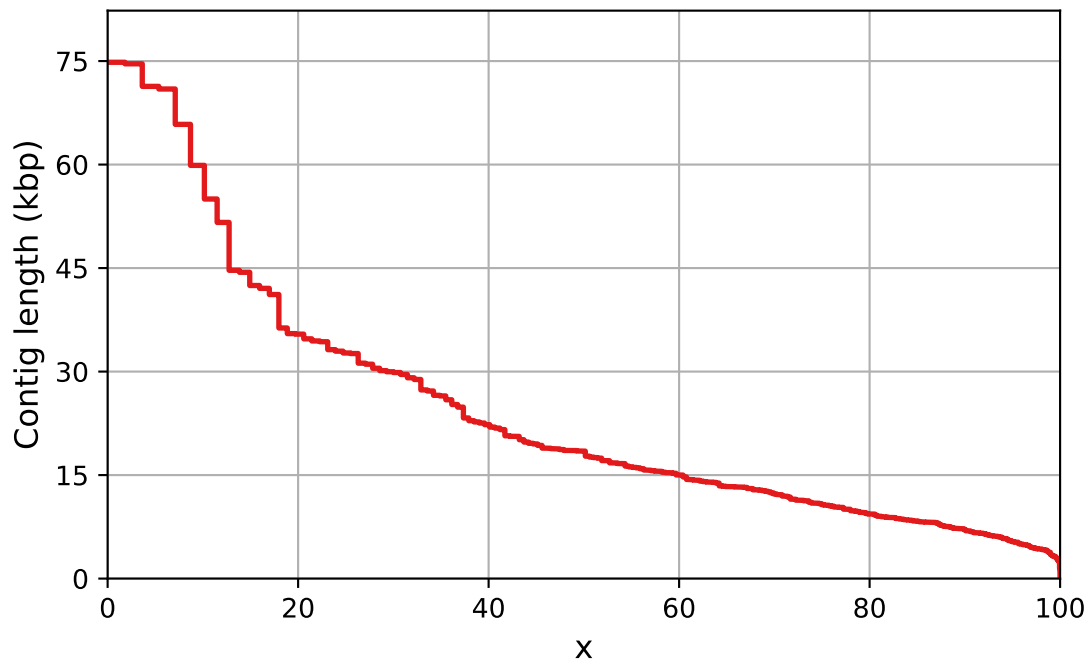


Report

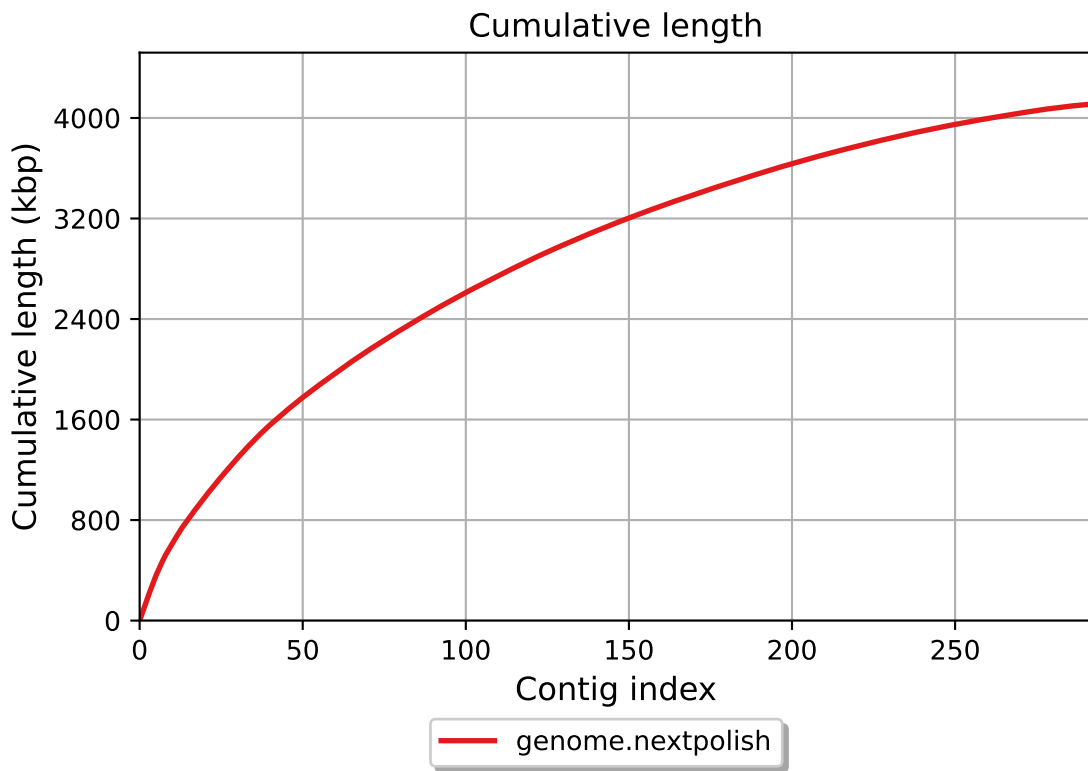
	genome.nextpolish
# contigs (>= 0 bp)	292
# contigs (>= 1000 bp)	291
# contigs (>= 5000 bp)	249
# contigs (>= 10000 bp)	150
# contigs (>= 25000 bp)	38
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	4109100
Total length (>= 1000 bp)	4108238
Total length (>= 5000 bp)	3943917
Total length (>= 10000 bp)	3204354
Total length (>= 25000 bp)	1509985
Total length (>= 50000 bp)	524042
# contigs	292
Largest contig	74825
Total length	4109100
GC (%)	48.93
N50	18469
N90	7097
auN	24745.4
L50	65
L90	209
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 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).

