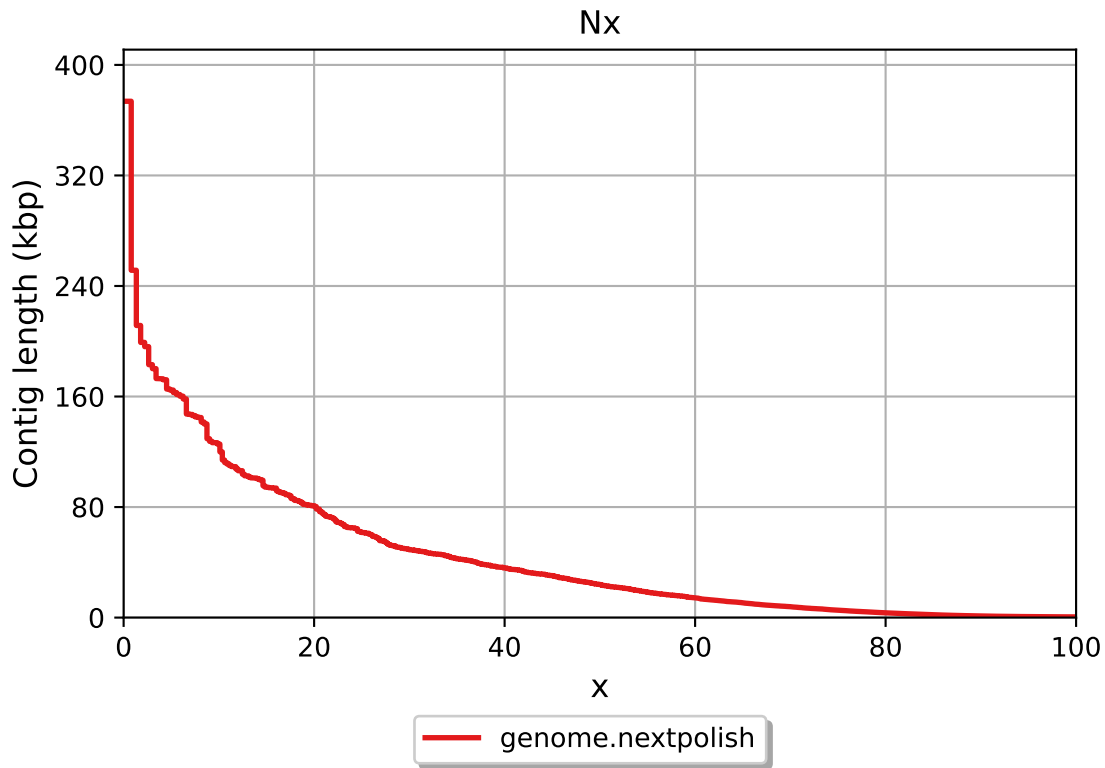


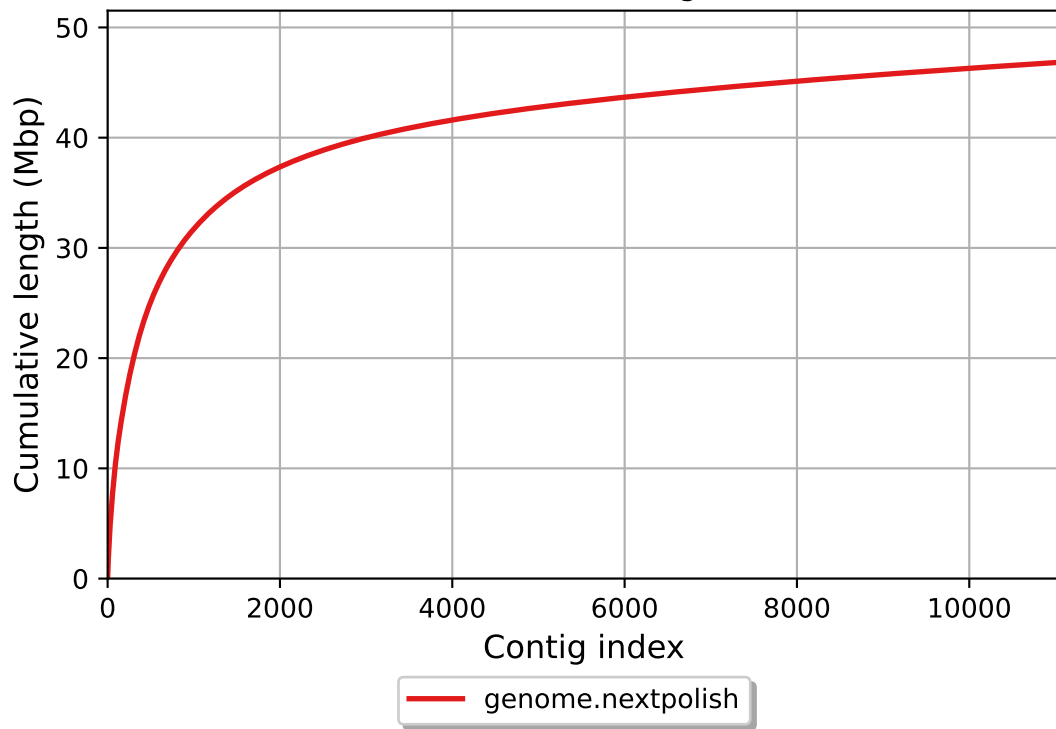
Report

	genome.nextpolish
# contigs (>= 0 bp)	47492
# contigs (>= 1000 bp)	5048
# contigs (>= 5000 bp)	1567
# contigs (>= 10000 bp)	915
# contigs (>= 25000 bp)	405
# contigs (>= 50000 bp)	148
Total length (>= 0 bp)	57227321
Total length (>= 1000 bp)	42809081
Total length (>= 5000 bp)	35532263
