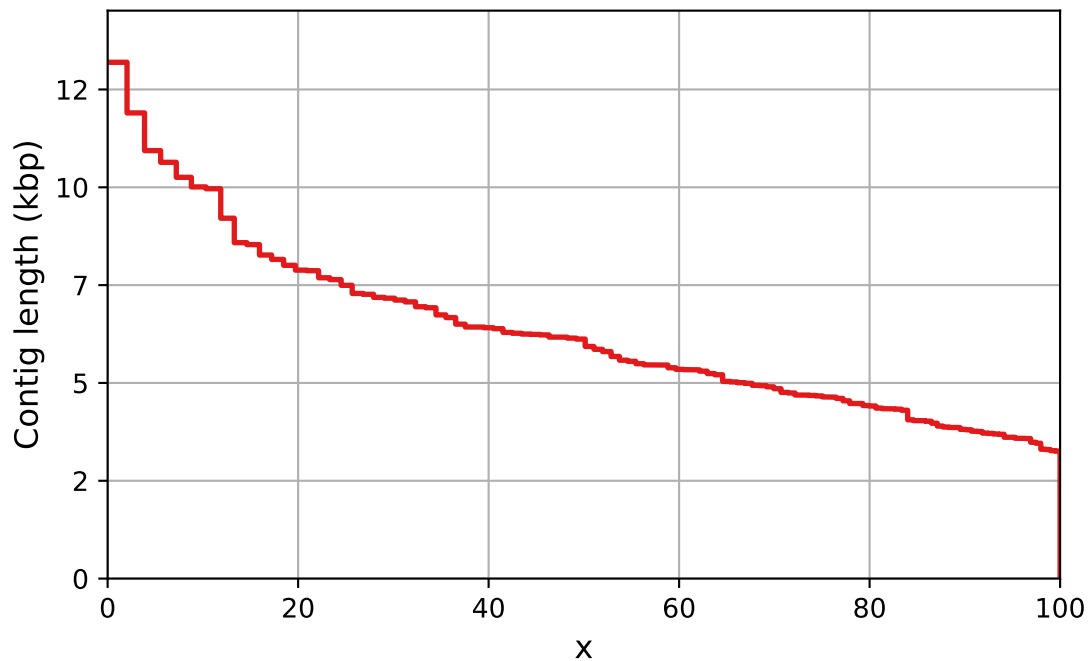


Report

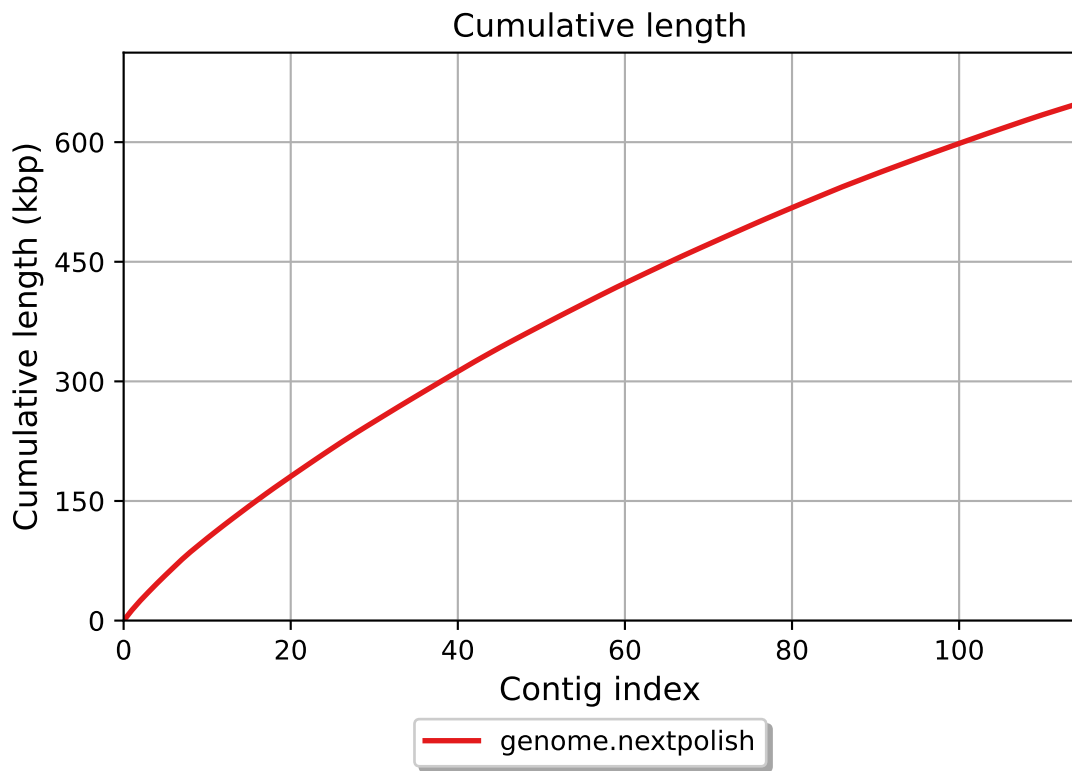
	genome.nextpolish
# contigs (>= 0 bp)	114
# contigs (>= 1000 bp)	114
# contigs (>= 5000 bp)	62
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	647517
Total length (>= 1000 bp)	647517
Total length (>= 5000 bp)	433170
Total length (>= 10000 bp)	66938
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	114
Largest contig	13195
Total length	647517
GC (%)	50.91
N50	6121
N90	3814
auN	6359.4
L50	42
L90	96
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 2 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).

