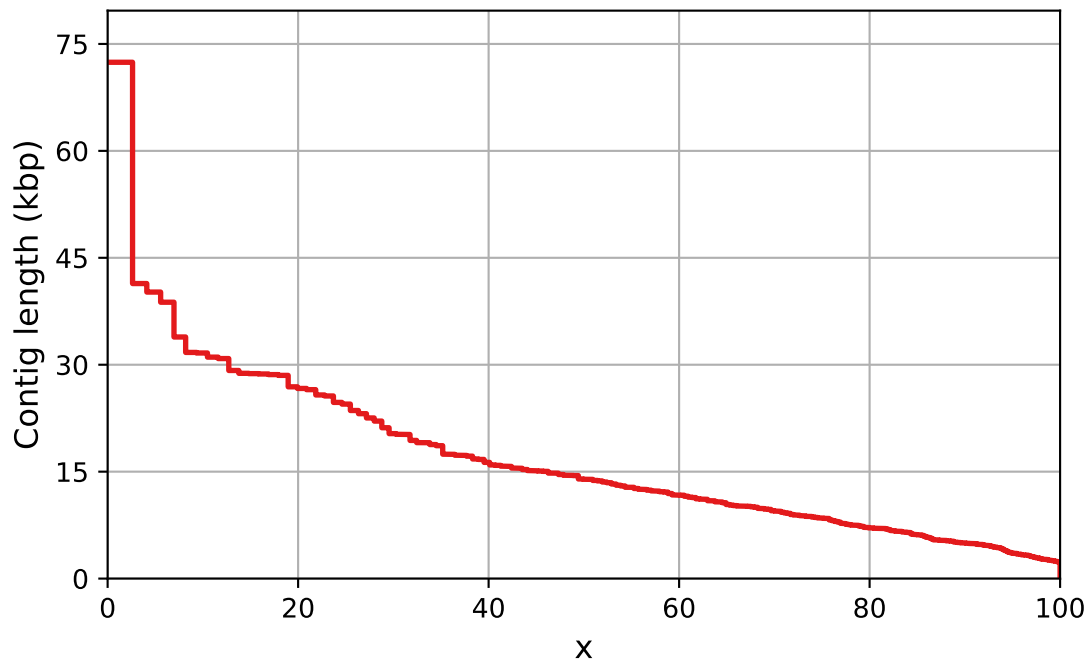


Report

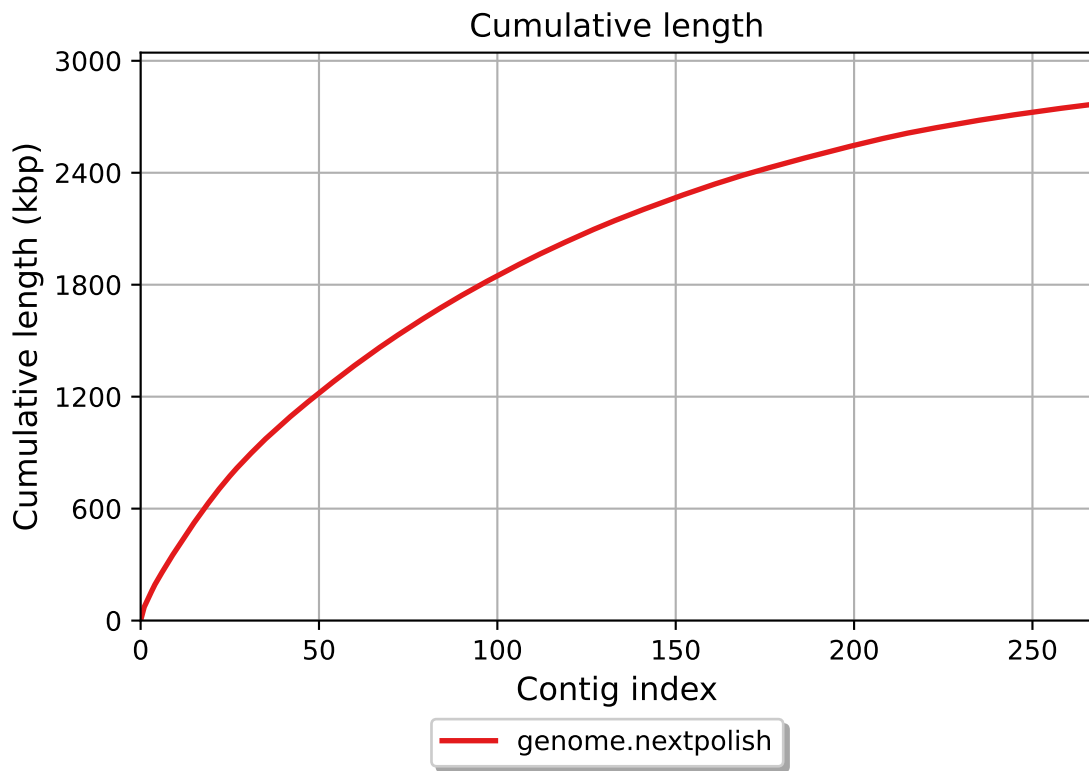
	genome.nextpolish
# contigs (>= 0 bp)	267
# contigs (>= 1000 bp)	267
# contigs (>= 5000 bp)	188
# contigs (>= 10000 bp)	104
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2767030
Total length (>= 1000 bp)	2767030
Total length (>= 5000 bp)	2488315
Total length (>= 10000 bp)	1888245
Total length (>= 25000 bp)	655995
Total length (>= 50000 bp)	72430
# contigs	267
Largest contig	72430
Total length	2767030
GC (%)	33.30
N50	13932
N90	4997
auN	17463.8
L50	62
L90	189
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

