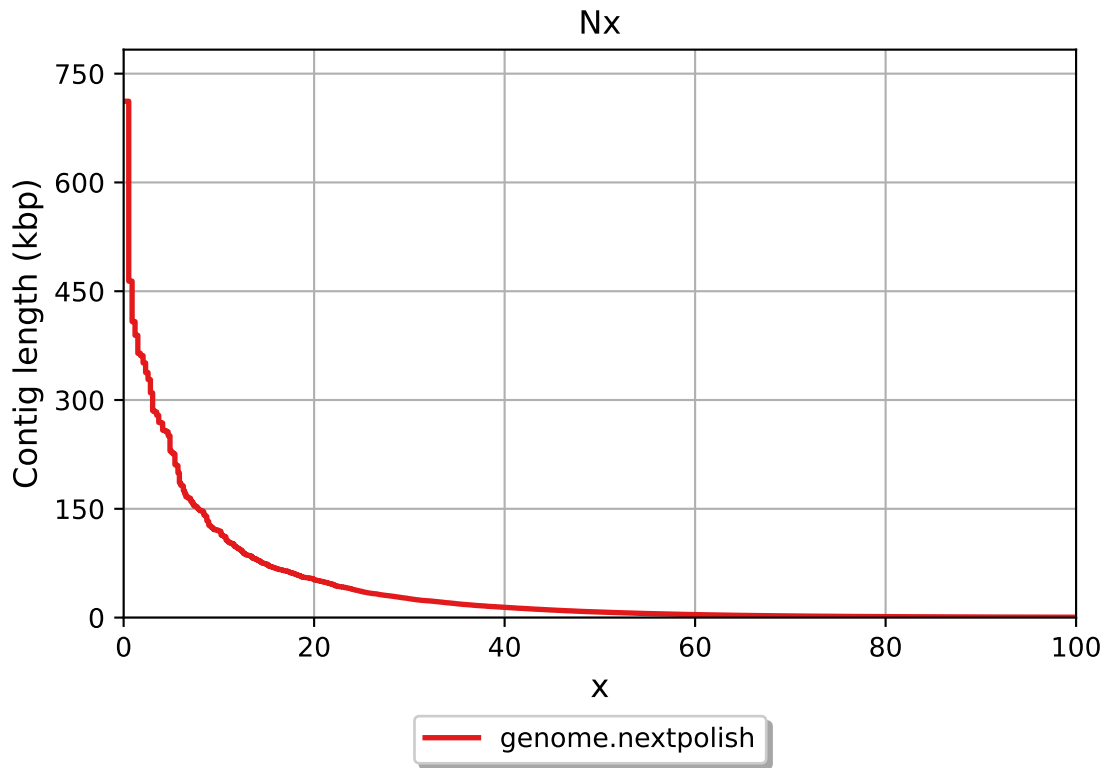


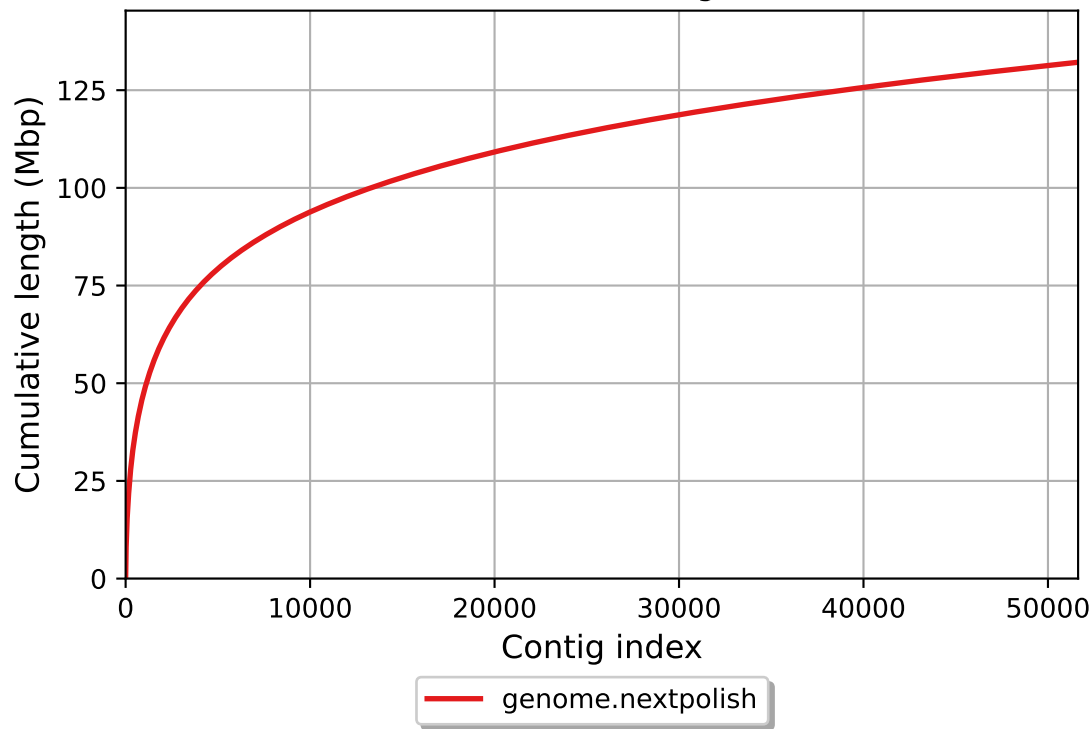
Report

	genome.nextpolish
# contigs (>= 0 bp)	188886
# contigs (>= 1000 bp)	23343
# contigs (>= 5000 bp)	4114
# contigs (>= 10000 bp)	1944
# contigs (>= 25000 bp)	637
# contigs (>= 50000 bp)	262
Total length (>= 0 bp)	171712530
Total length (>= 1000 bp)	112775514
Total length (>= 5000 bp)	75301960
