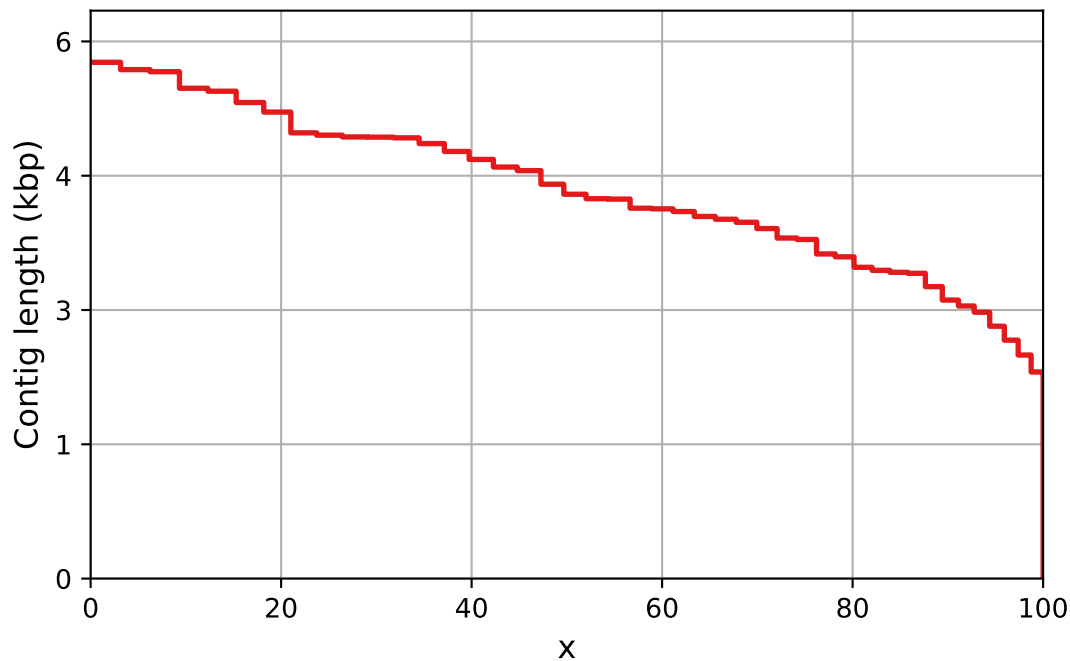


Report

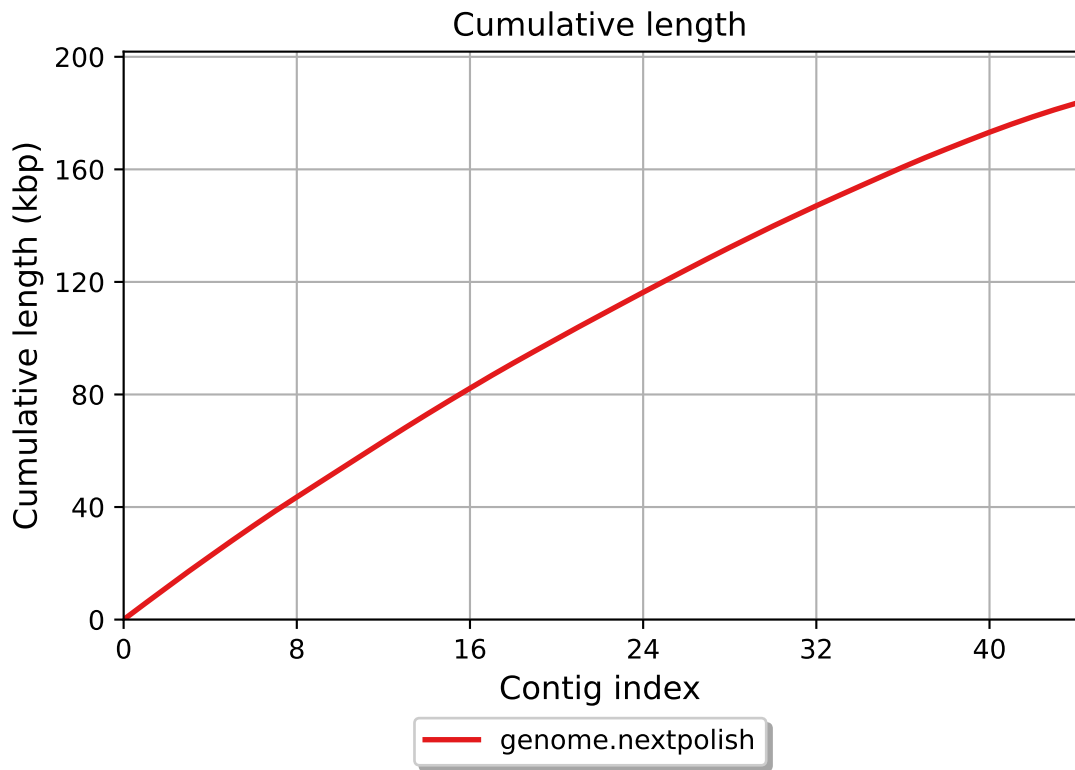
	genome.nextpolish
# contigs (>= 0 bp)	44
# contigs (>= 1000 bp)	44
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	183487
Total length (>= 1000 bp)	183487
Total length (>= 5000 bp)	38564
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	44
Largest contig	5767
Total length	183487
GC (%)	52.09
N50	4293
N90	3111
auN	4363.7
L50	19
L90	38
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

