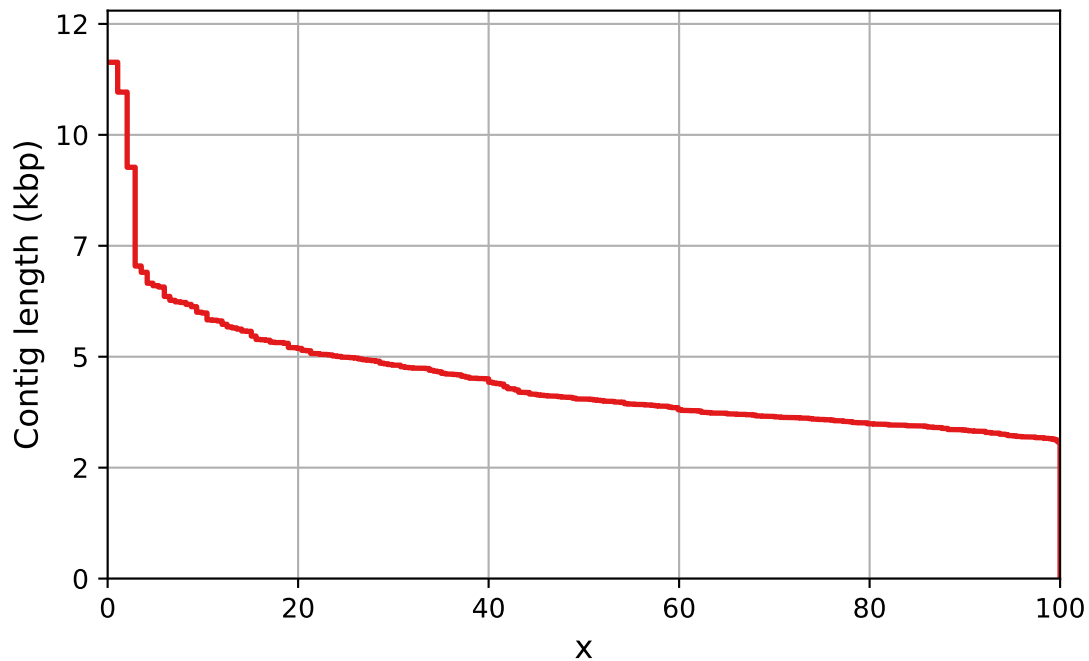


Report

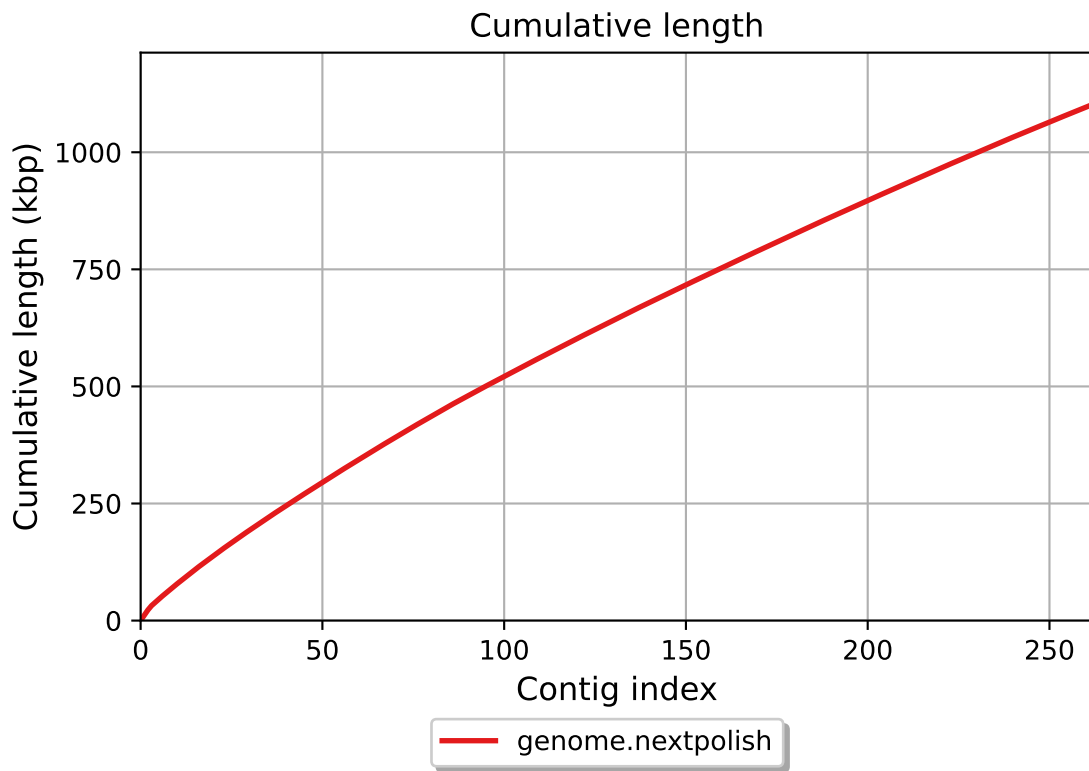
	genome.nextpolish
# contigs (>= 0 bp)	262
# contigs (>= 1000 bp)	262
# contigs (>= 5000 bp)	45
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1102470
Total length (>= 1000 bp)	1102470
Total length (>= 5000 bp)	270357
Total length (>= 10000 bp)	22598
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	262
Largest contig	11635
Total length	1102470
GC (%)	40.44
N50	4046
N90	3350
auN	4494.0
L50	108
L90	228
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 3 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).

