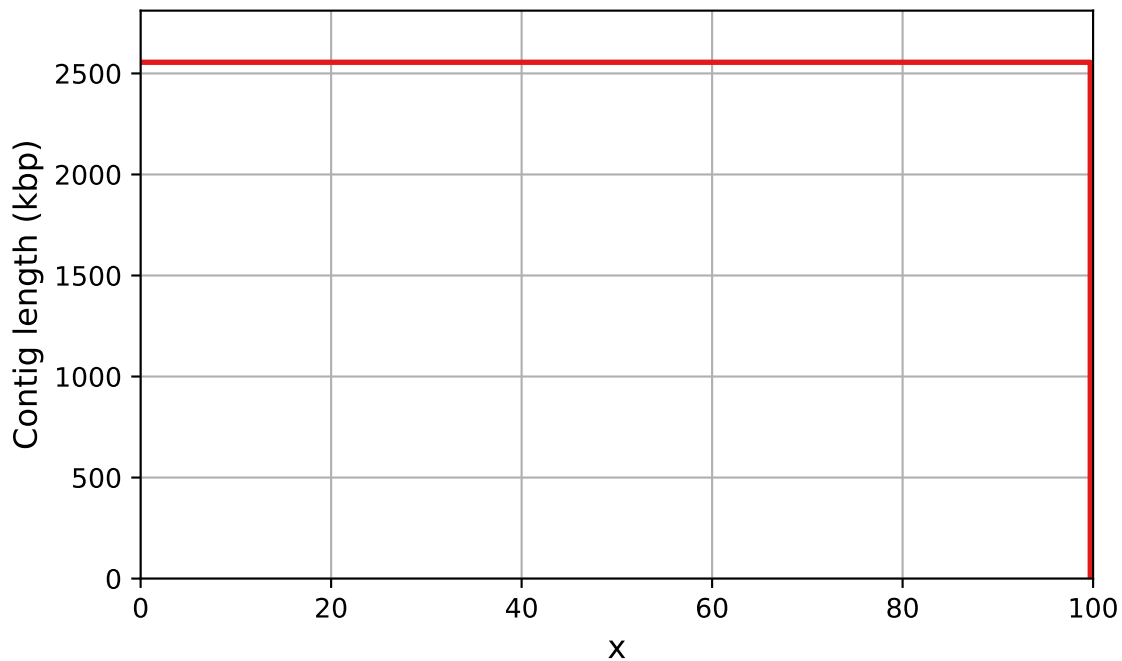


Report

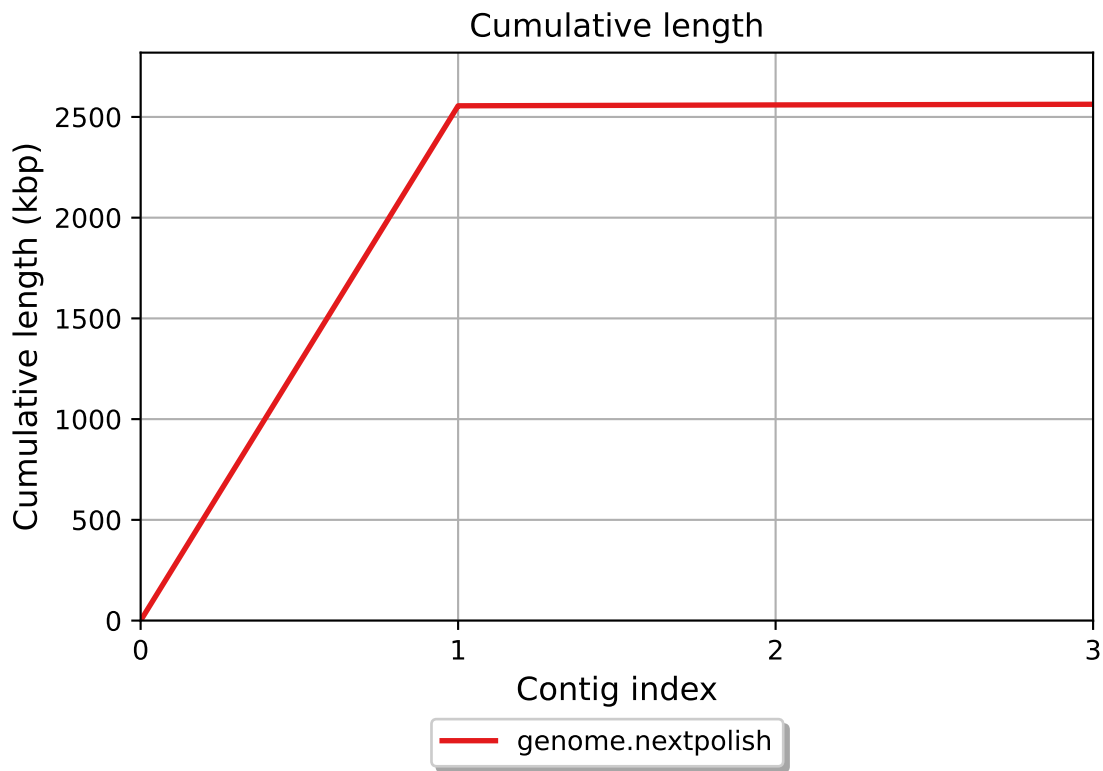
	genome.nextpolish
# contigs (>= 0 bp)	3
# contigs (>= 1000 bp)	3
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2562899
Total length (>= 1000 bp)	2562899
Total length (>= 5000 bp)	2555258
Total length (>= 10000 bp)	2555258
Total length (>= 25000 bp)	2555258
Total length (>= 50000 bp)	2555258
# contigs	3
Largest contig	2555258
Total length	2562899
GC (%)	33.87
N50	2555258
N90	2555258
auN	2547651.4
L50	1
L90	1
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 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).

