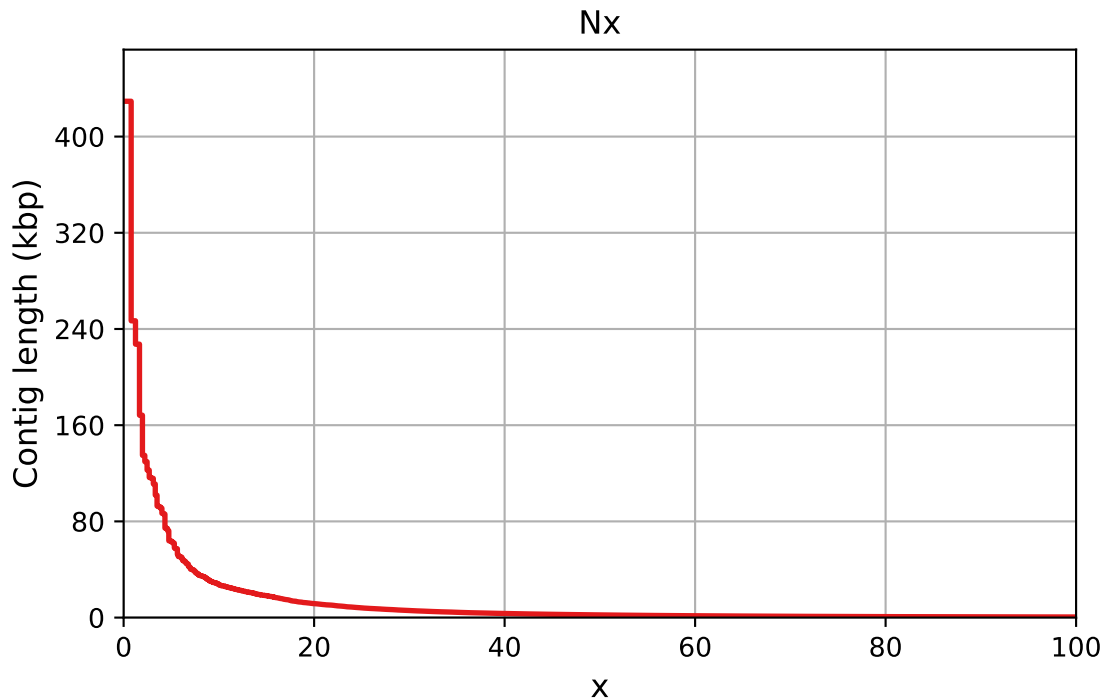


Report

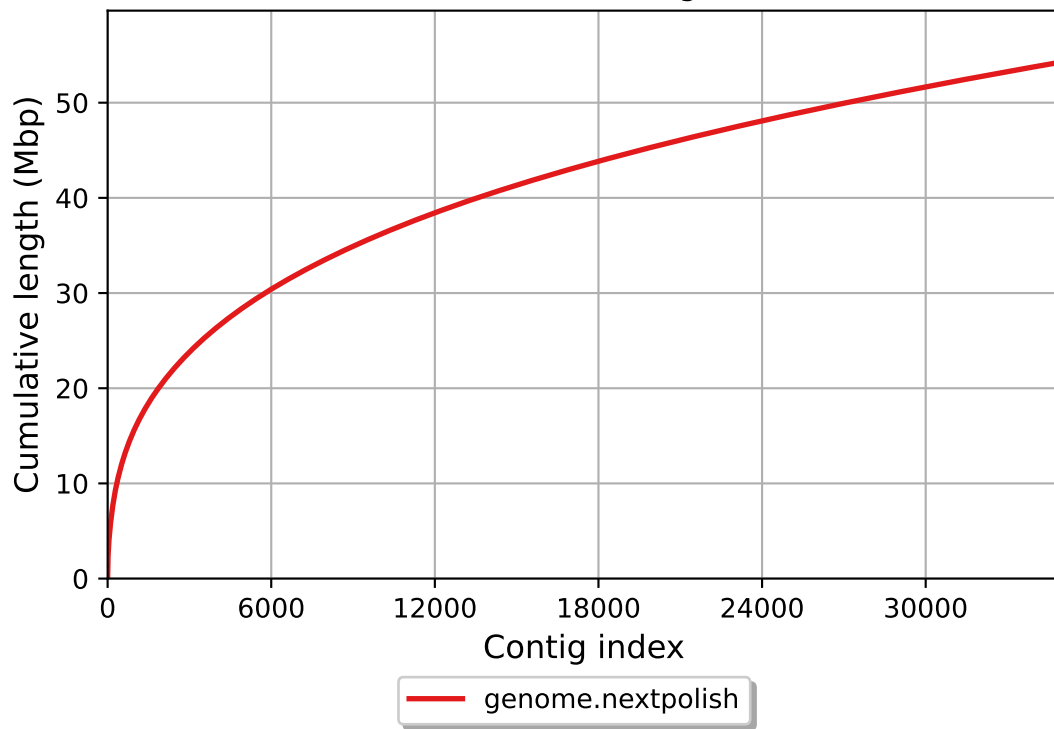
	genome.nextpolish
# contigs (>= 0 bp)	165779
# contigs (>= 1000 bp)	12902
# contigs (>= 5000 bp)	1350
# contigs (>= 10000 bp)	515
# contigs (>= 25000 bp)	113
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	91959252
Total length (>= 1000 bp)	39361601
Total length (>= 5000 bp)	17726368
Total length (>= 10000 bp)	12036325
Total length (>= 25000 bp)	5981023
Total length (>= 50000 bp)	3316219
# contigs	34929
Largest contig	429365
Total length	54247596
GC (%)	52.23
N50	2213
N90	622
auN	15009.1
L50	4326
L90	25155
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 14 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).

