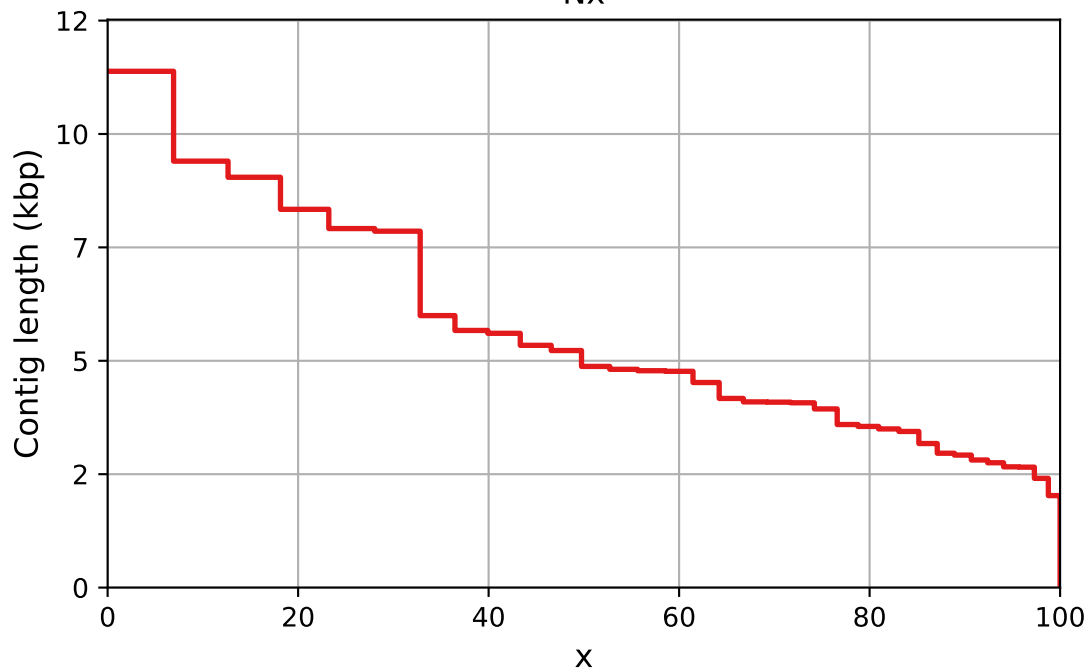


# Report

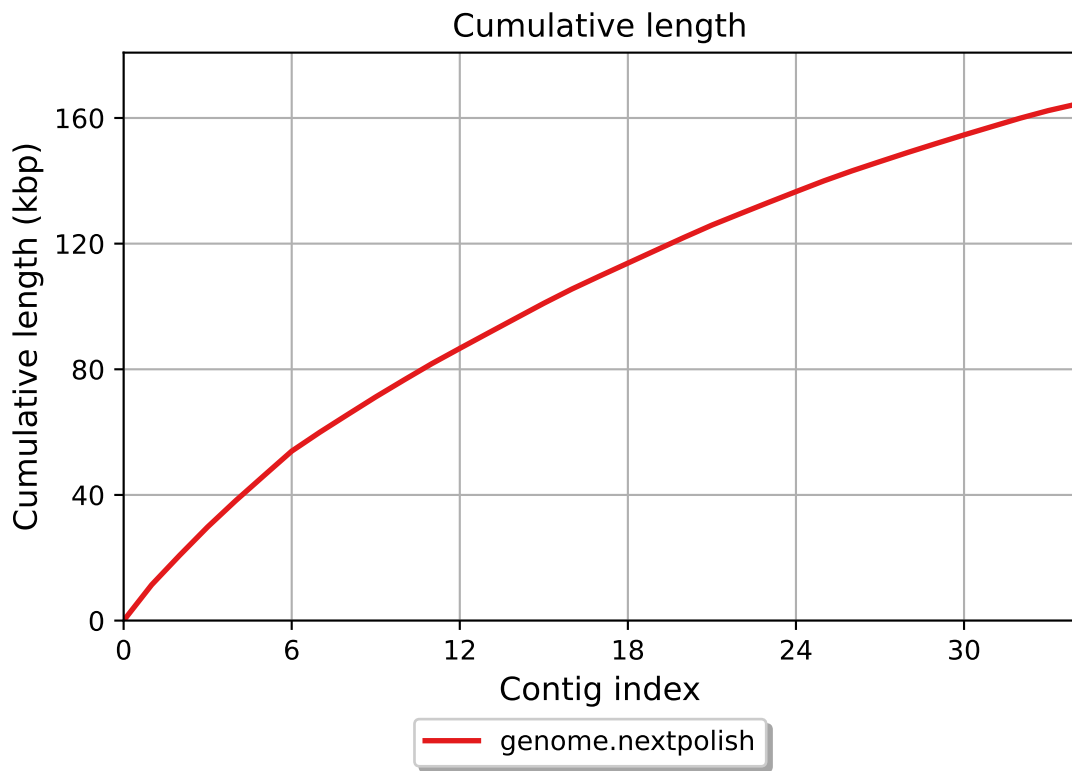
	genome.nextpolish
# contigs (>= 0 bp)	34
# contigs (>= 1000 bp)	34
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	164366
Total length (>= 1000 bp)	164366
Total length (>= 5000 bp)	81778
Total length (>= 10000 bp)	11382
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	34
Largest contig	11382
Total length	164366
GC (%)	40.33
N50	4876
N90	2921
auN	5849.9
L50	12
L90	28
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

