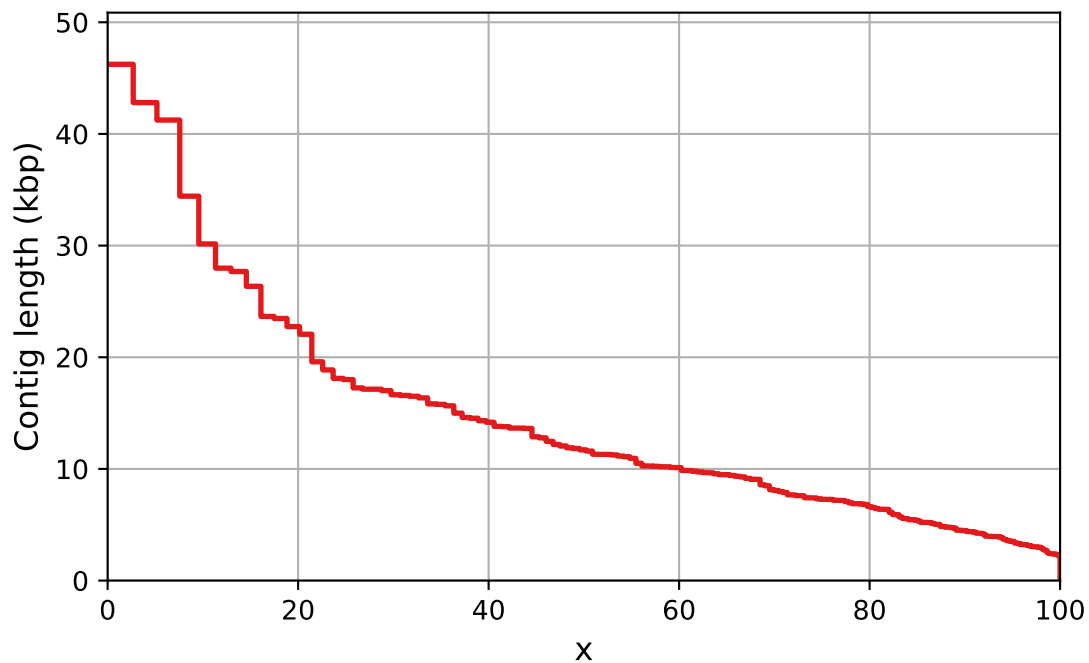


# Report

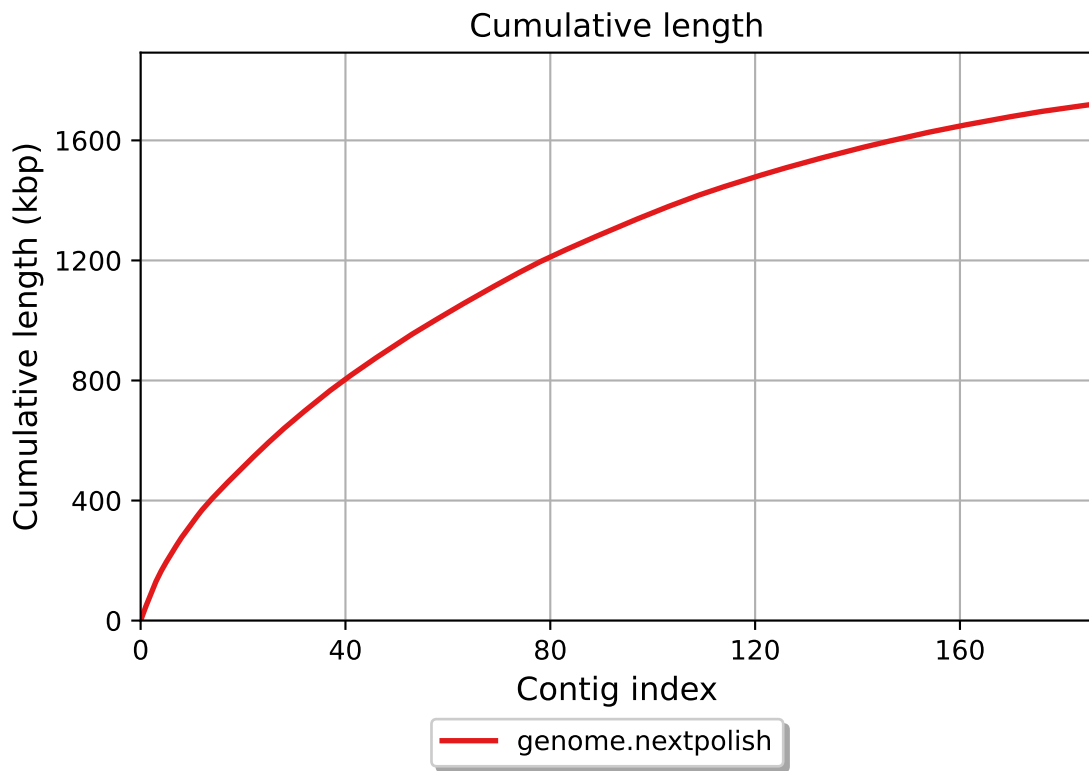
	genome.nextpolish
# contigs (>= 0 bp)	186
# contigs (>= 1000 bp)	186
# contigs (>= 5000 bp)	125
# contigs (>= 10000 bp)	61
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1720349
Total length (>= 1000 bp)	1720349
Total length (>= 5000 bp)	1503835
Total length (>= 10000 bp)	1036095
Total length (>= 25000 bp)	276875
Total length (>= 50000 bp)	0
# contigs	186
Largest contig	46238
Total length	1720349
GC (%)	38.14
N50	11709
N90	4478
auN	15076.9
L50	45
L90	135
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 part