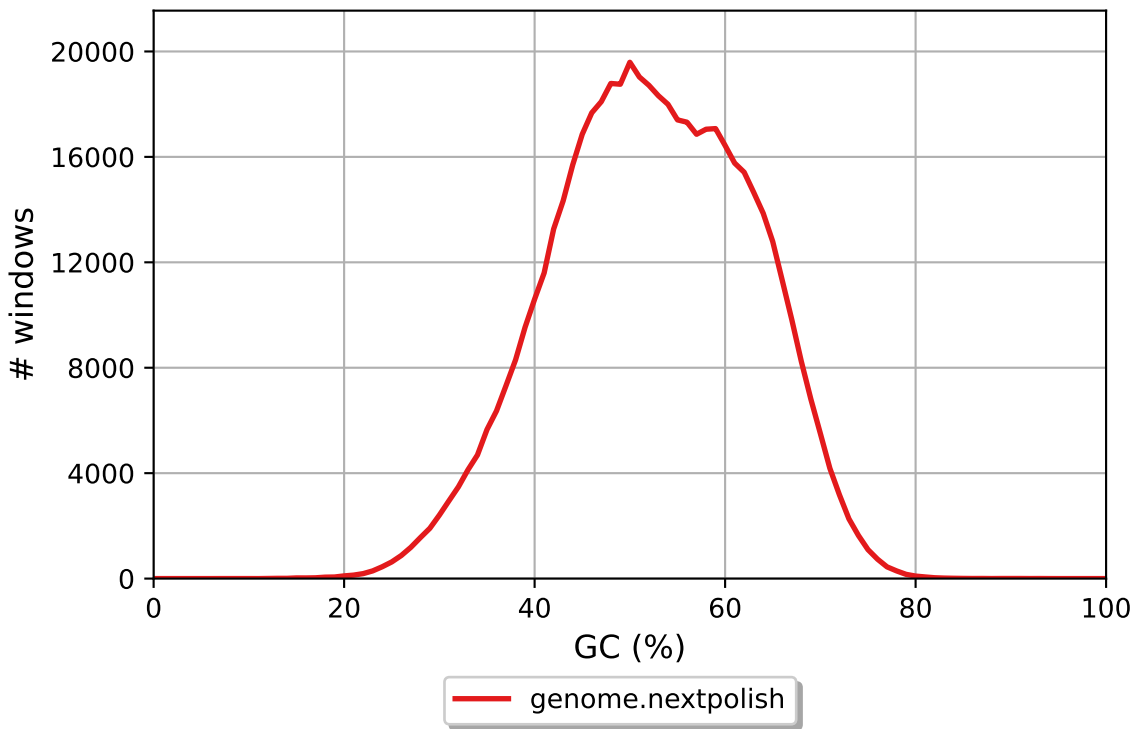


— genome.nextpolish

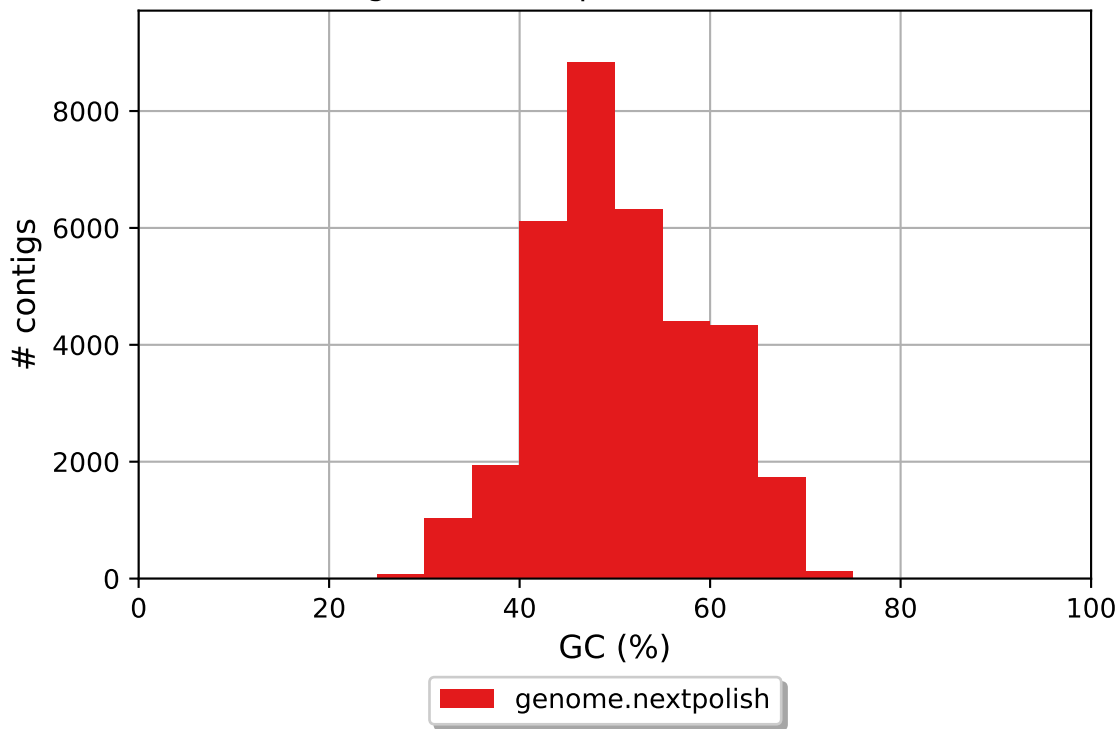
Cumulative length