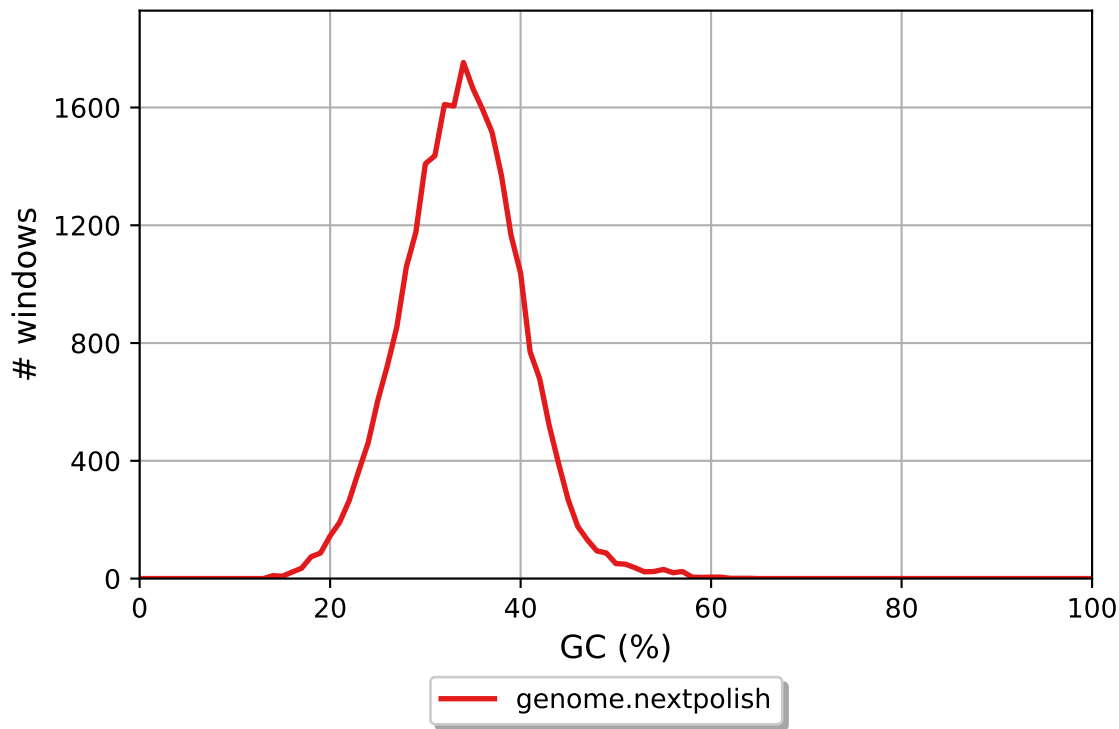
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

