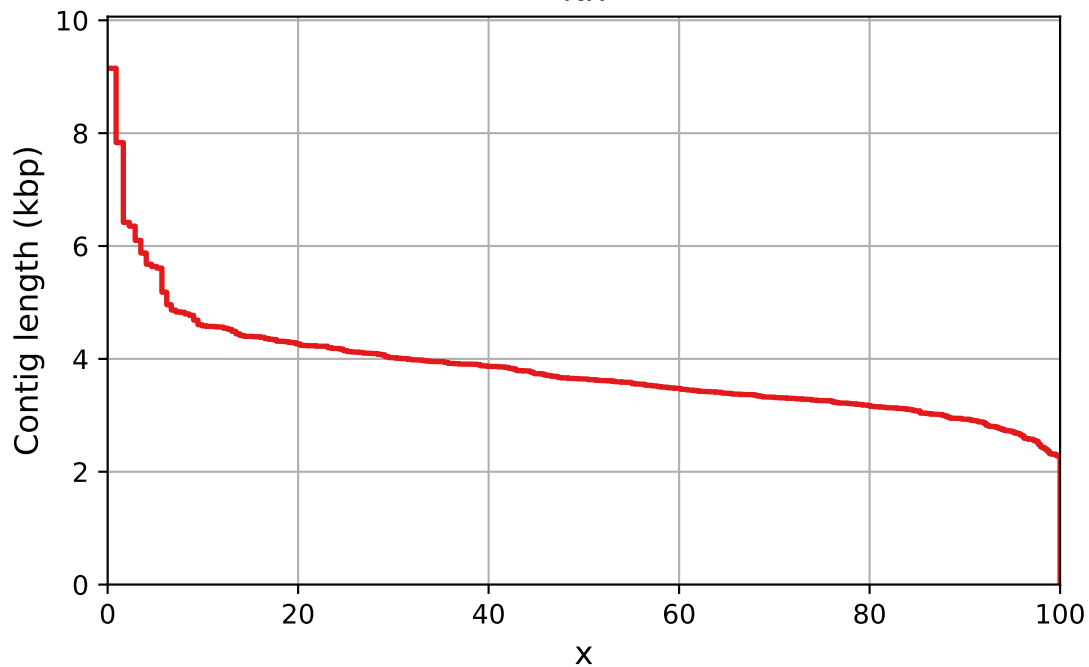


Report

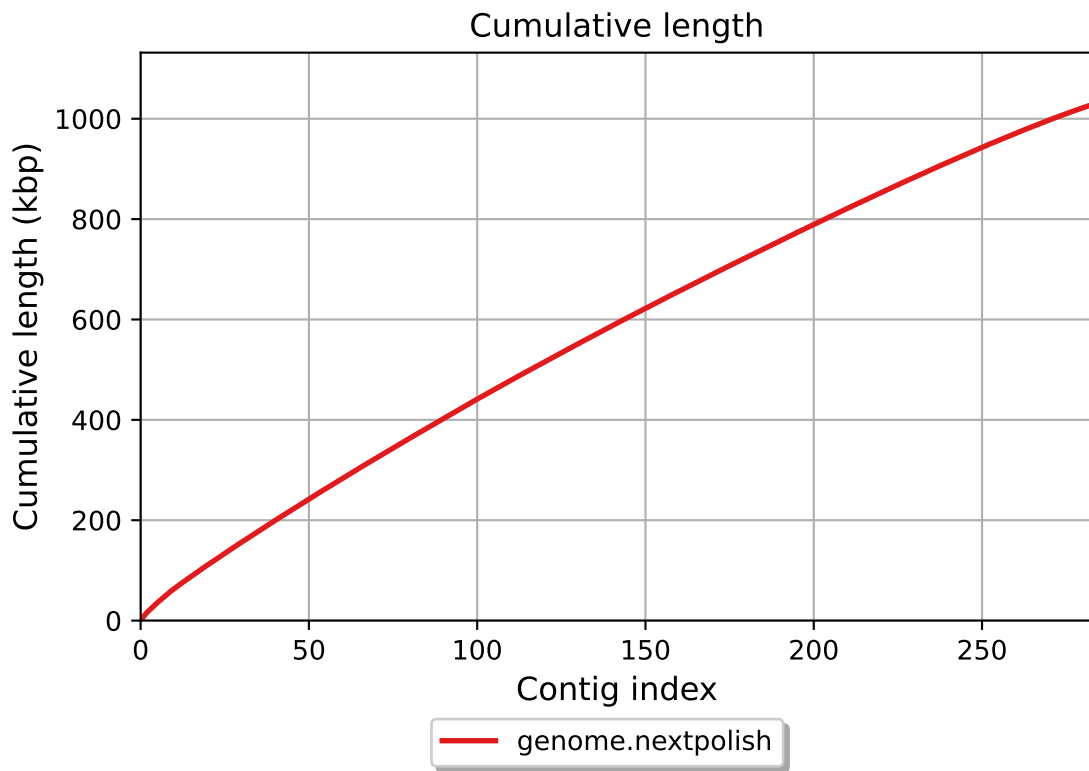
	genome.nextpolish
# contigs (>= 0 bp)	283
# contigs (>= 1000 bp)	283
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1028841
Total length (>= 1000 bp)	1028841
Total length (>= 5000 bp)	63836
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	283
Largest contig	9150
Total length	1028841
GC (%)	44.66
N50	3645
N90	2930
auN	3811.9
L50	120
L90	245
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 1 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).

