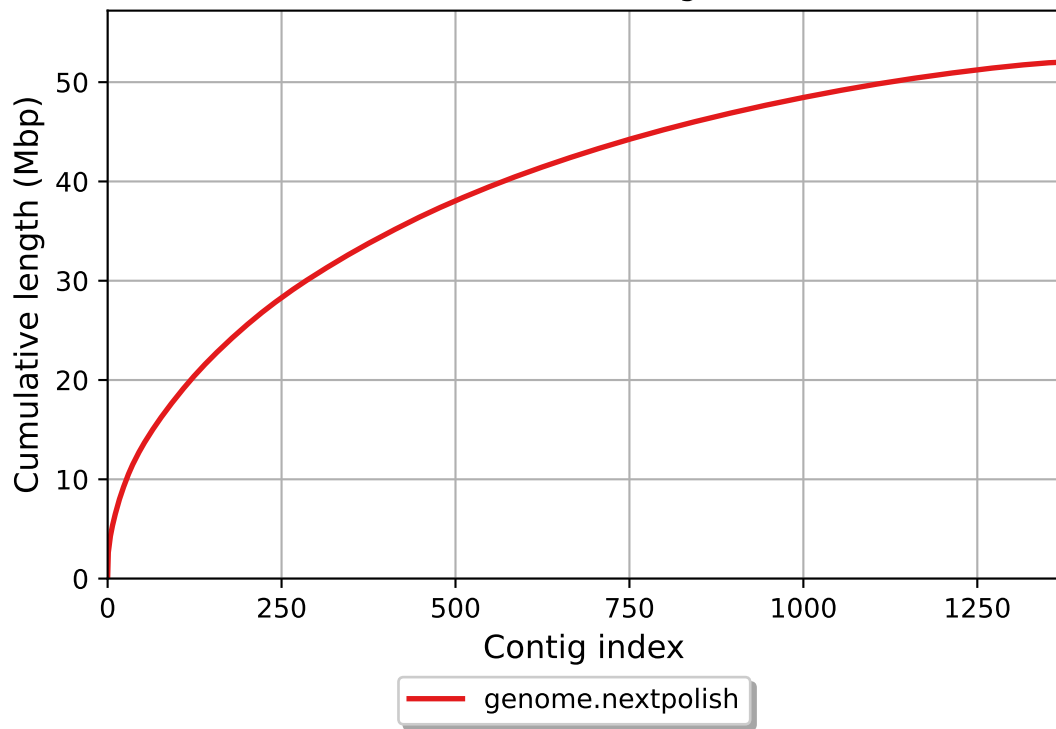
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

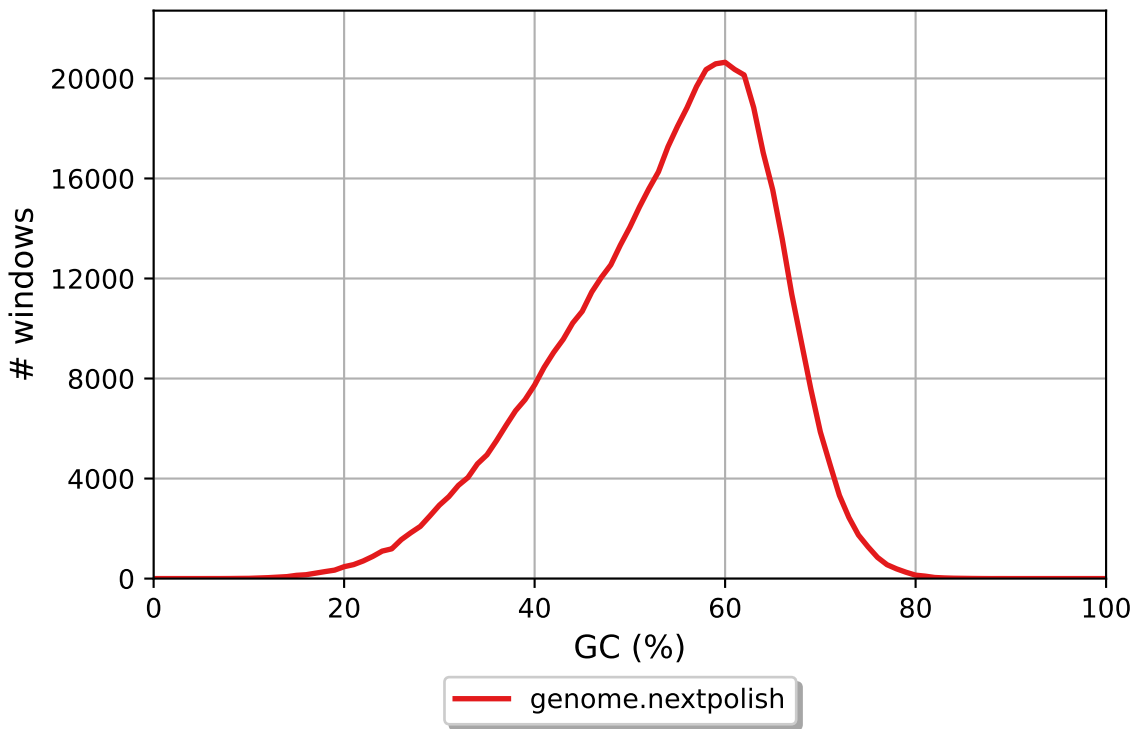


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