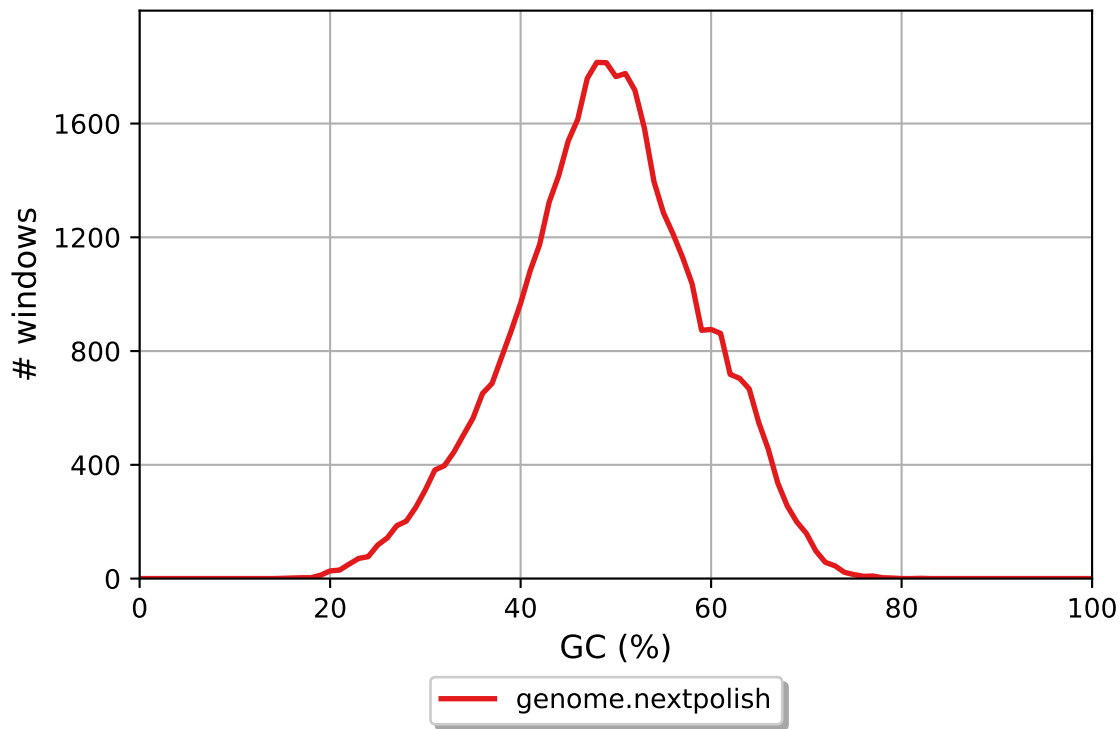
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

