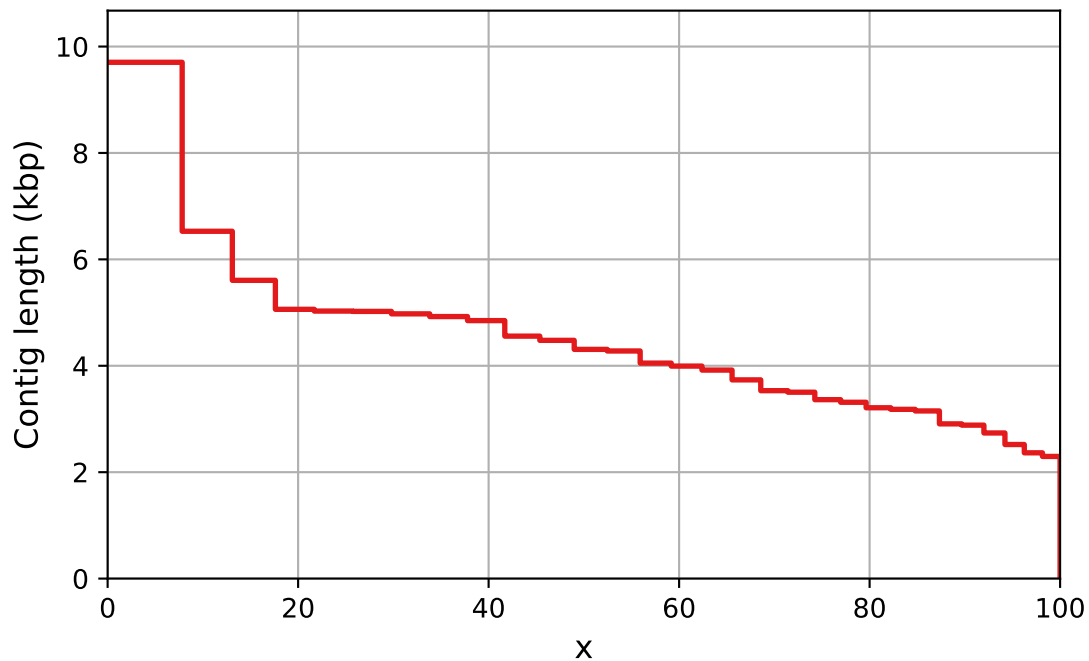


# Report

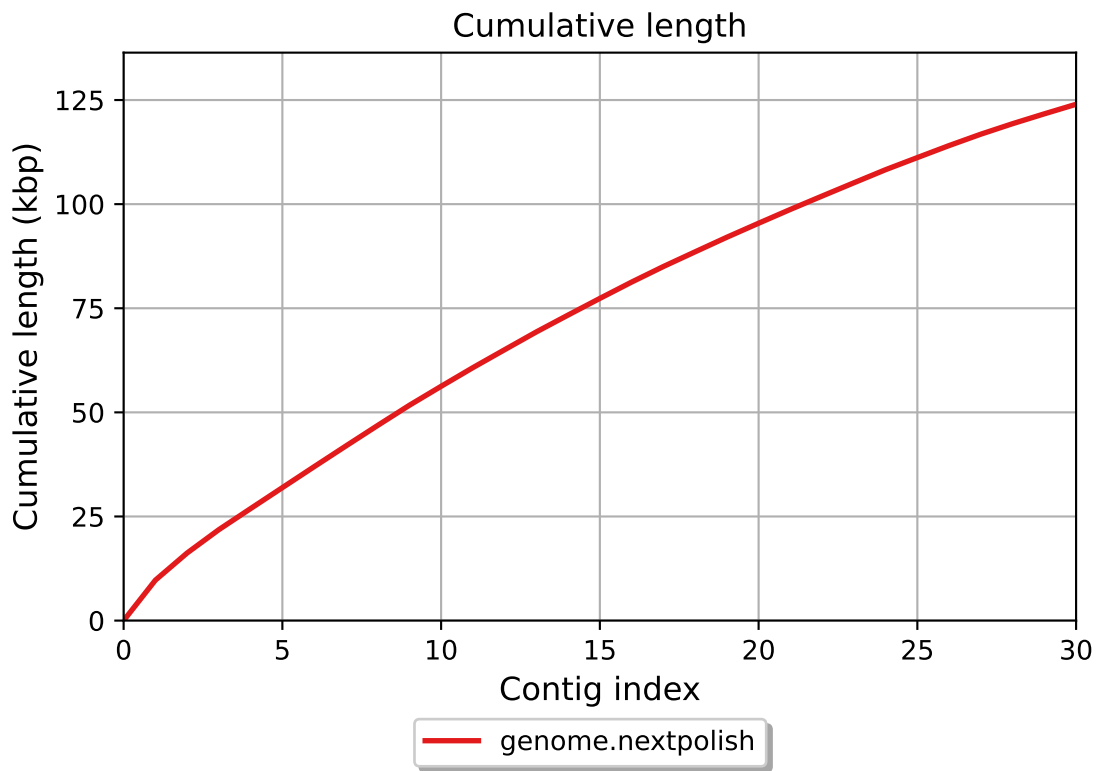
	genome.nextpolish
# contigs (>= 0 bp)	30
# contigs (>= 1000 bp)	30
# contigs (>= 5000 bp)	6
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	123982
Total length (>= 1000 bp)	123982
Total length (>= 5000 bp)	36949
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	30
Largest contig	9704
Total length	123982
GC (%)	34.61
N50	4308
N90	2884
auN	4638.3
L50	12
L90	26
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

