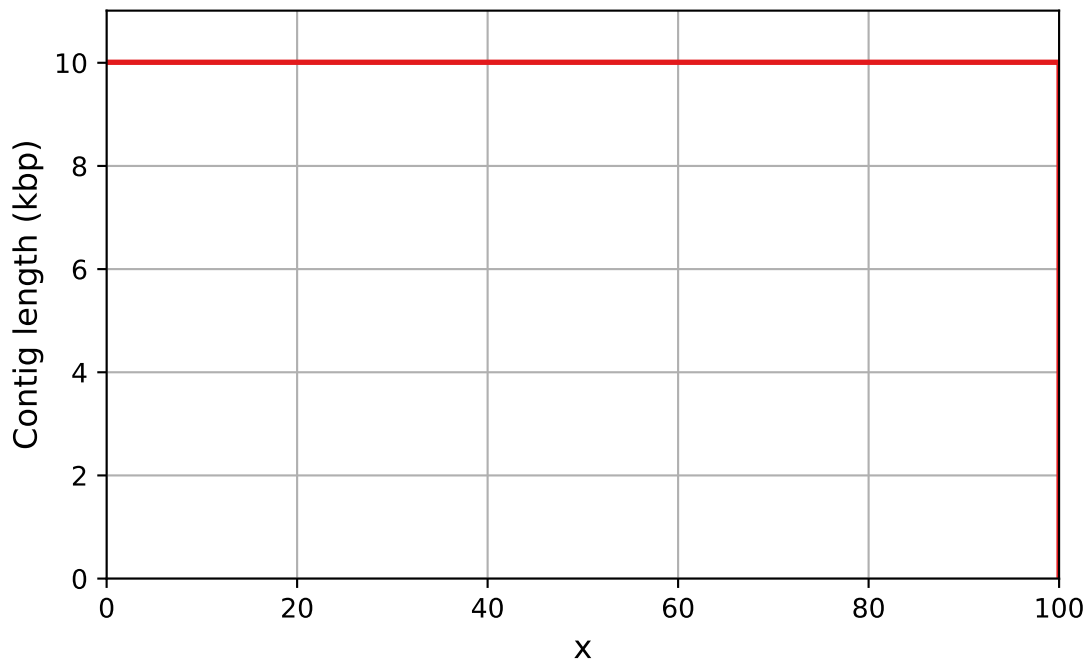


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1
# contigs (>= 1000 bp)	1
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	10010
Total length (>= 1000 bp)	10010
Total length (>= 5000 bp)	10010
Total length (>= 10000 bp)	10010
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1
Largest contig	10010
Total length	10010
GC (%)	42.42
N50	10010
N90	10010
auN	10010.0
L50	1
L90	1
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

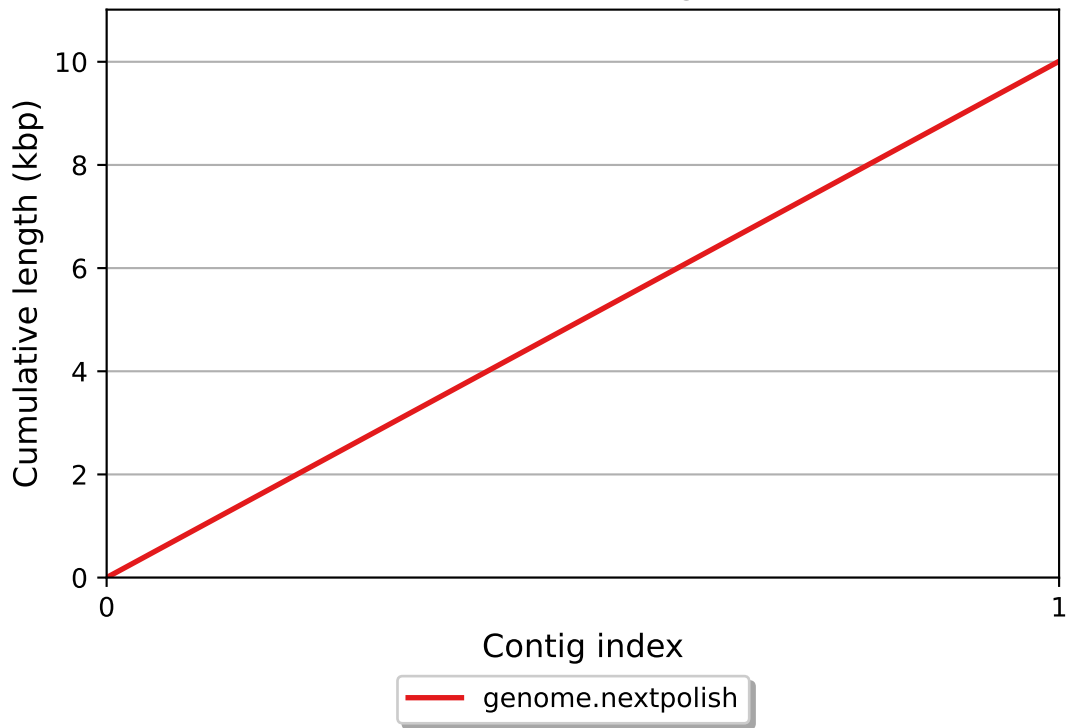
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

