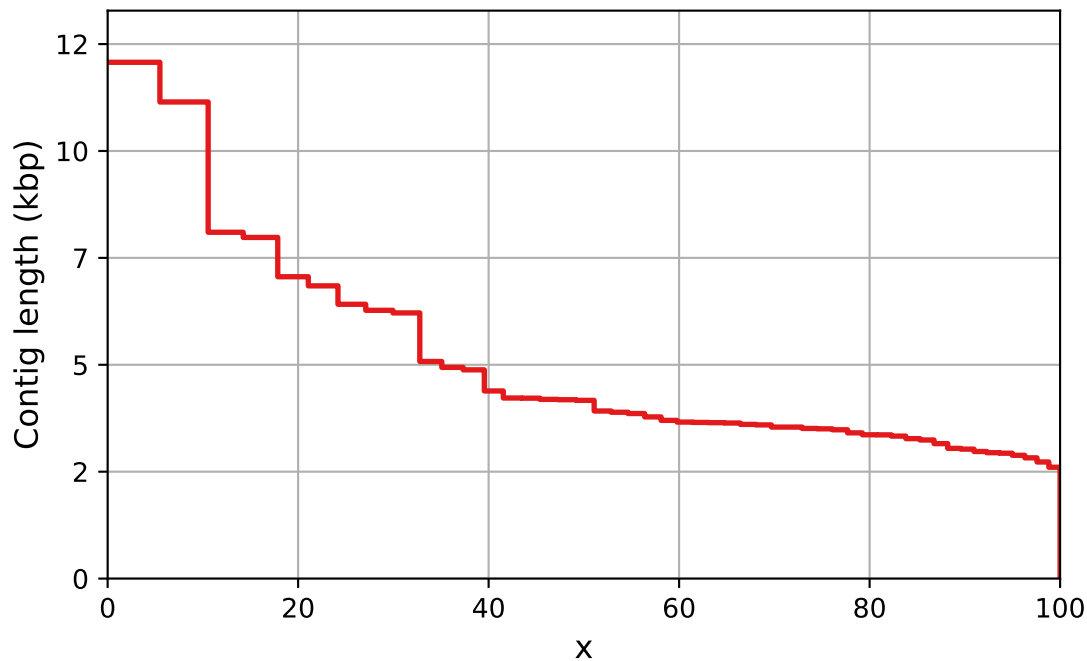


Report

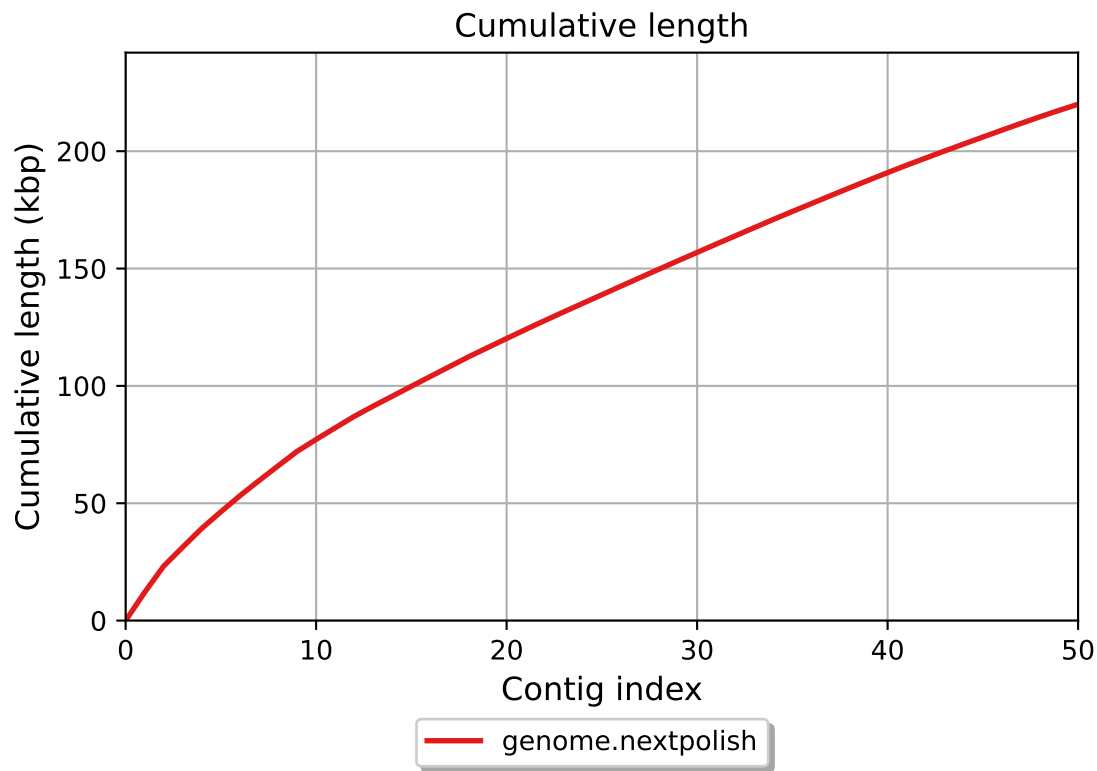
	genome.nextpolish
# contigs (>= 0 bp)	50
# contigs (>= 1000 bp)	50
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	219991
Total length (>= 1000 bp)	219991
Total length (>= 5000 bp)	77181
Total length (>= 10000 bp)	23220
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	50
Largest contig	12074
Total length	219991
GC (%)	56.46
N50	4168
N90	3027
auN	5283.4
L50	18
L90	43
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

