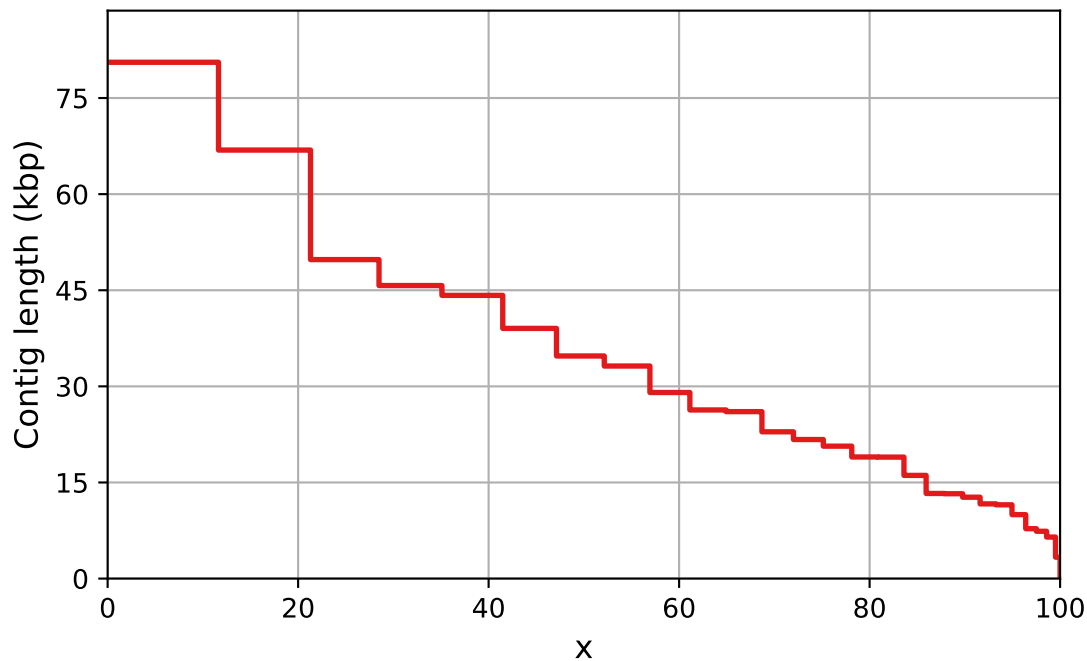


Report

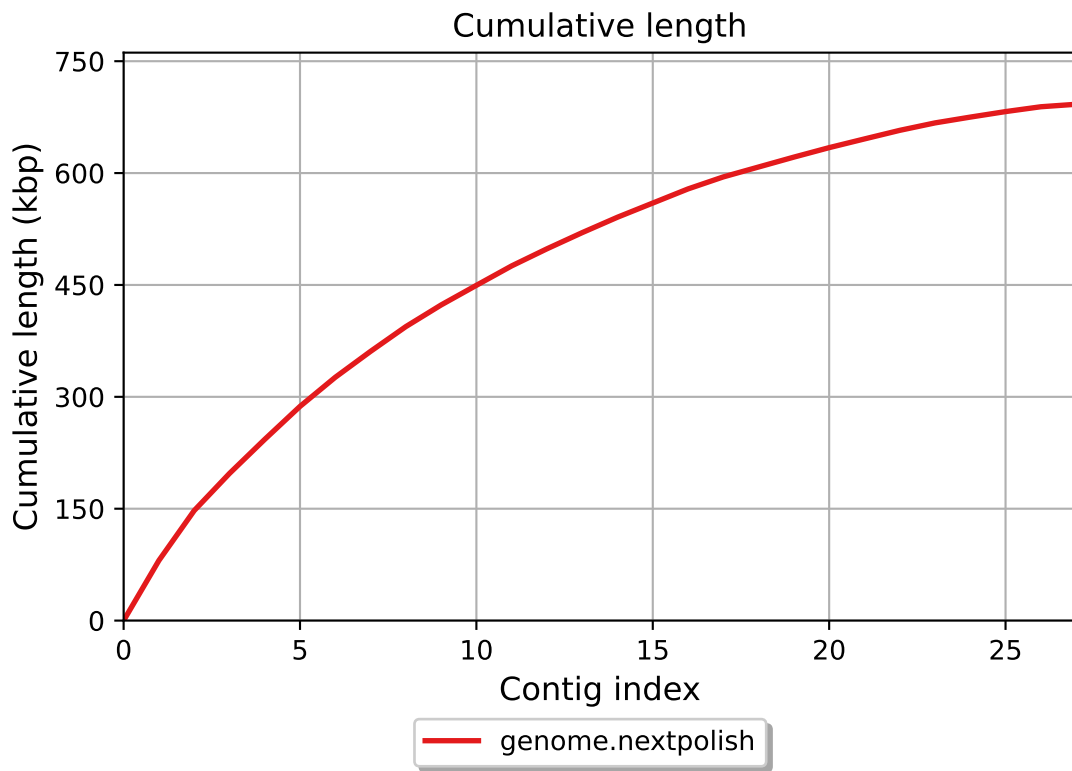
	genome.nextpolish
# contigs (>= 0 bp)	27
# contigs (>= 1000 bp)	27
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	692251
Total length (>= 1000 bp)	692251
Total length (>= 5000 bp)	688913
Total length (>= 10000 bp)	657280
Total length (>= 25000 bp)	475551
Total length (>= 50000 bp)	147458
# contigs	27
Largest contig	80580
Total length	692251
GC (%)	54.70
N50	34741
N90	12702
auN	38985.9
L50	7
L90	20
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