Total length (>= 10000 bp)	60327885
Total length (>= 25000 bp)	40306638
Total length (>= 50000 bp)	27427070
# contigs	51634
Largest contig	711919
Total length	132145064
GC (%)	47.17
N50	7651
N90	792
auN	42098.1
L50	2605
L90	30280
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	37 + 21 part

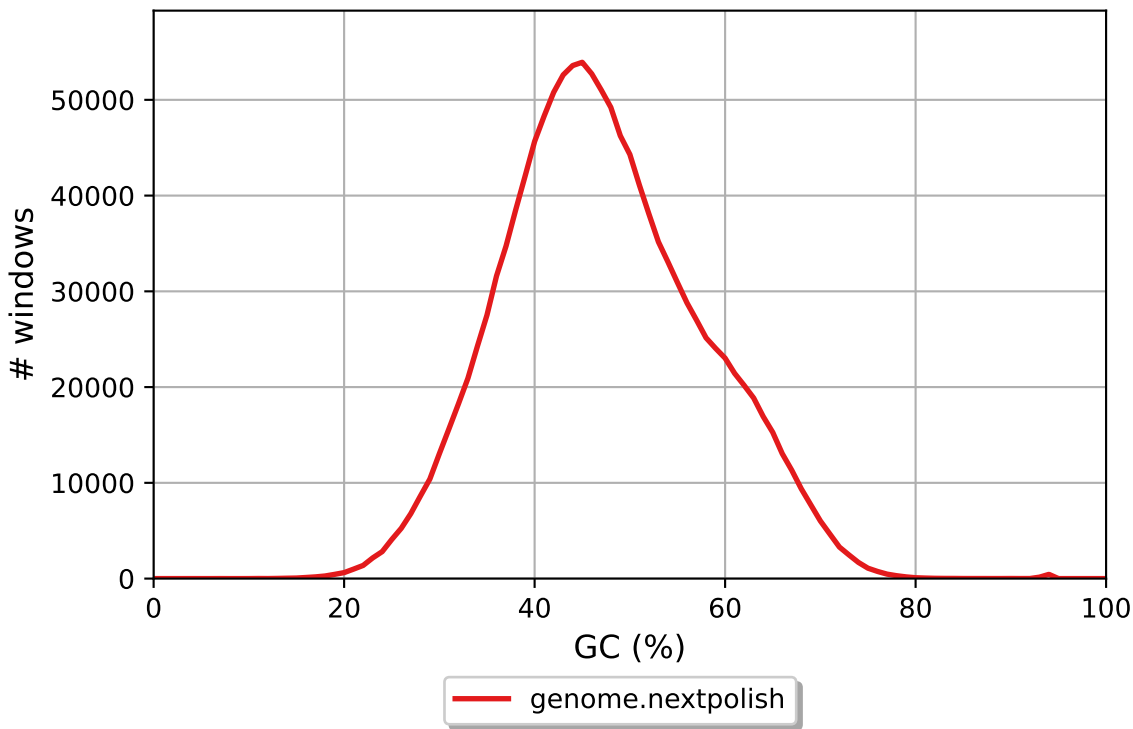
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



