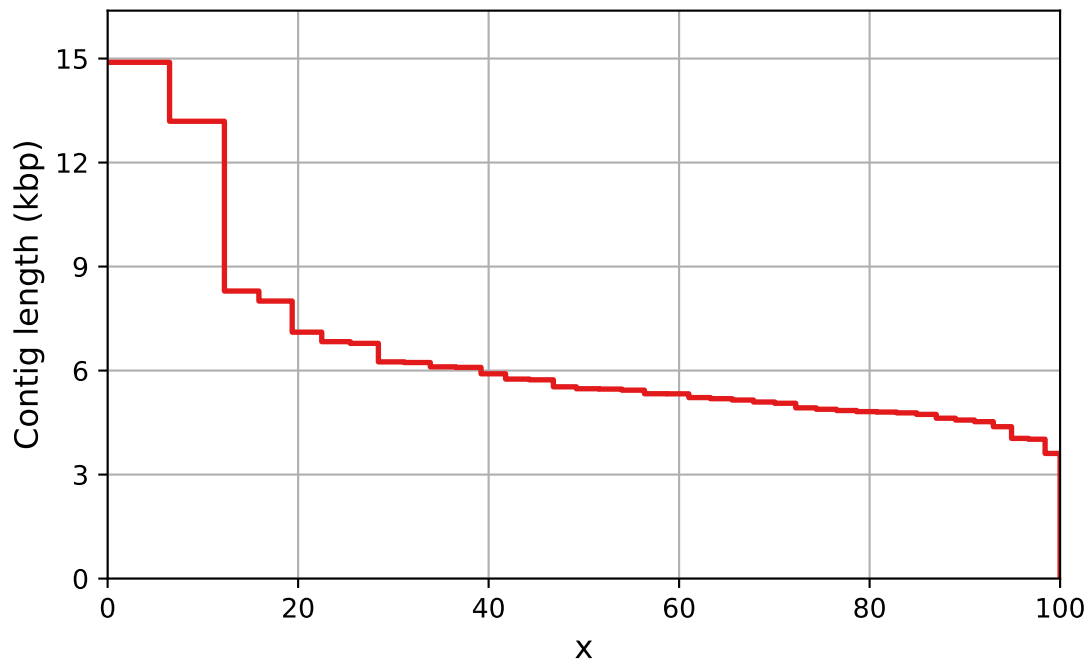


Report

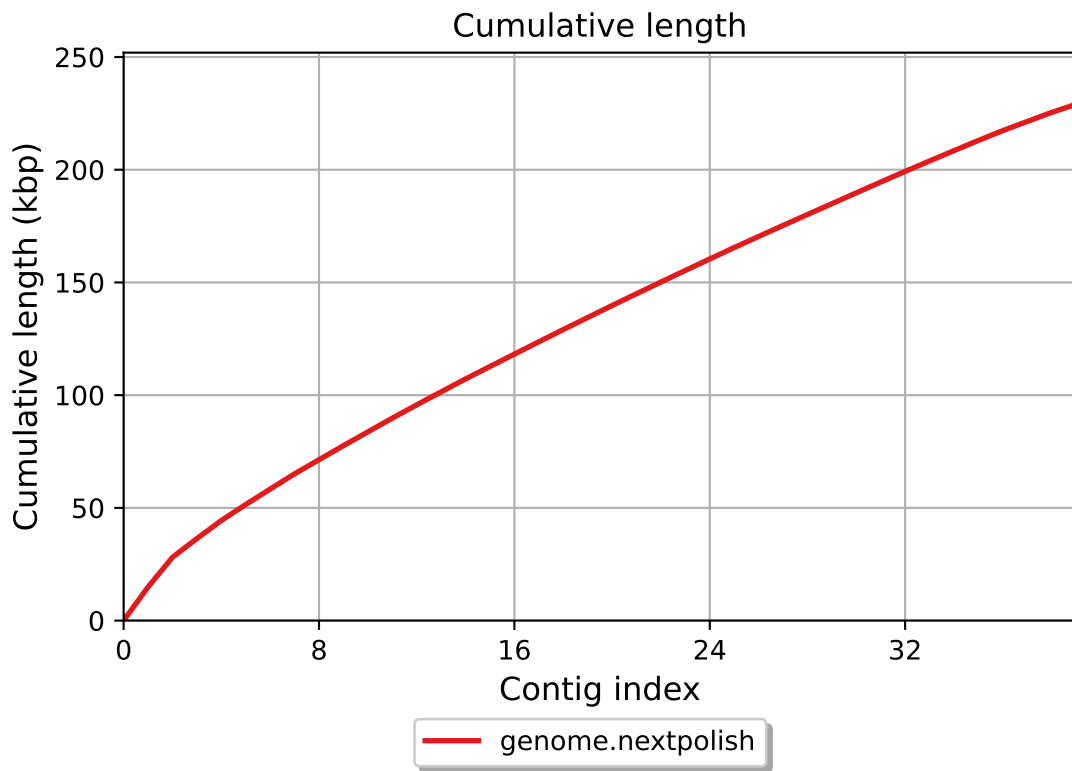
	genome.nextpolish
# contigs (>= 0 bp)	39
# contigs (>= 1000 bp)	39
# contigs (>= 5000 bp)	25
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	229036
Total length (>= 1000 bp)	229036
Total length (>= 5000 bp)	165464
Total length (>= 10000 bp)	28083
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	39
Largest contig	14892
Total length	229036
GC (%)	54.16
N50	5476
N90	4573
auN	6657.6
L50	16
L90	34
# 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).