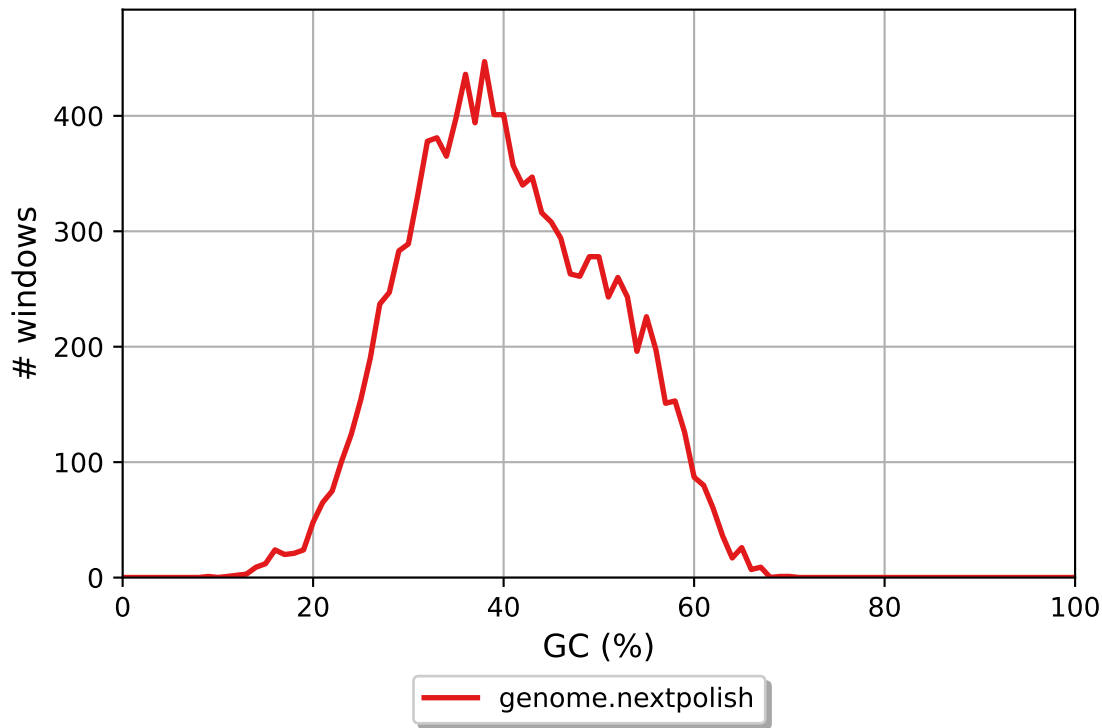
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