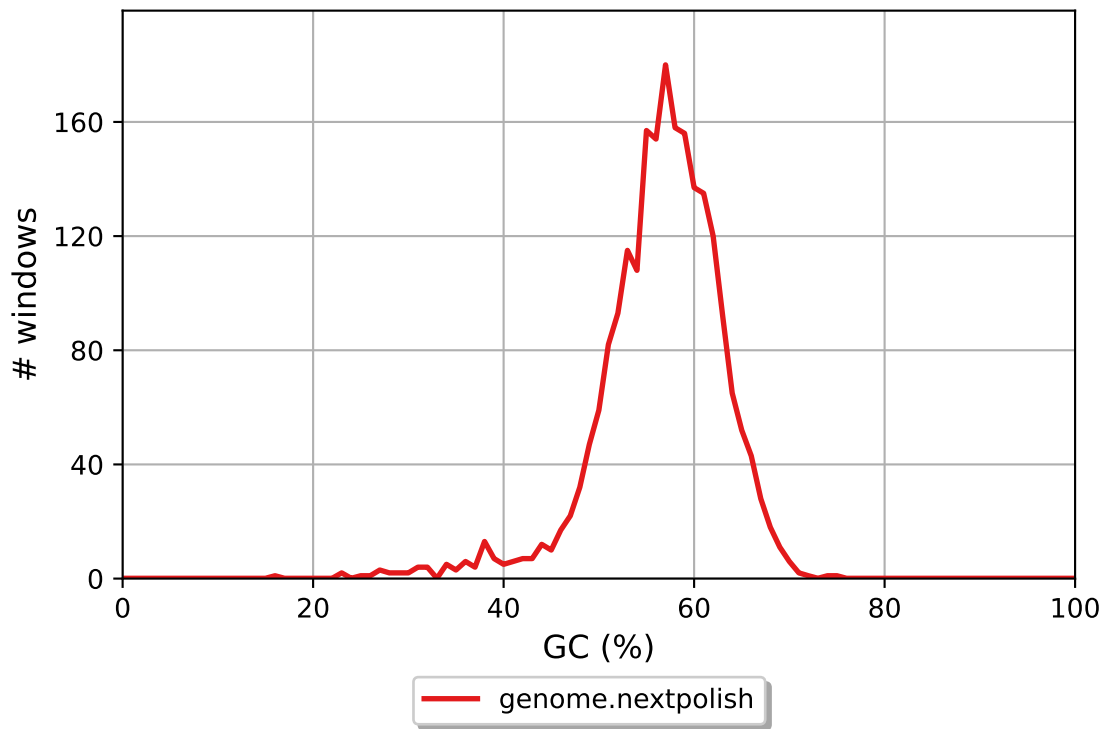
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

