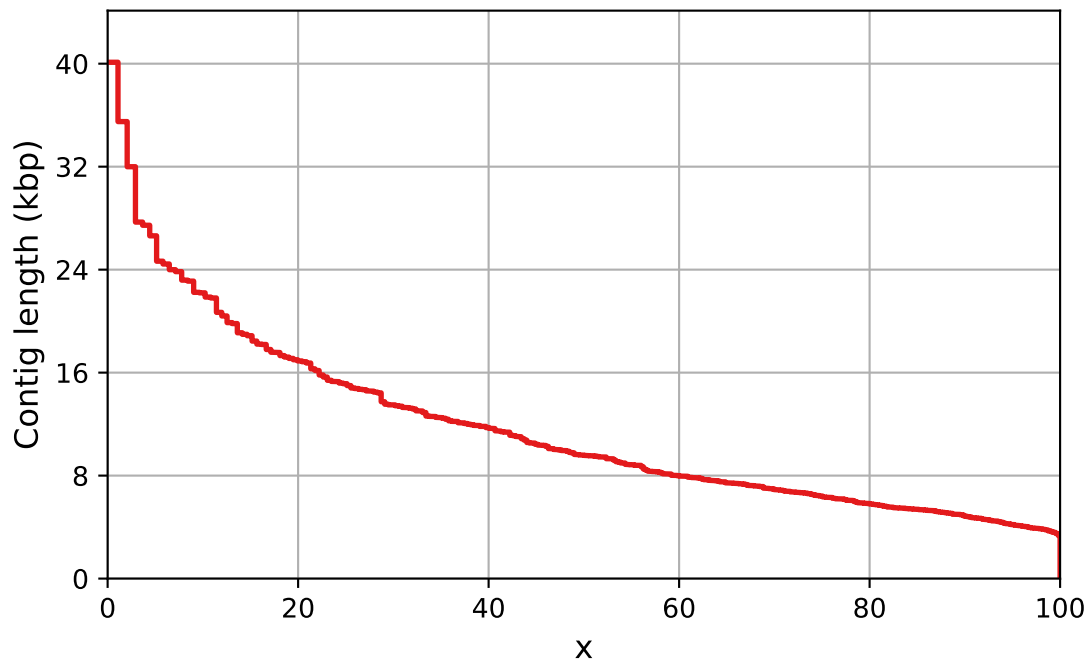


Report

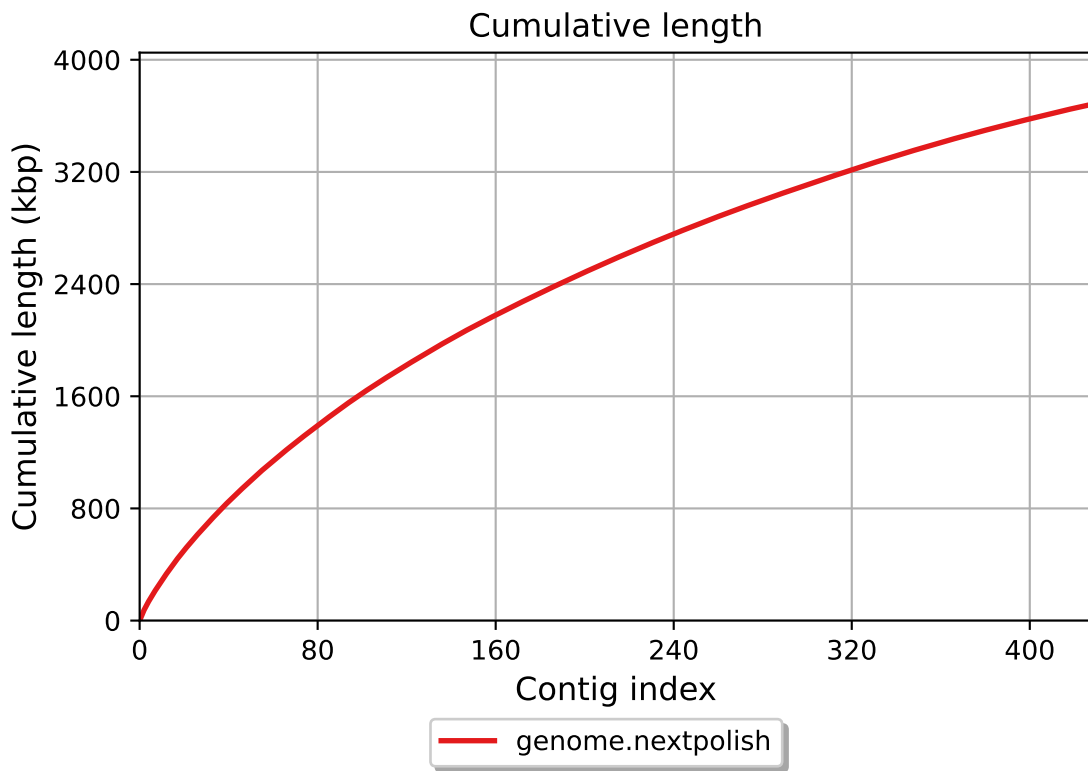
	genome.nextpolish
# contigs (>= 0 bp)	428
# contigs (>= 1000 bp)	428
# contigs (>= 5000 bp)	331
# contigs (>= 10000 bp)	112
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3682621
Total length (>= 1000 bp)	3682621
Total length (>= 5000 bp)	3271118
Total length (>= 10000 bp)	1744375
Total length (>= 25000 bp)	189377
Total length (>= 50000 bp)	0
# contigs	428
Largest contig	40110
Total length	3682621
GC (%)	55.91
N50	9588
N90	4854
auN	11761.8
L50	122
L90	340
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

