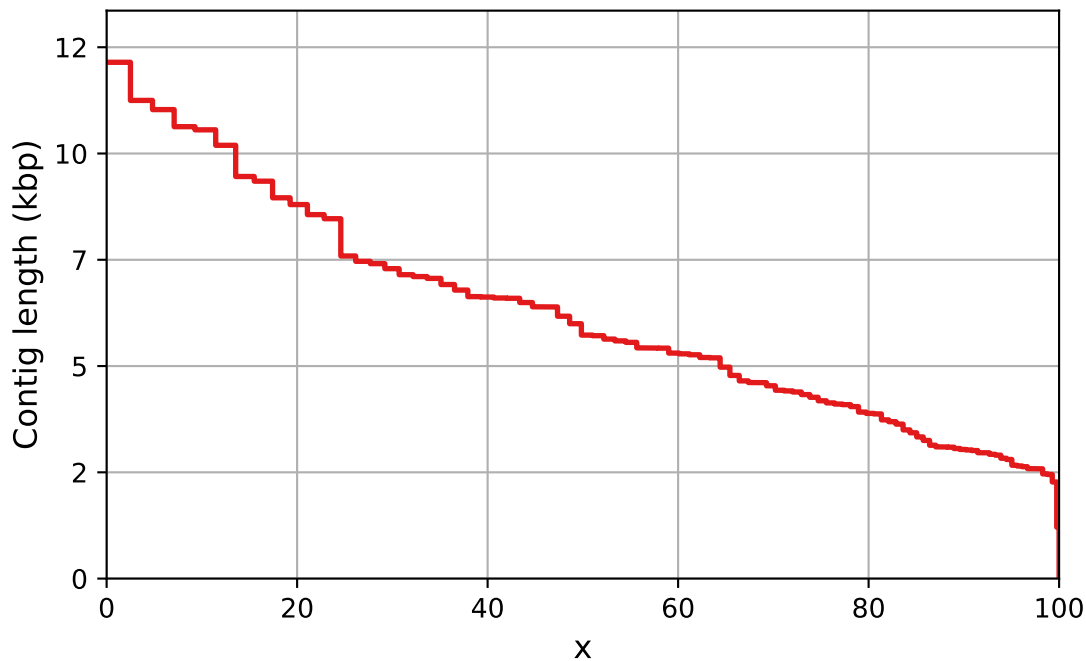


Report

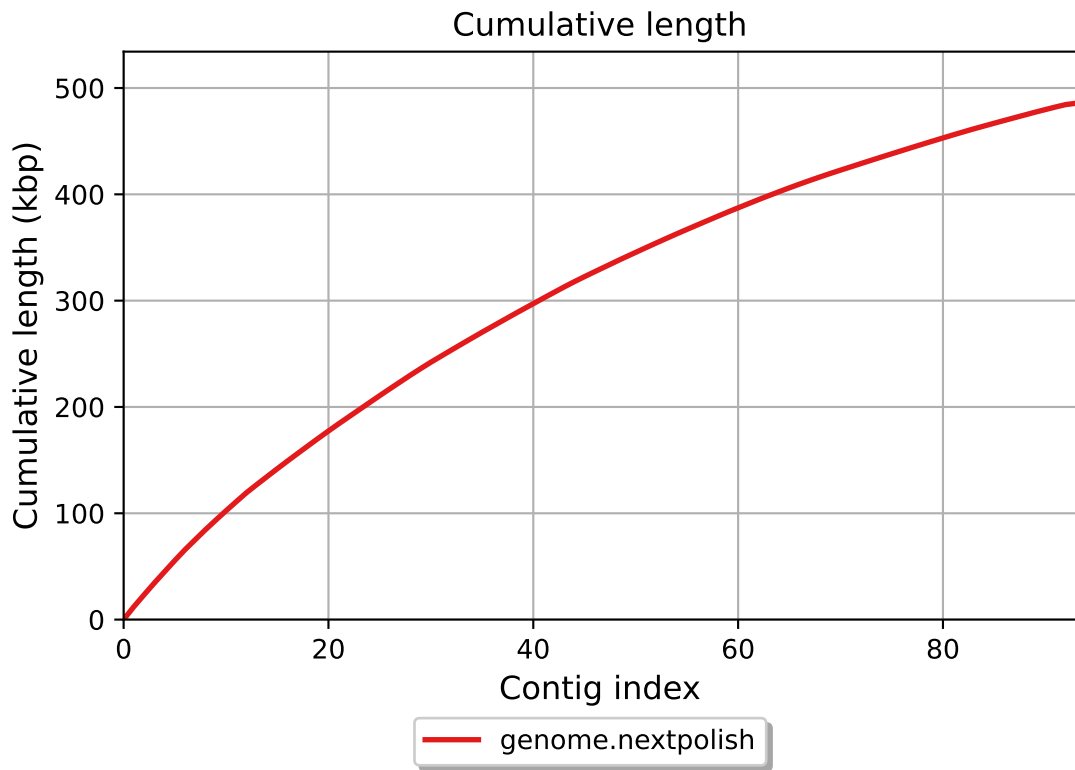
	genome.nextpolish
# contigs (>= 0 bp)	93
# contigs (>= 1000 bp)	93
# contigs (>= 5000 bp)	43
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	485651
Total length (>= 1000 bp)	485651
Total length (>= 5000 bp)	312821
Total length (>= 10000 bp)	65809
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	93
Largest contig	12146
Total length	485651
GC (%)	40.77
N50	5728
N90	3038
auN	6297.4
L50	31
L90	75
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

