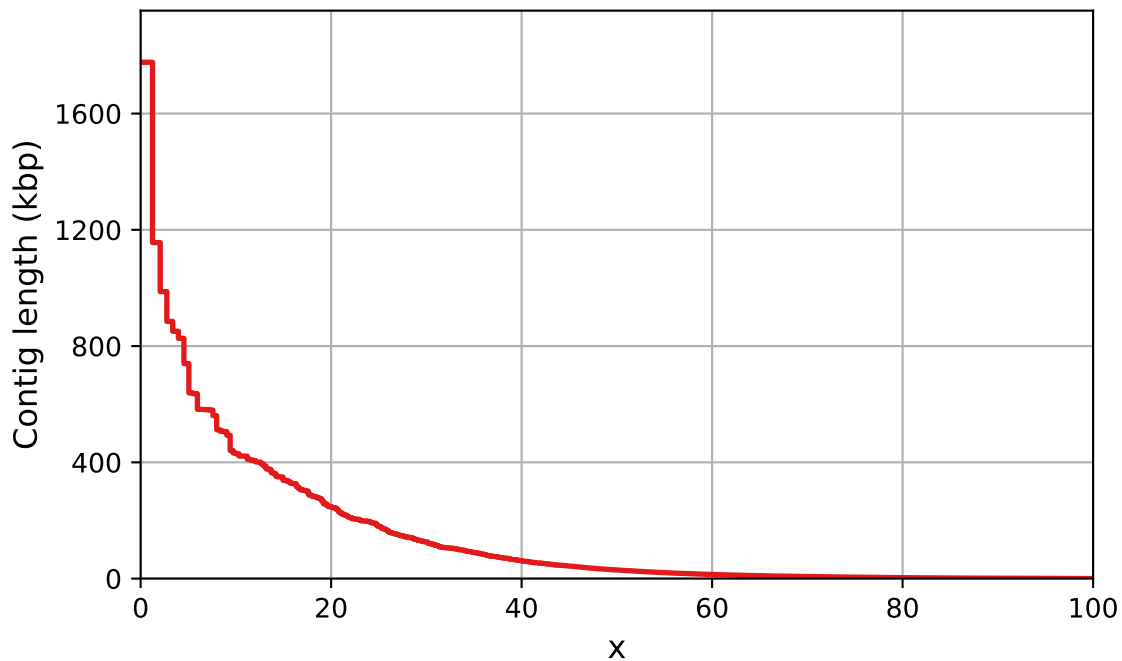


Report

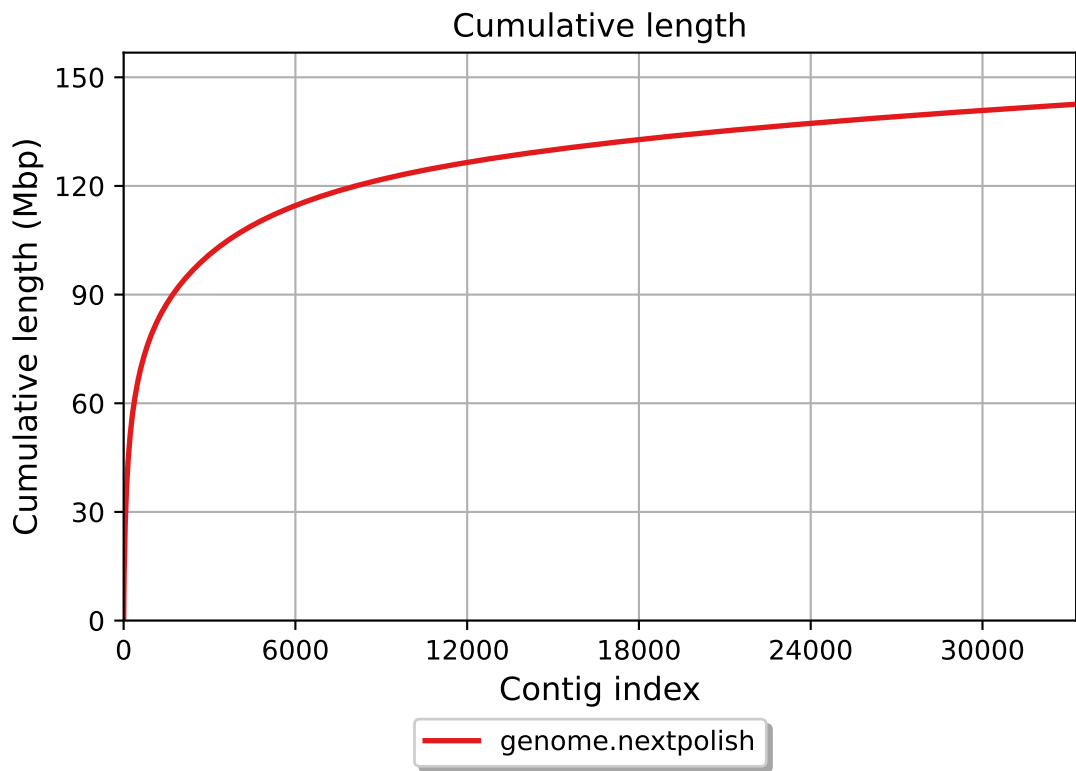
	genome.nextpolish
# contigs (>= 0 bp)	149311
# contigs (>= 1000 bp)	15480
# contigs (>= 5000 bp)	3938
# contigs (>= 10000 bp)	1957
# contigs (>= 25000 bp)	774
# contigs (>= 50000 bp)	381
Total length (>= 0 bp)	175378146
Total length (>= 1000 bp)	130463937
Total length (>= 5000 bp)	106500847
Total length (>= 10000 bp)	92581257
Total length (>= 25000 bp)	74626958
Total length (>= 50000 bp)	60962976
# contigs	33270
Largest contig	1776463
Total length	142549652
GC (%)	45.16
N50	29714
N90	1155
auN	154342.8
L50	651
L90	13458
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	81 + 26 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).