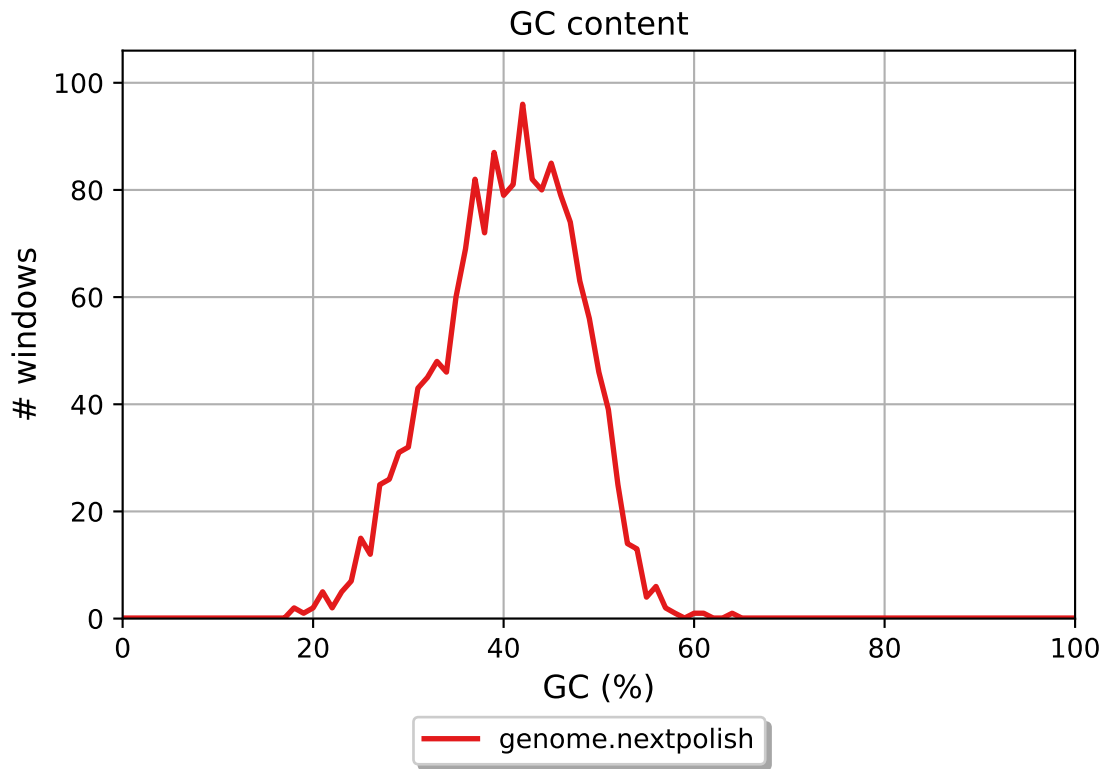
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