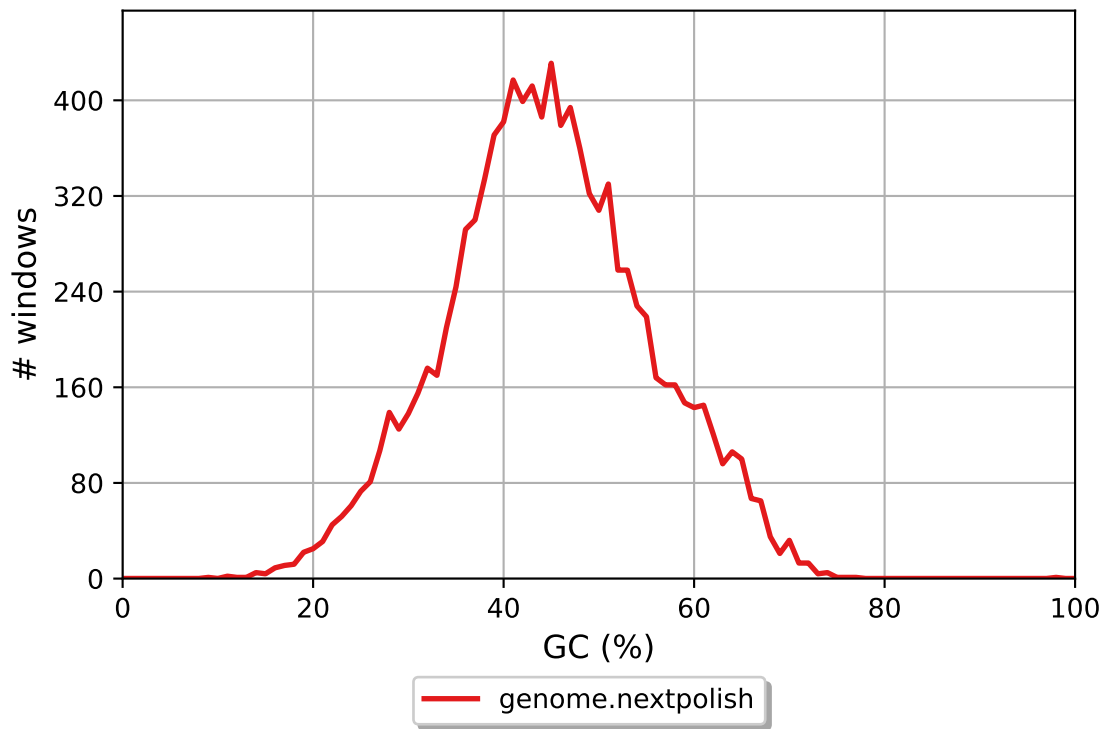
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