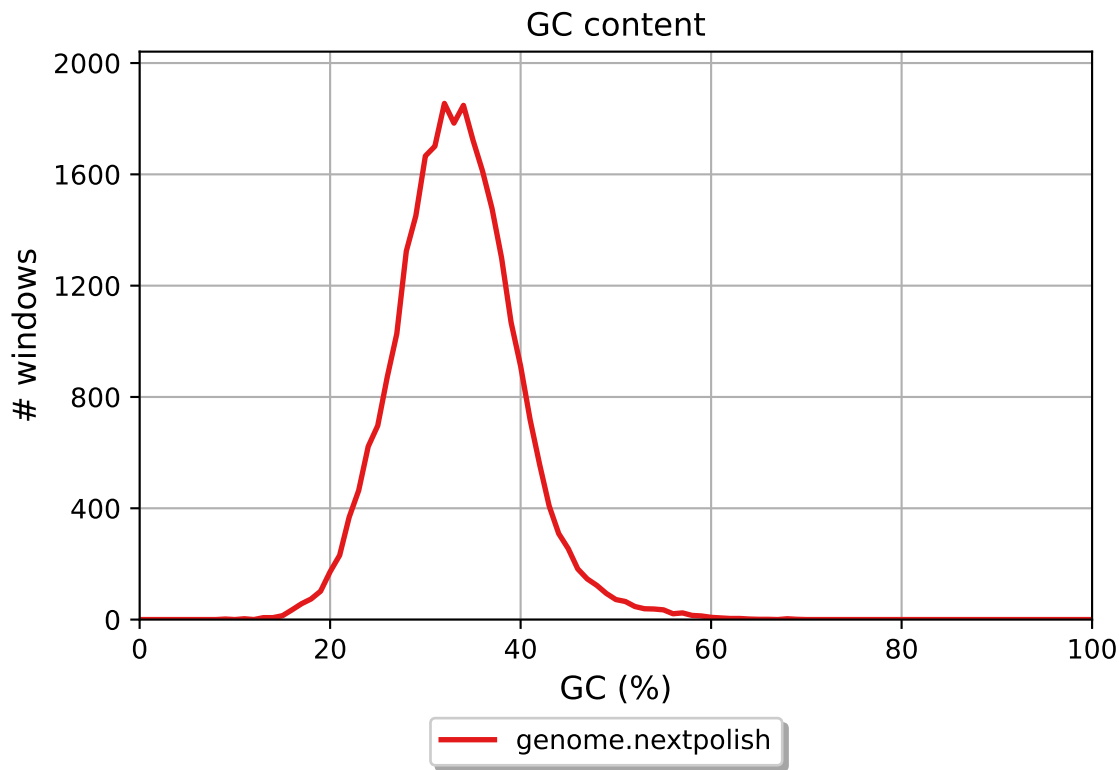
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



genome.nextpolish





genome.nextpolish GC content