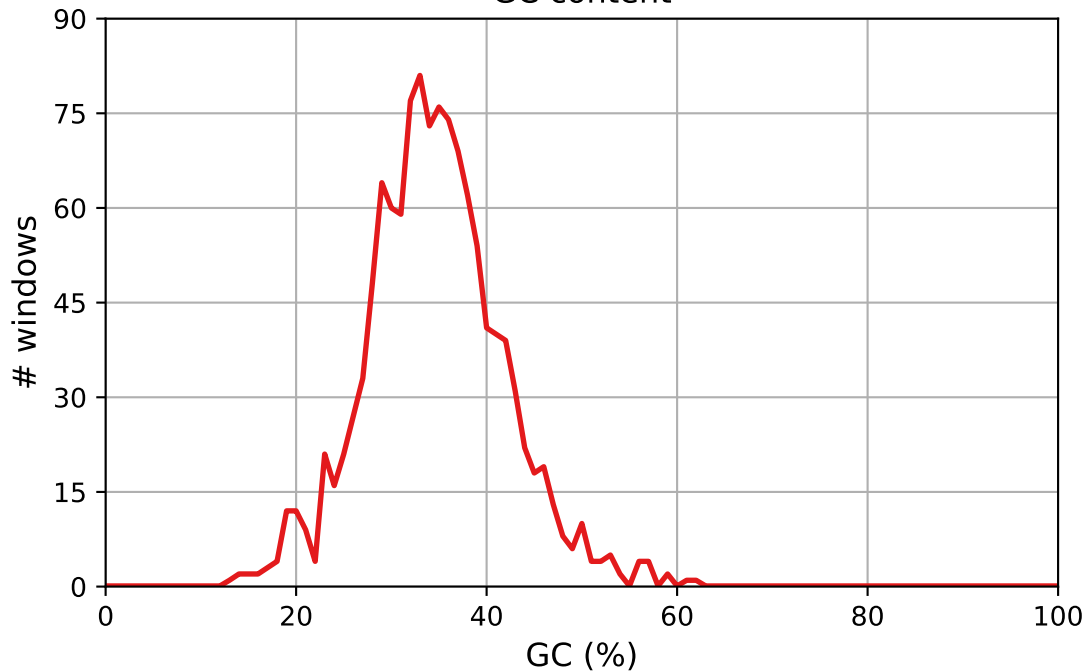
Nx



— genome.nextpolish



GC content



— genome.nextpolish

genome.nextpolish GC content