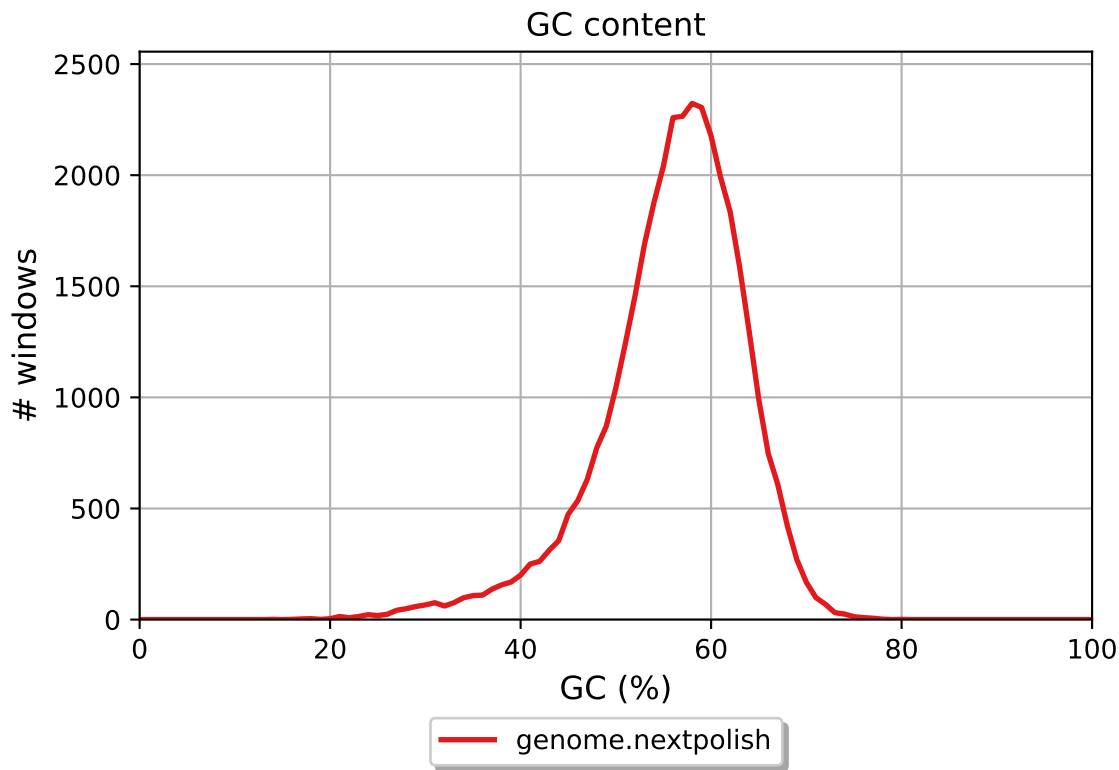
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