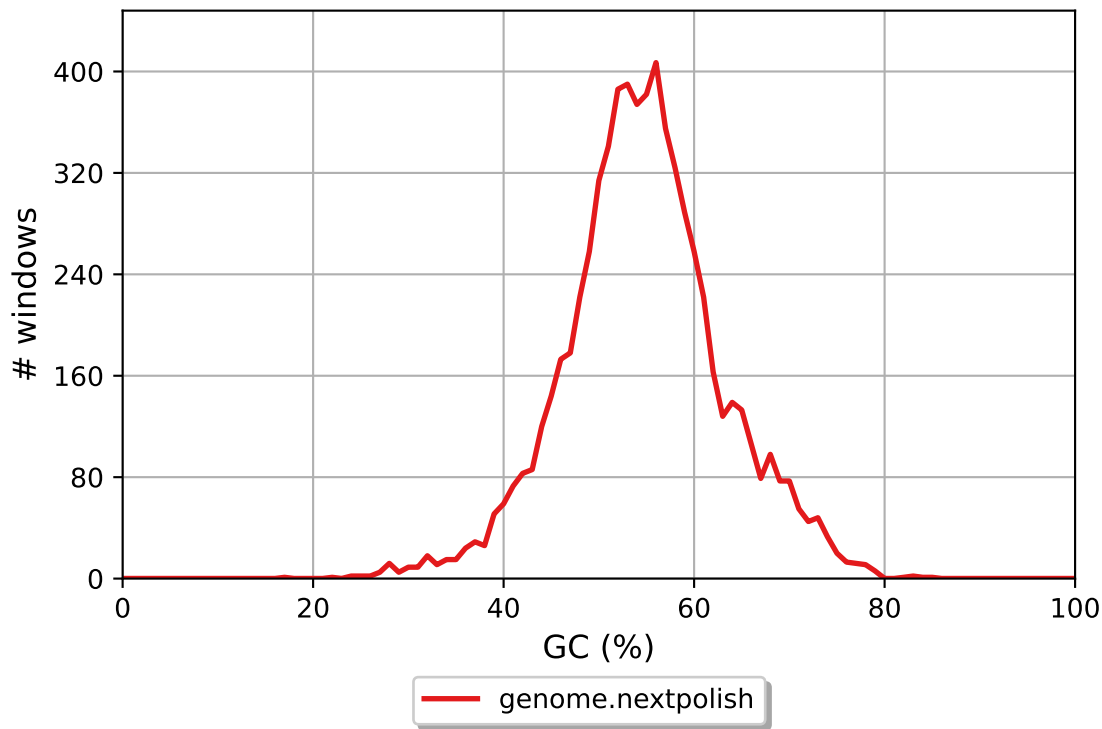
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