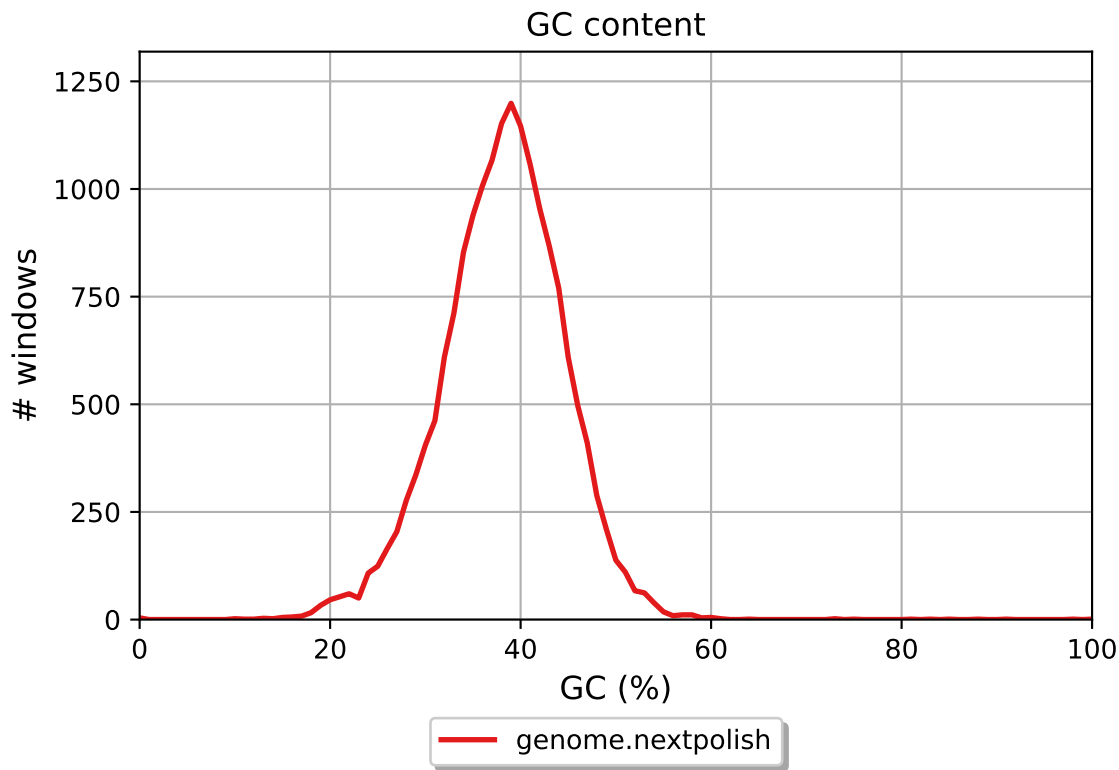
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