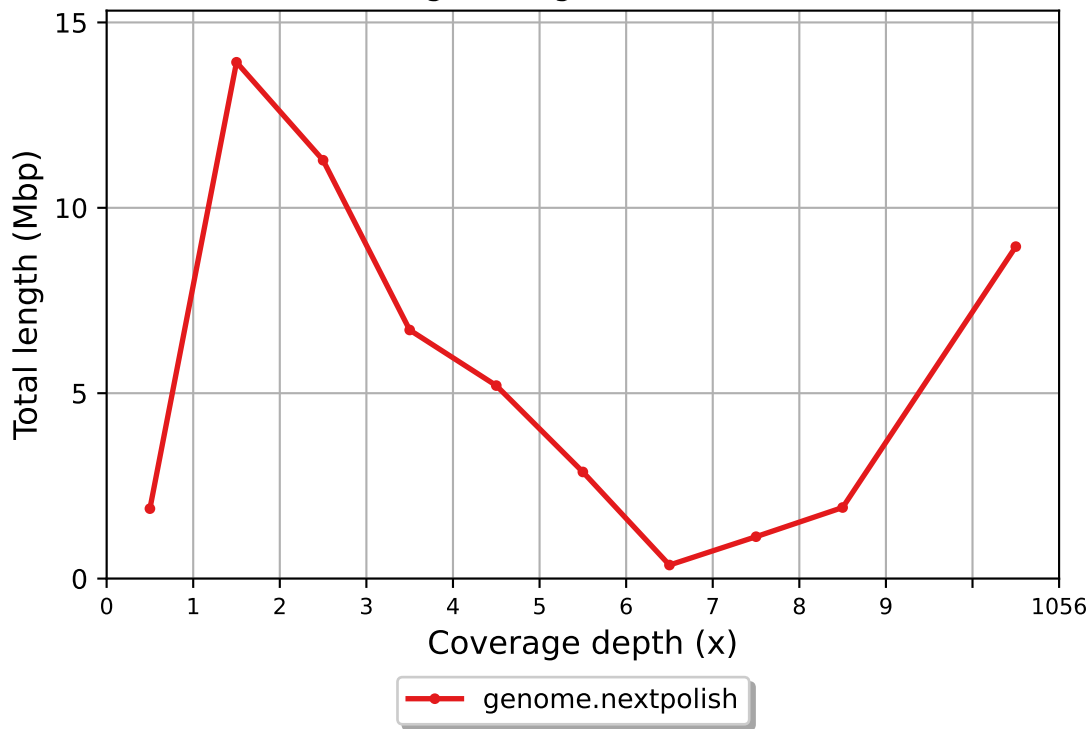
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



genome.nextpolish coverage histogram (bin size: 1x)

