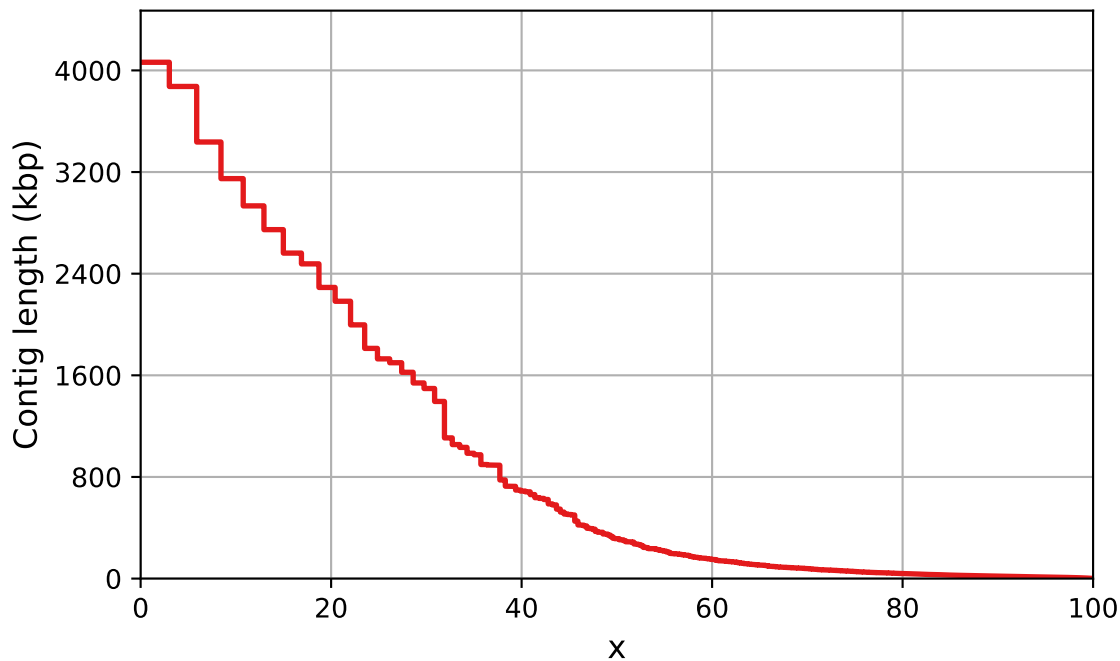


Report

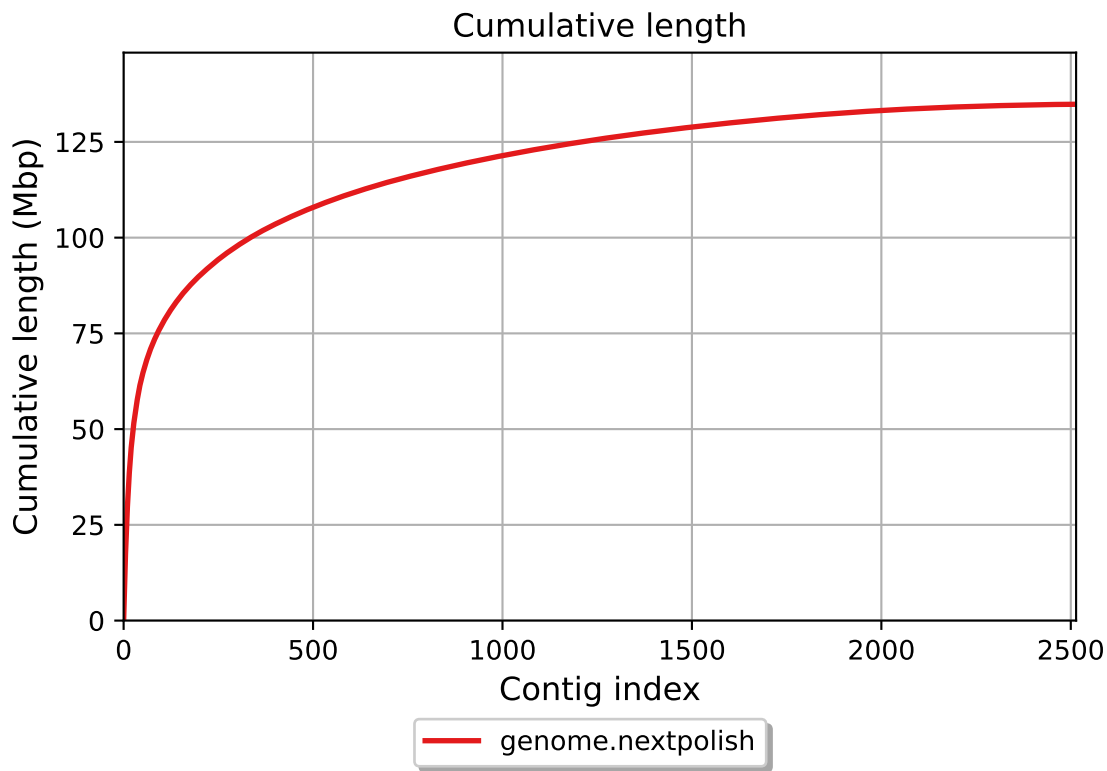
	genome.nextpolish
# contigs (>= 0 bp)	2517
# contigs (>= 1000 bp)	2482
# contigs (>= 5000 bp)	2059
# contigs (>= 10000 bp)	1635
# contigs (>= 25000 bp)	773
# contigs (>= 50000 bp)	388
Total length (>= 0 bp)	134828223
Total length (>= 1000 bp)	134804004
Total length (>= 5000 bp)	133527374
Total length (>= 10000 bp)	130315304
Total length (>= 25000 bp)	116453473
Total length (>= 50000 bp)	102924357
# contigs	2514
Largest contig	4064014
Total length	134826836
GC (%)	50.07
N50	314248
N90	19223
auN	1015601.7
L50	59
L90	997
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	255 + 8 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).