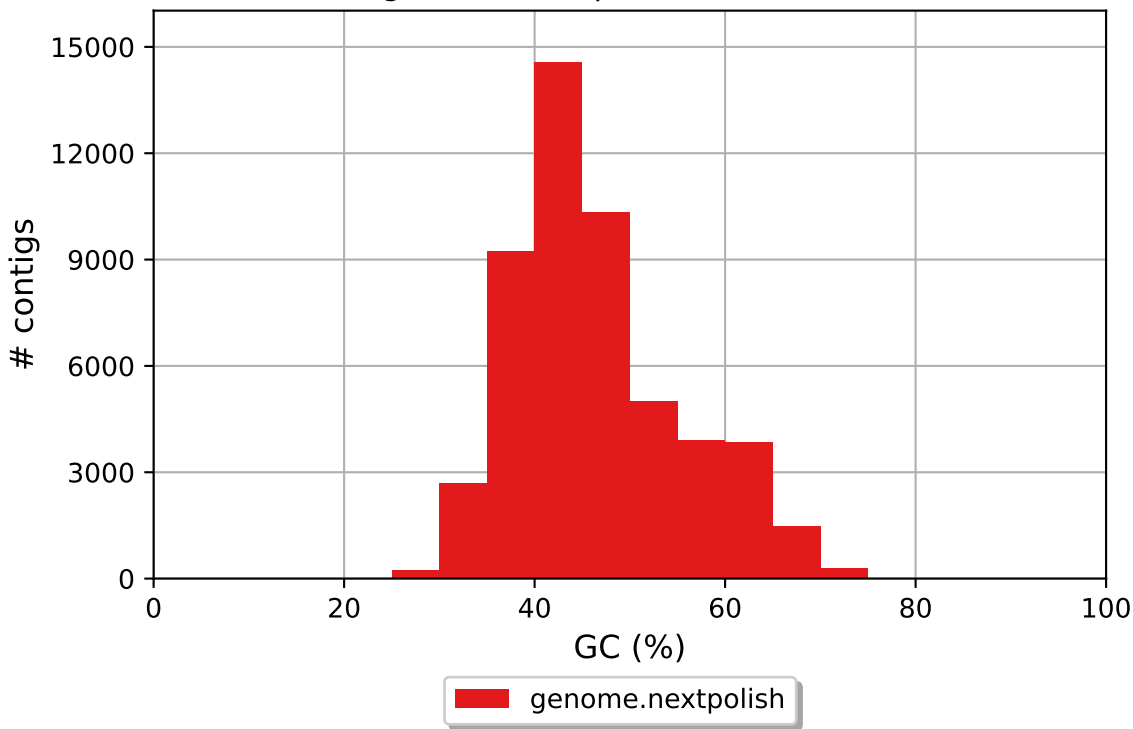
Cumulative length



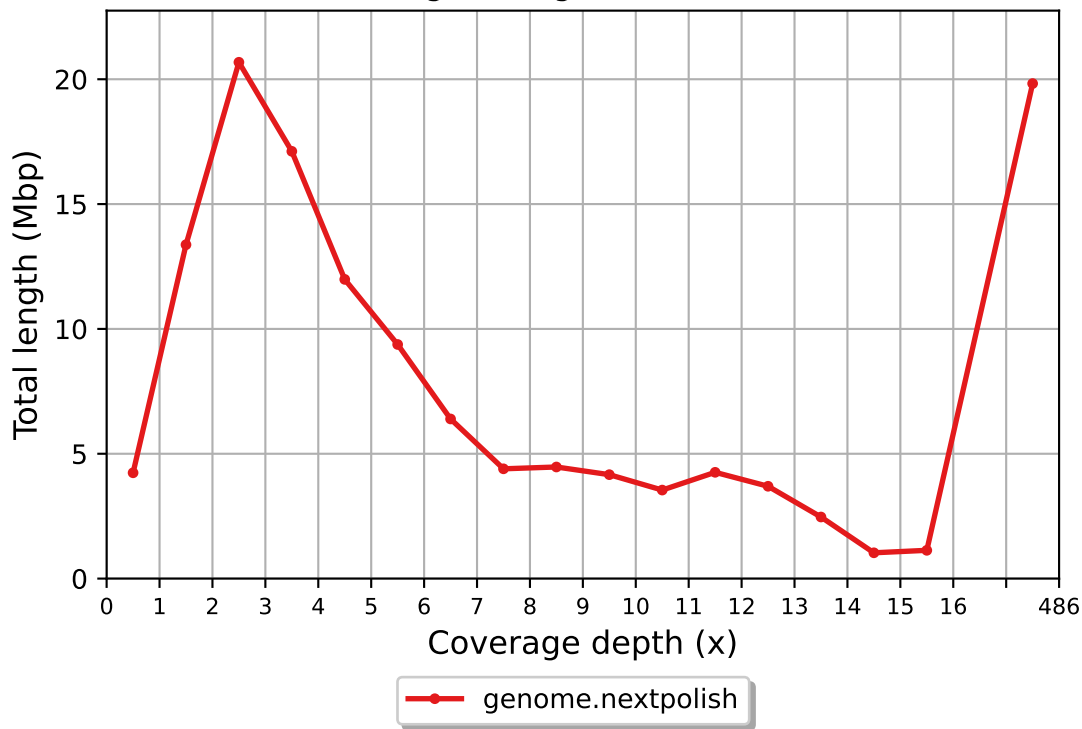
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