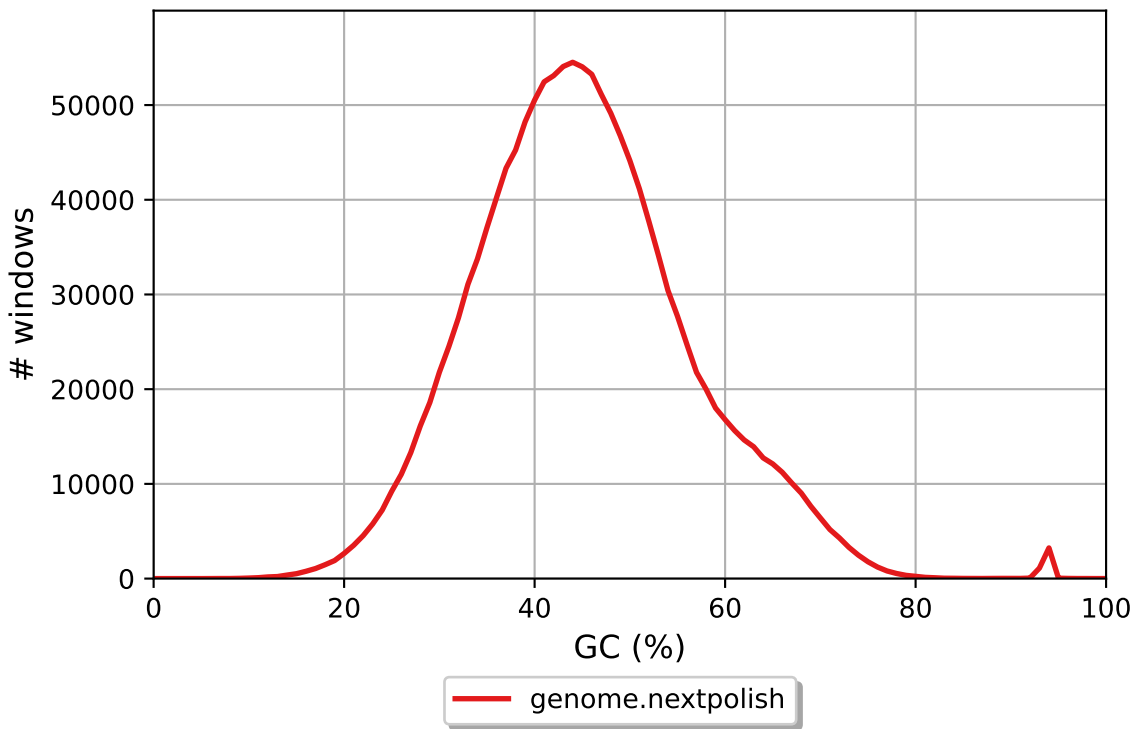
Nx



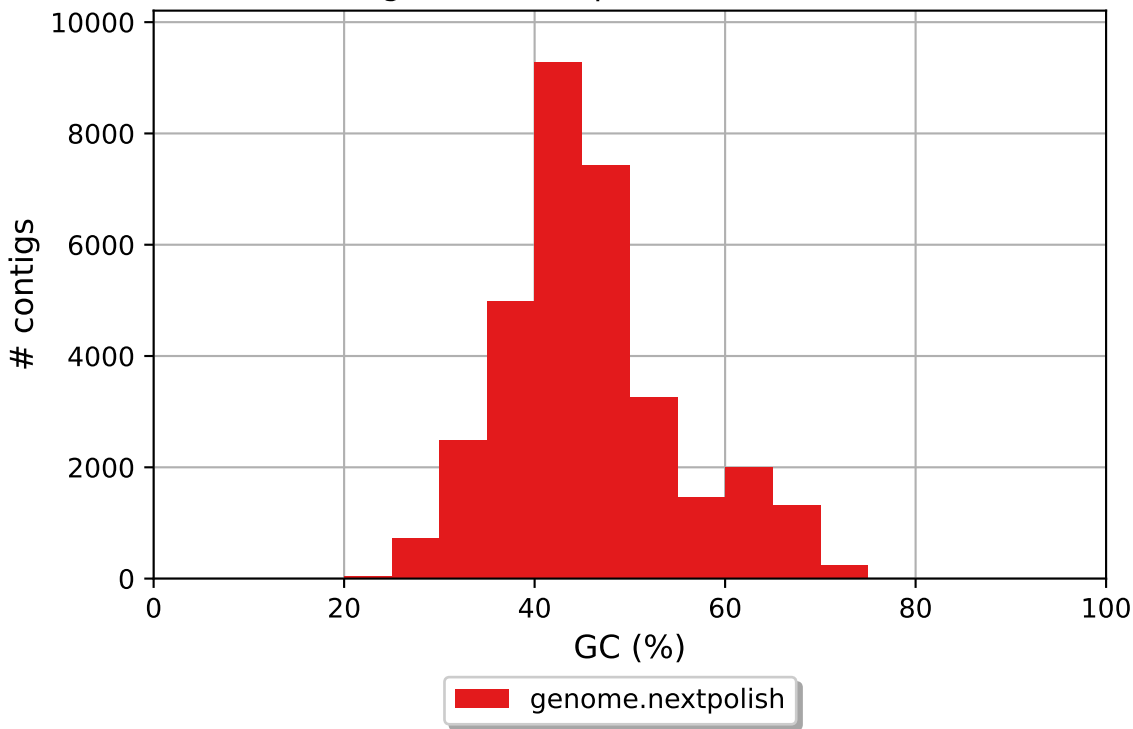
— genome.nextpolish