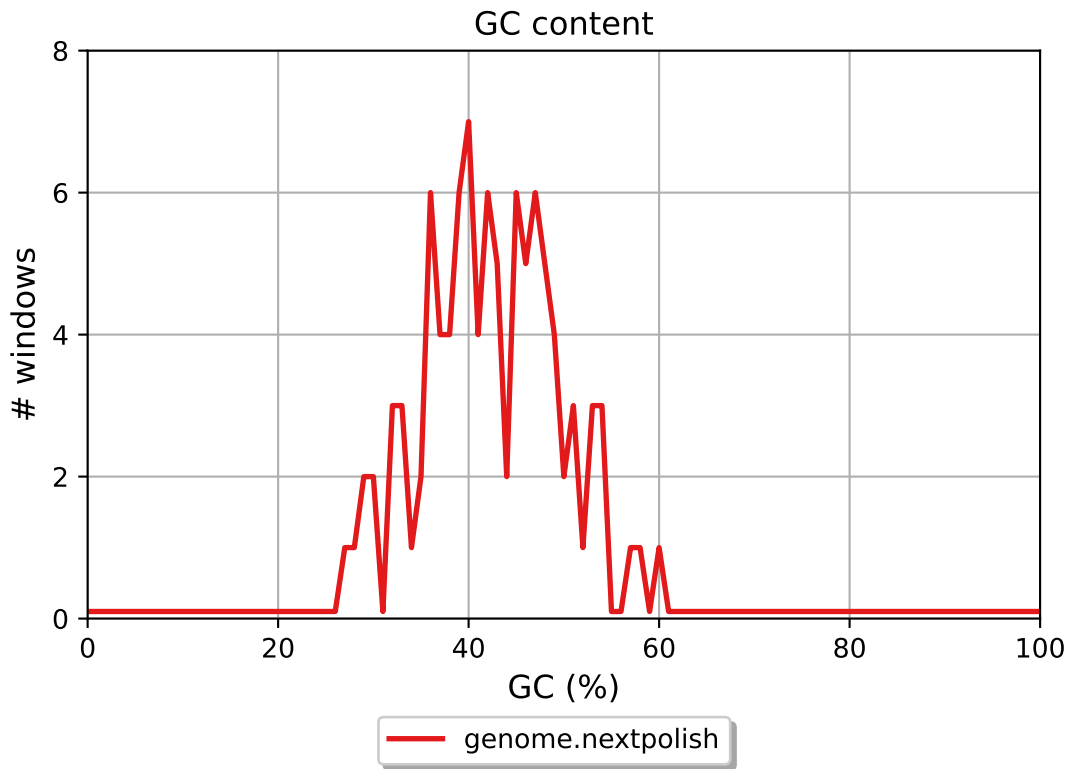
Nx



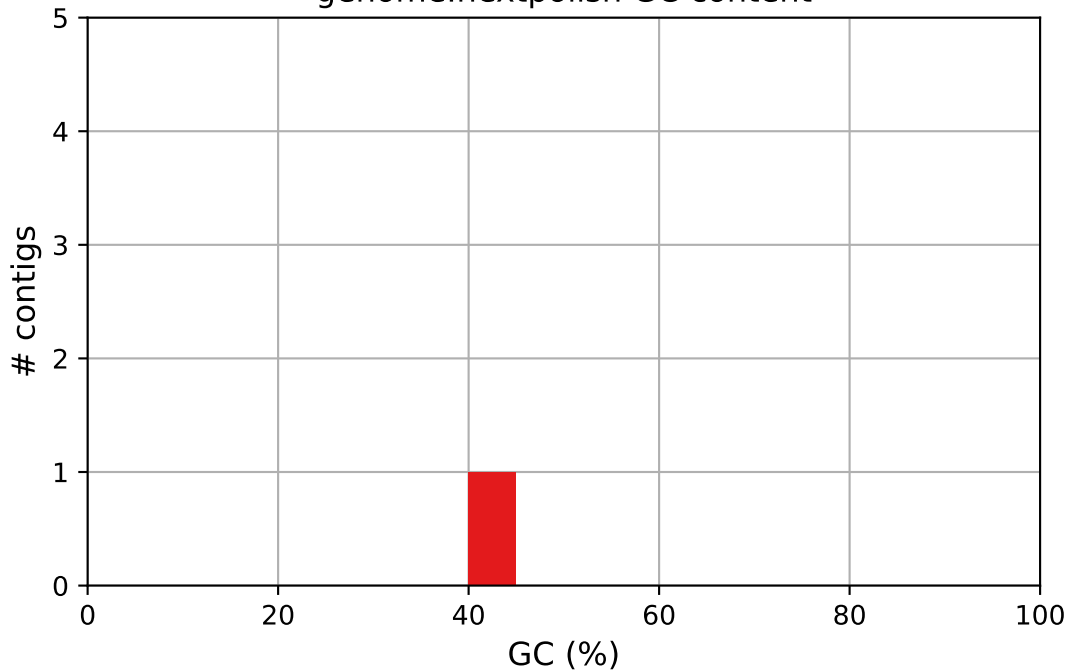
— genome.nextpolish

Cumulative length





genome.nextpolish GC content



genome.nextpolish