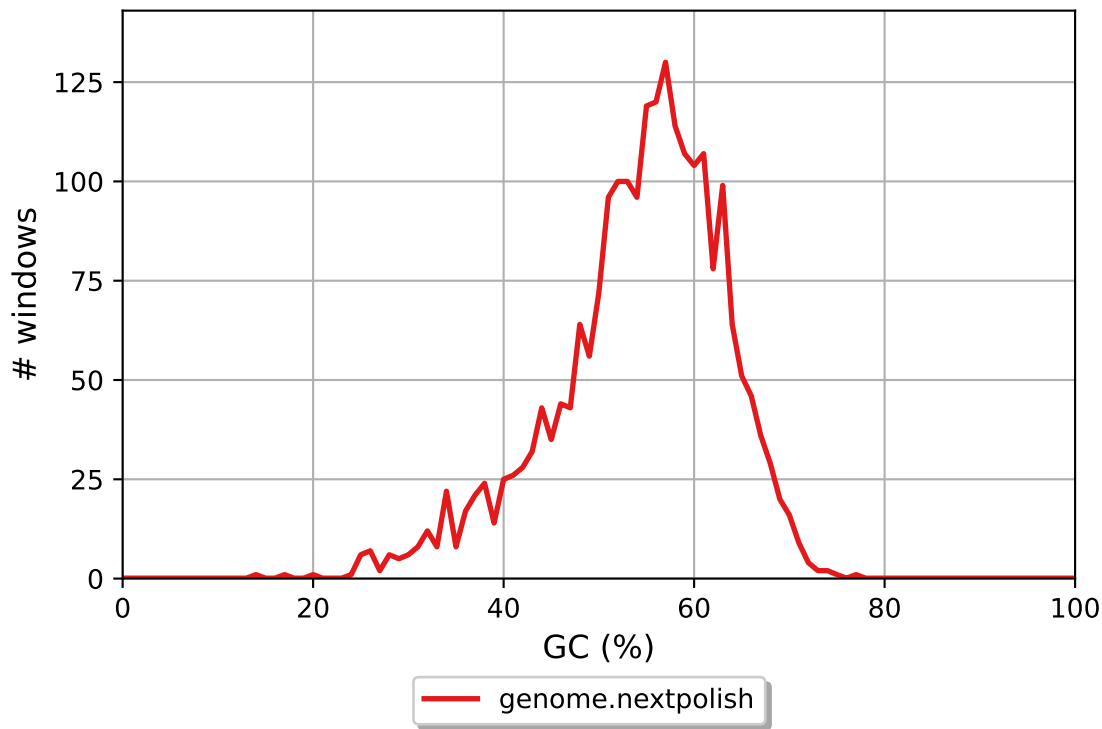
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

