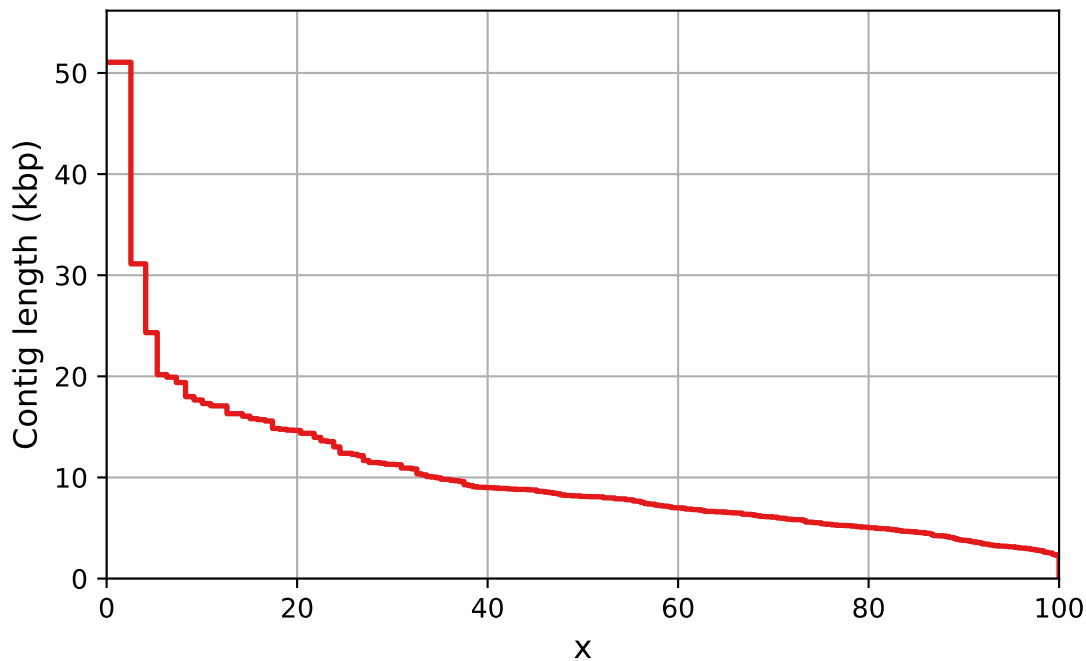


Report

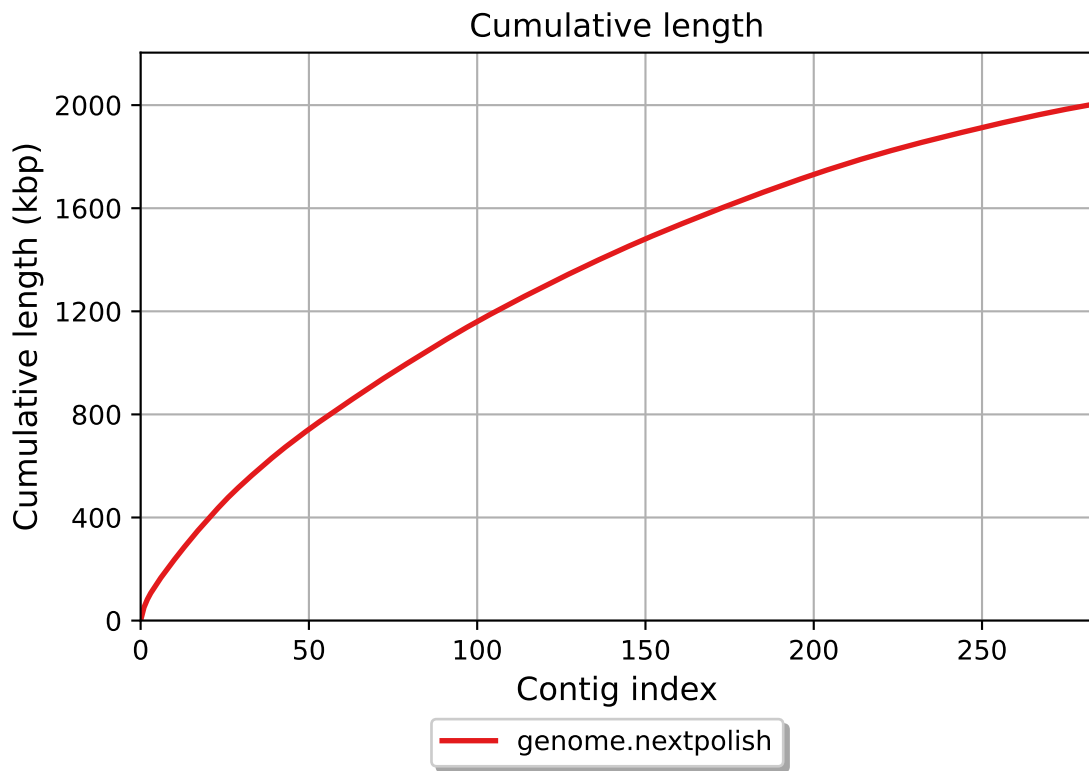
	genome.nextpolish
# contigs (>= 0 bp)	283
# contigs (>= 1000 bp)	283
# contigs (>= 5000 bp)	176
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2003332
Total length (>= 1000 bp)	2003332
Total length (>= 5000 bp)	1617875
Total length (>= 10000 bp)	693288
Total length (>= 25000 bp)	82185
Total length (>= 50000 bp)	51059
# contigs	283
Largest contig	51059
Total length	2003332
GC (%)	39.69
N50	8132
N90	3782
auN	10449.0
L50	80
L90	218
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