Total length (>= 10000 bp)	30920340
Total length (>= 25000 bp)	22999943
Total length (>= 50000 bp)	13796043
# contigs	11054
Largest contig	373686
Total length	46840141
GC (%)	45.31
N50	23934
N90	1170
auN	45326.4
L50	423
L90	4443
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 4 part

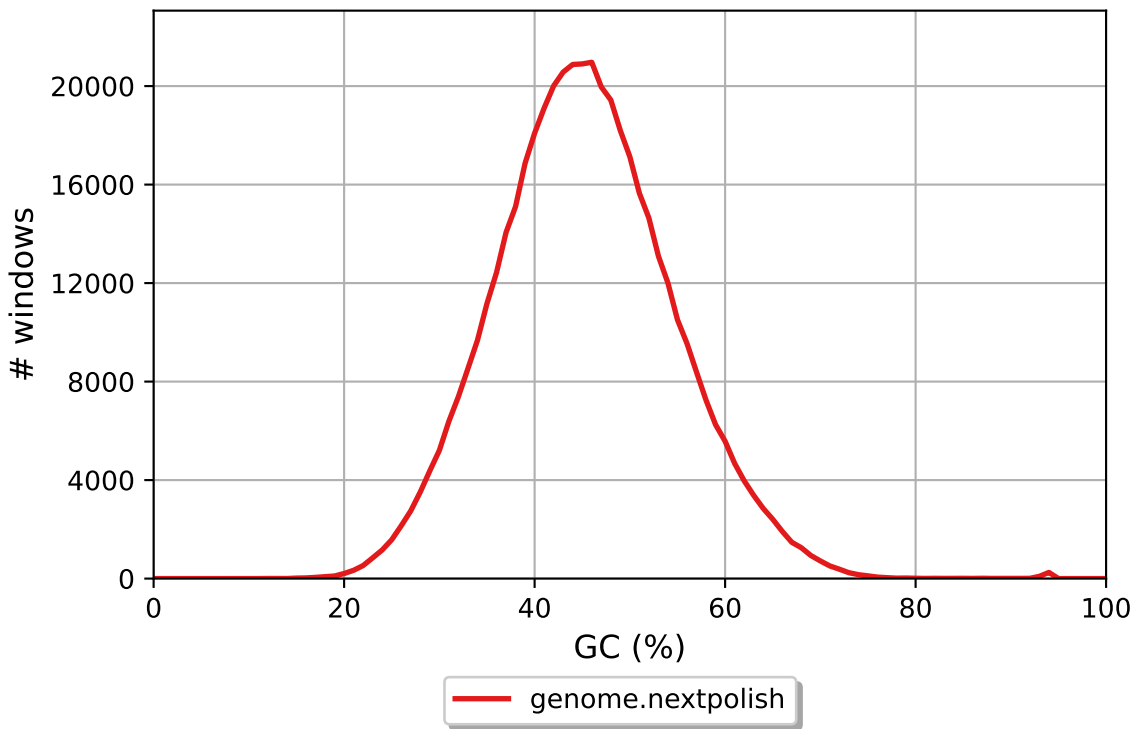
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



