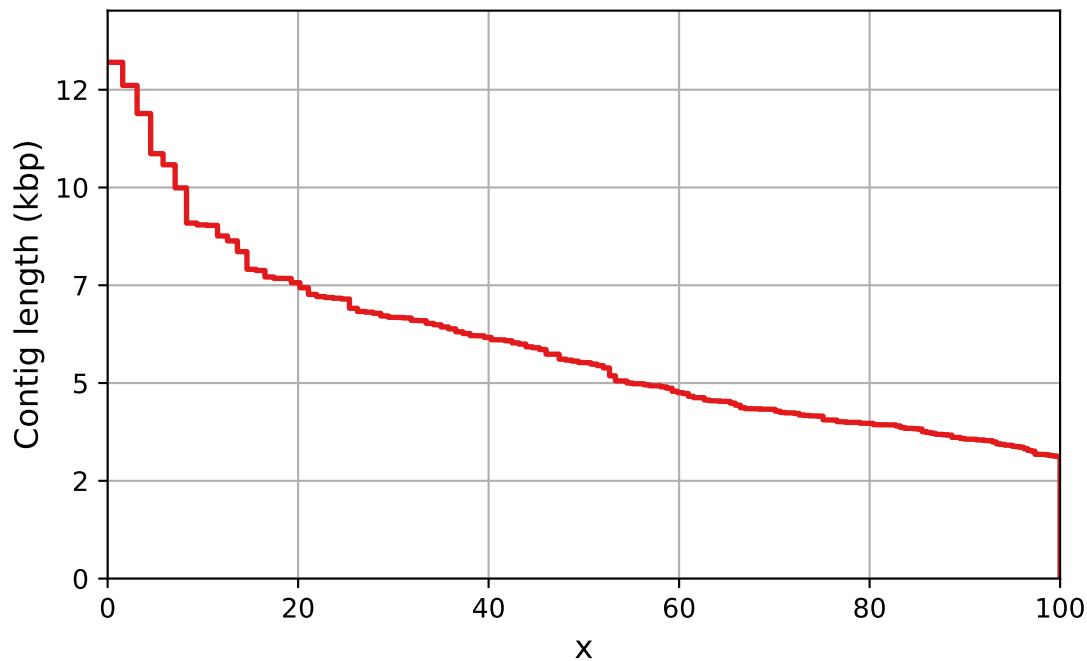


# Report

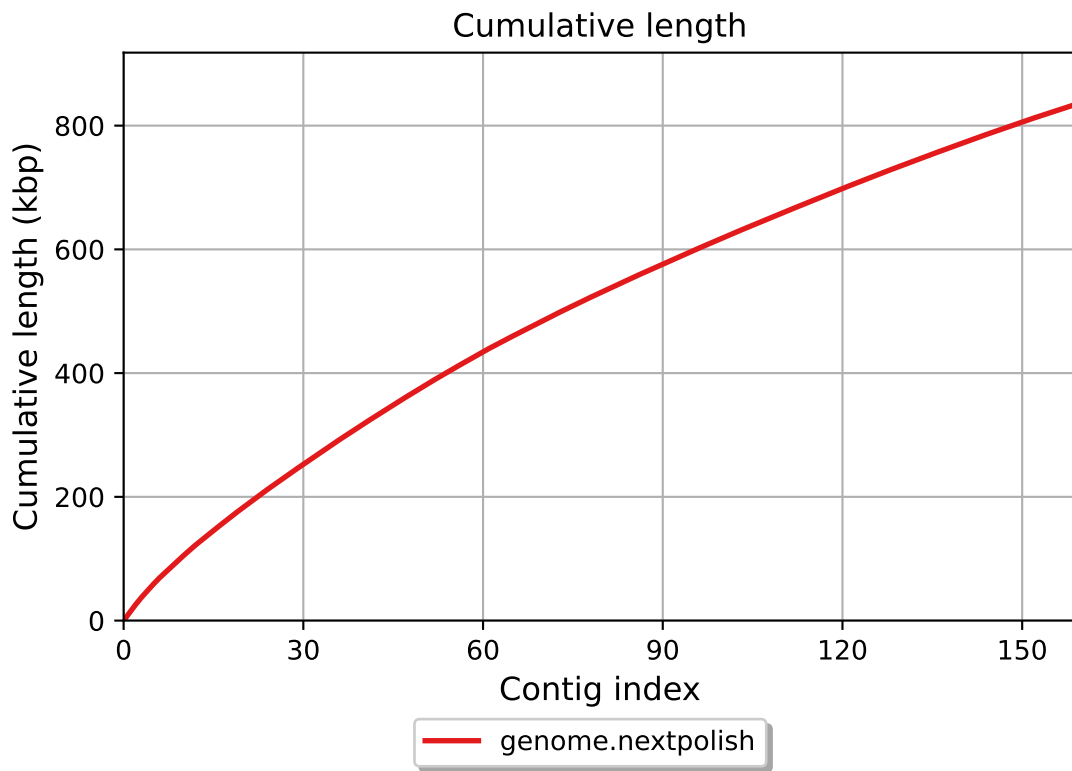
	genome.nextpolish
# contigs (>= 0 bp)	159
# contigs (>= 1000 bp)	159
# contigs (>= 5000 bp)	65
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	834532
Total length (>= 1000 bp)	834532
Total length (>= 5000 bp)	460040
Total length (>= 10000 bp)	59147
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	159
Largest contig	13201
Total length	834532
GC (%)	51.48
N50	5526
N90	3567
auN	5961.5
L50	57
L90	135
# 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).

