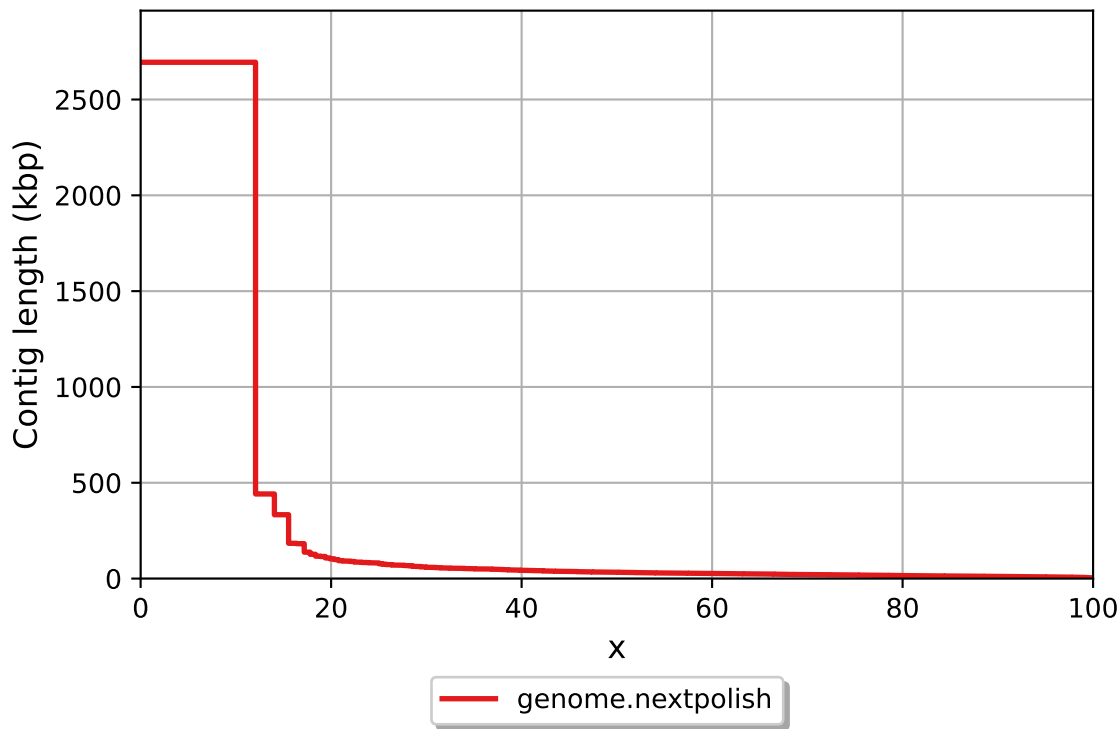


Report

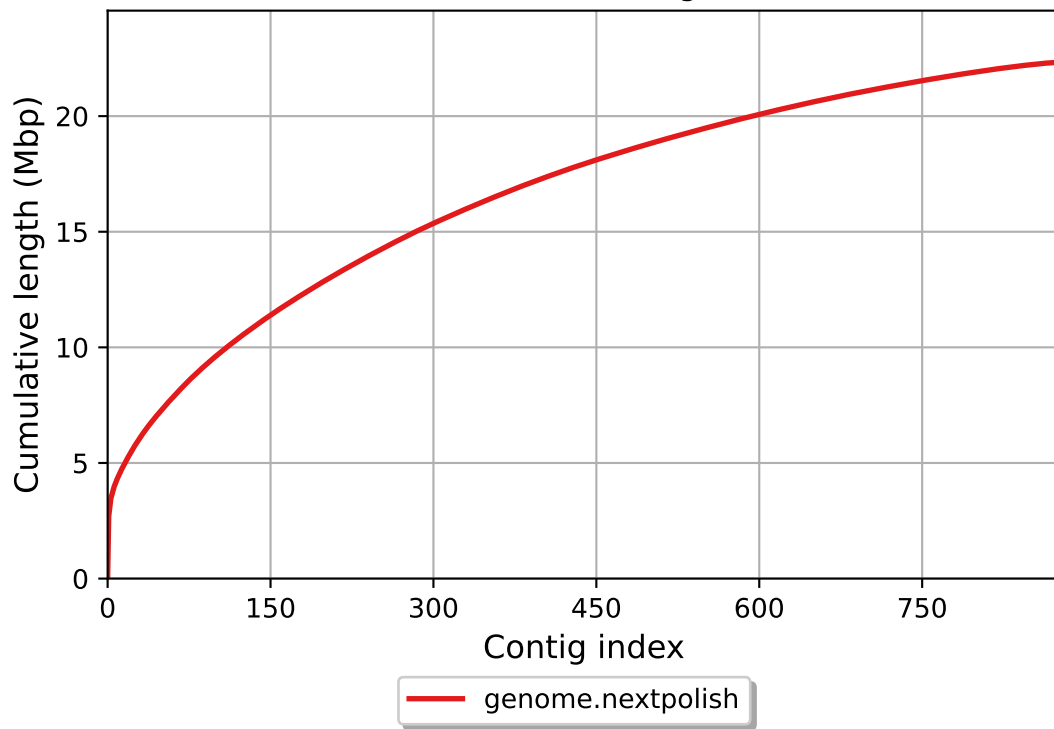
	genome.nextpolish
# contigs (>= 0 bp)	877
# contigs (>= 1000 bp)	874
# contigs (>= 5000 bp)	852
# contigs (>= 10000 bp)	661
# contigs (>= 25000 bp)	245
# contigs (>= 50000 bp)	64
Total length (>= 0 bp)	22329175
Total length (>= 1000 bp)	22327533
Total length (>= 5000 bp)	22236504
Total length (>= 10000 bp)	20730196
Total length (>= 25000 bp)	14072442
Total length (>= 50000 bp)	8038854
# contigs	877
Largest contig	2694323
Total length	22329175
GC (%)	52.76
N50	33036
N90	11362
auN	371409.3
L50	143
L90	602
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	77 + 5 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).