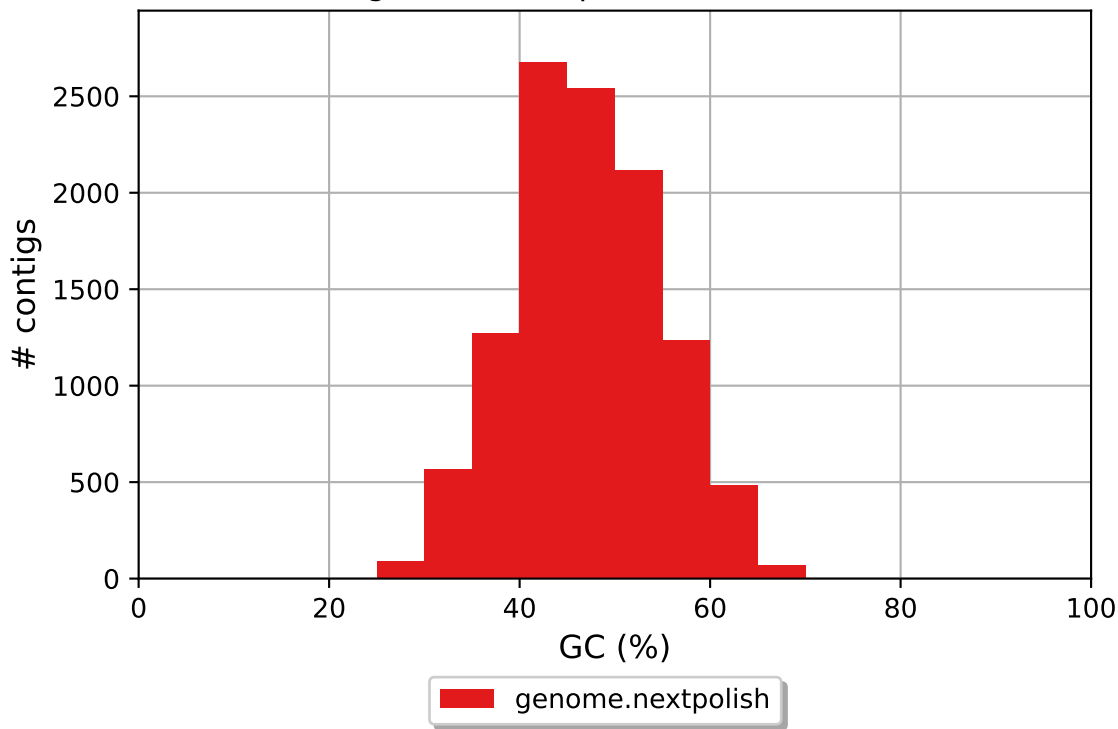
Cumulative length



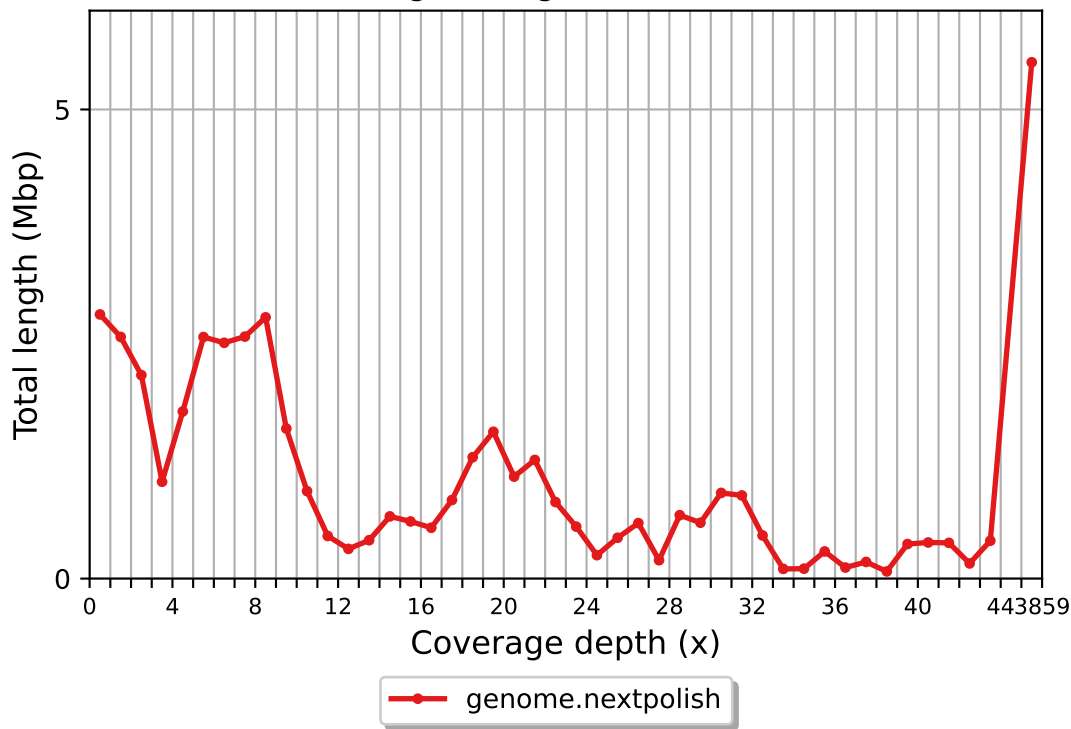
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