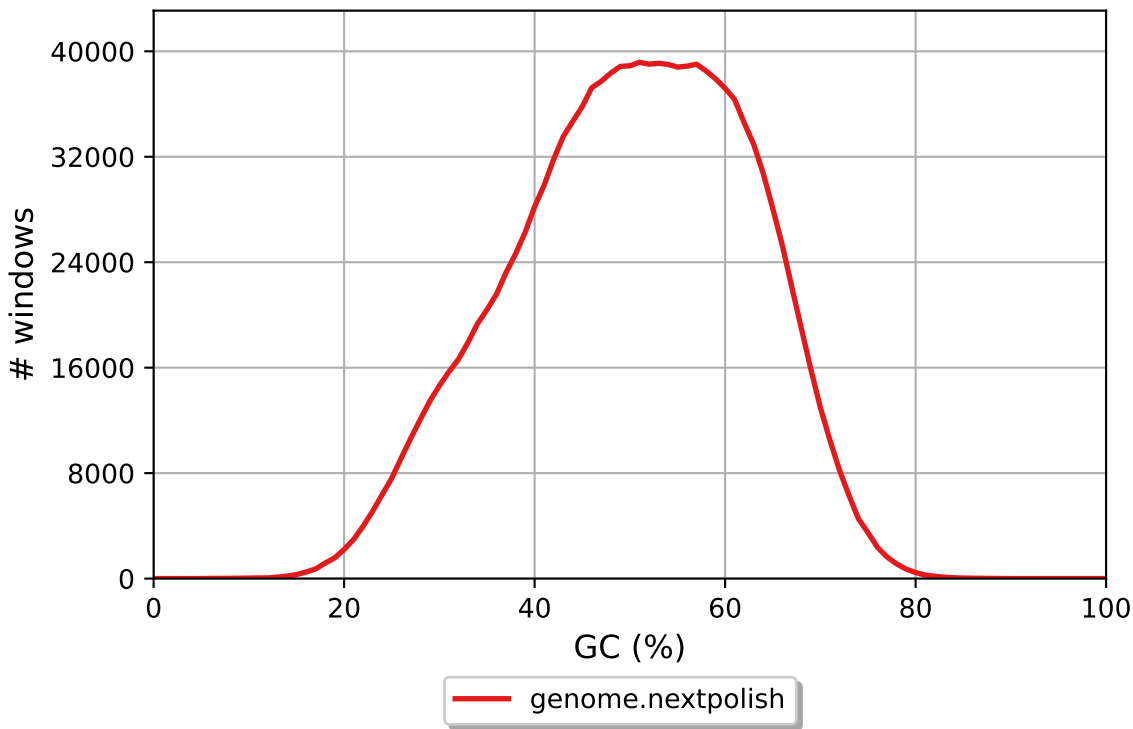
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