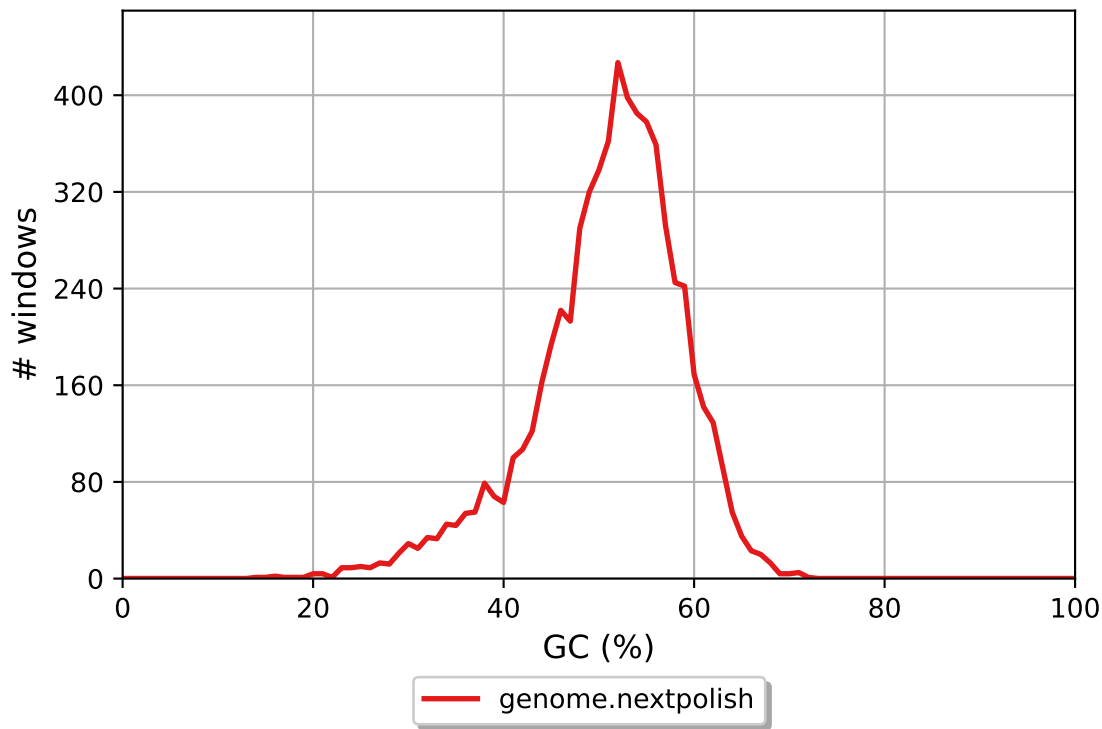
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