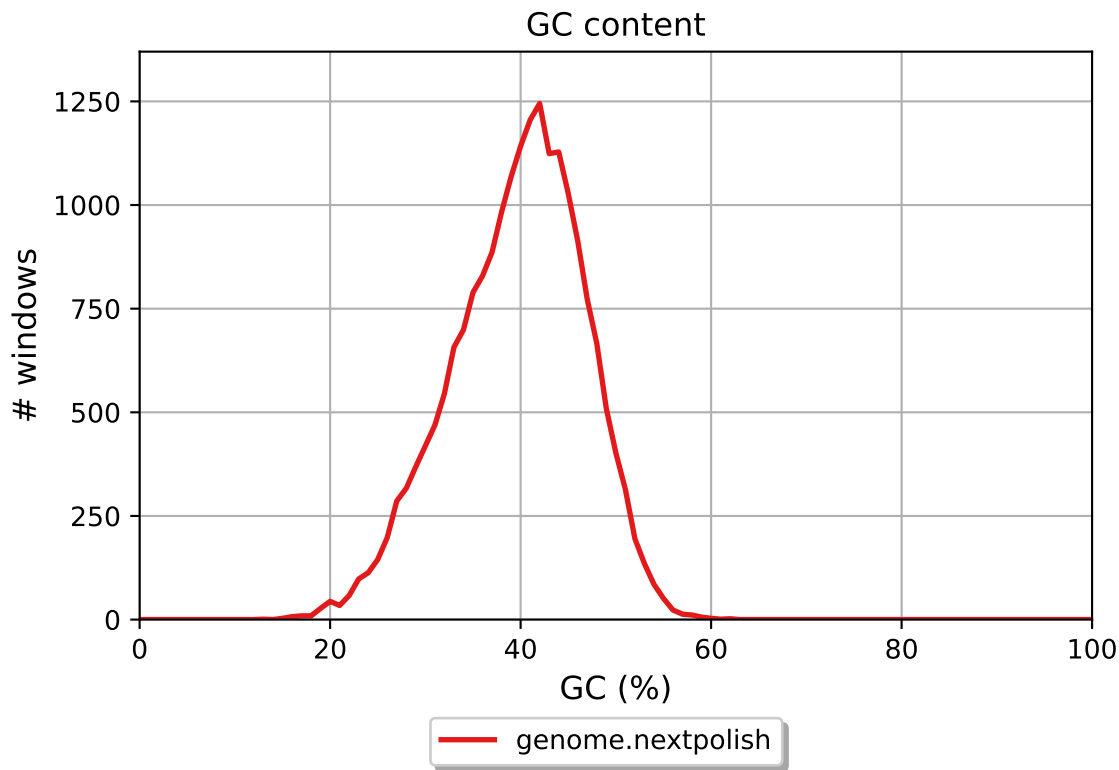
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

