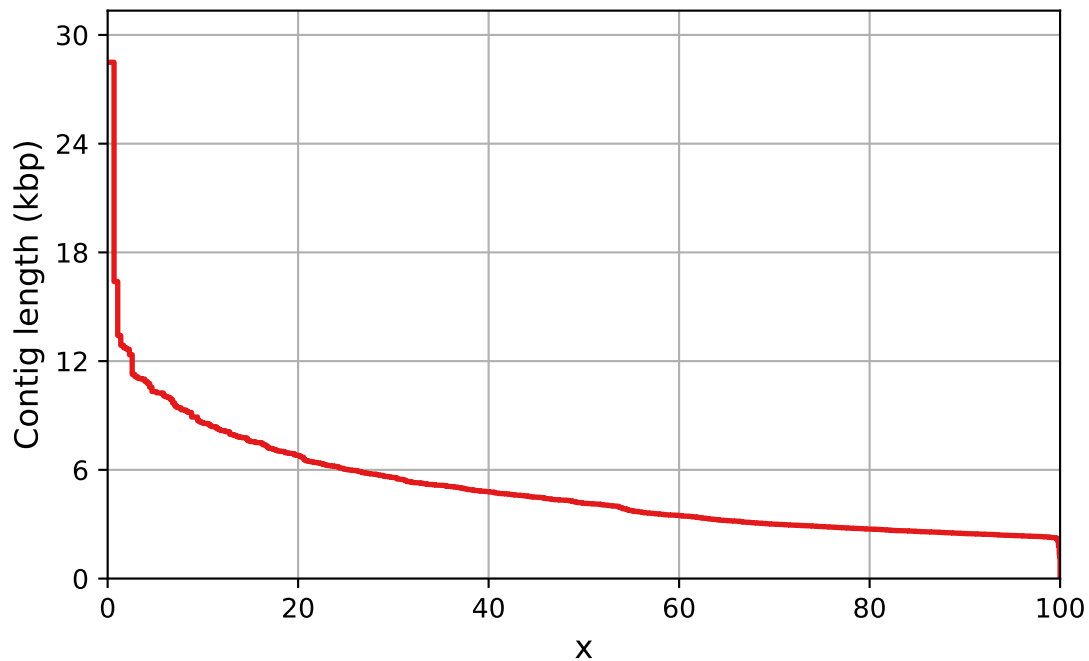


Report

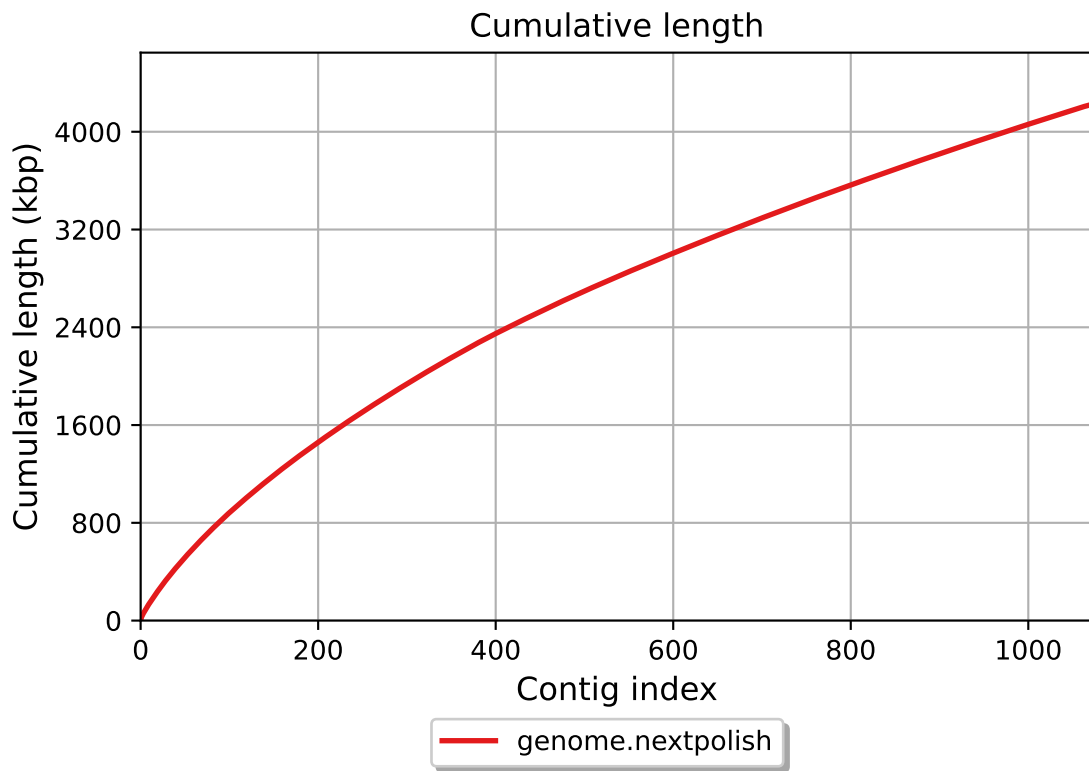
	genome.nextpolish
# contigs (>= 0 bp)	1073
# contigs (>= 1000 bp)	1073
# contigs (>= 5000 bp)	221
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4224950
Total length (>= 1000 bp)	4224950
Total length (>= 5000 bp)	1566169
Total length (>= 10000 bp)	268140
Total length (>= 25000 bp)	28491
Total length (>= 50000 bp)	0
# contigs	1073
Largest contig	28491
Total length	4224950
GC (%)	54.91
N50	4152
N90	2486
auN	5009.0
L50	341
L90	894
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