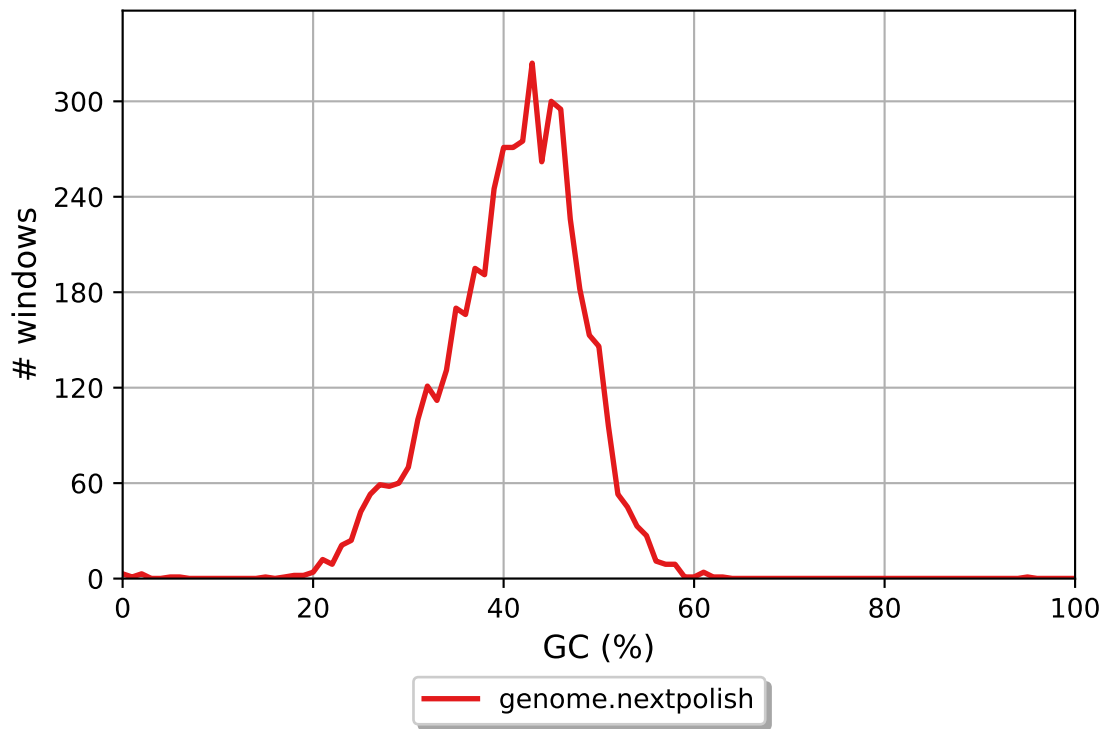
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