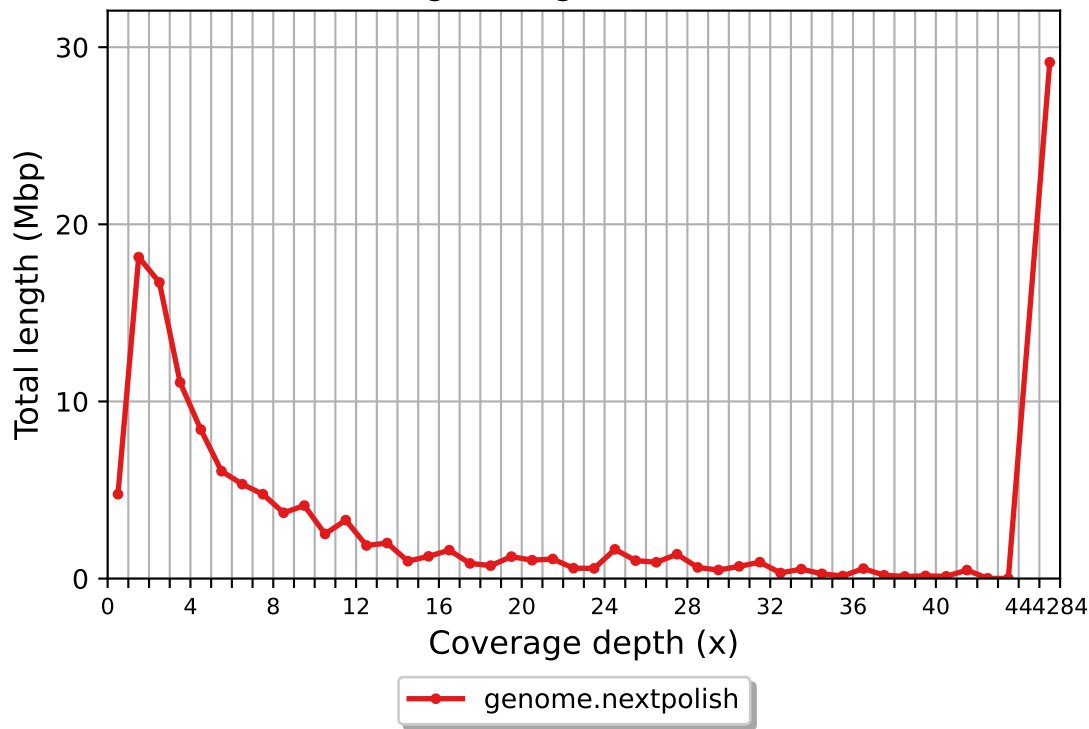
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