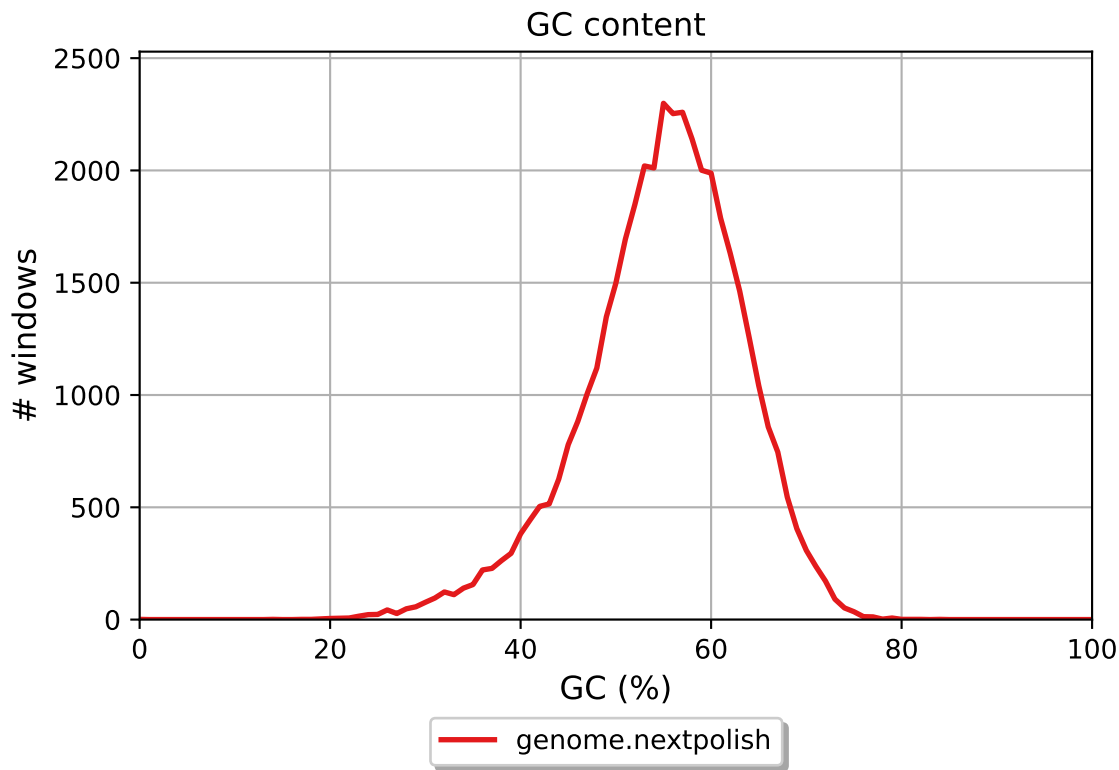
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