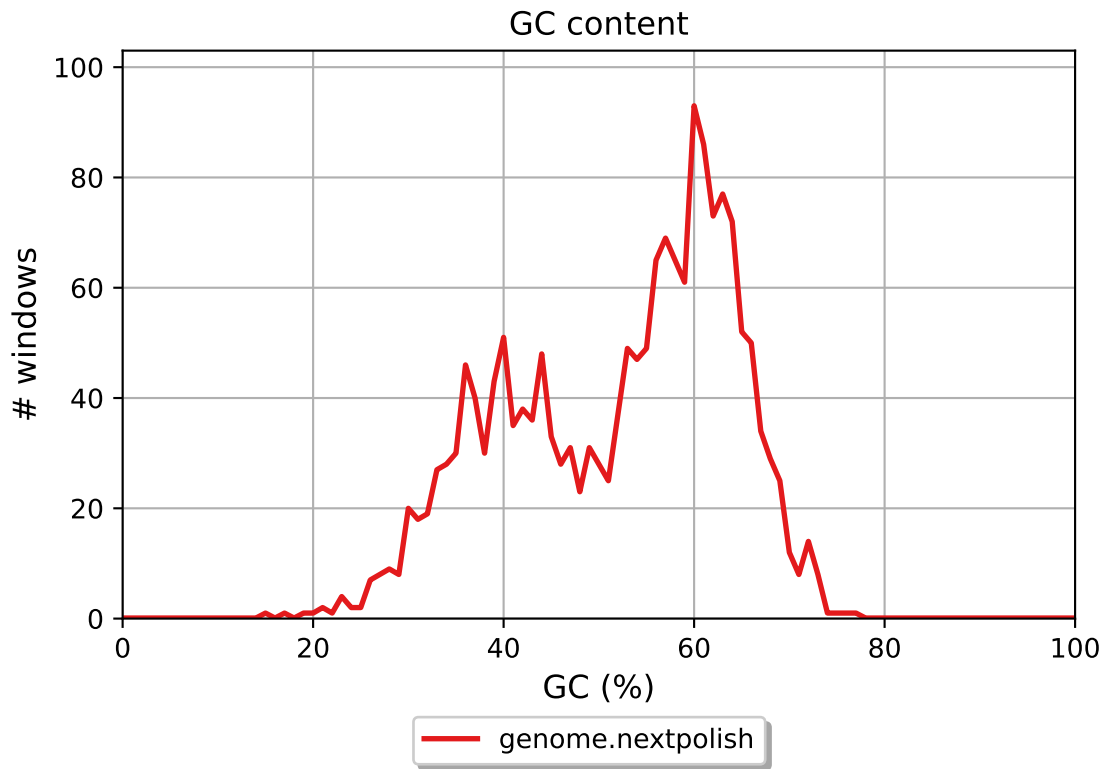
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