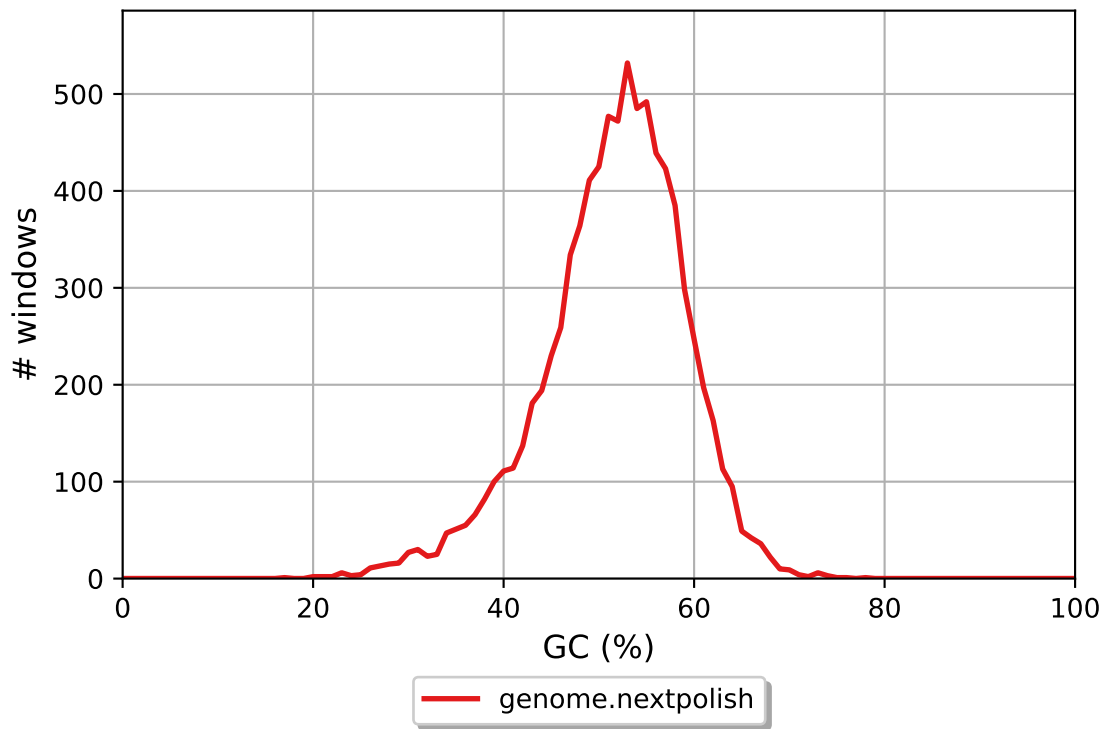
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