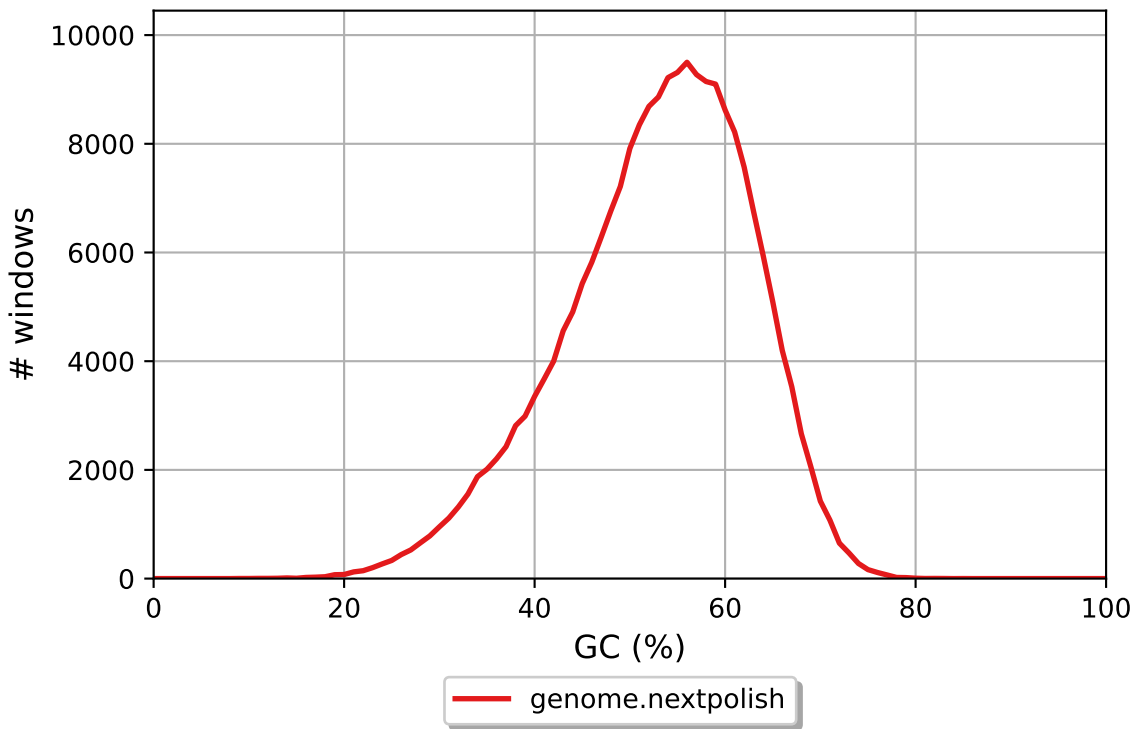
Nx



Cumulative length



GC content



genome.nextpolish GC content

