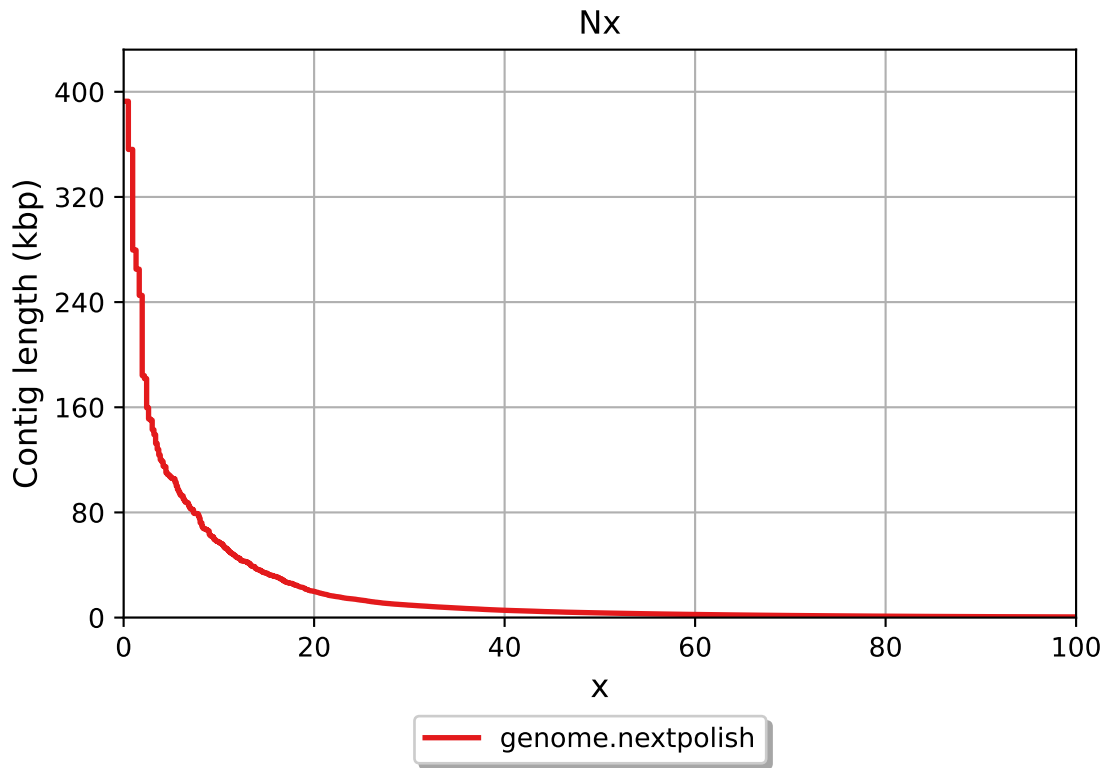


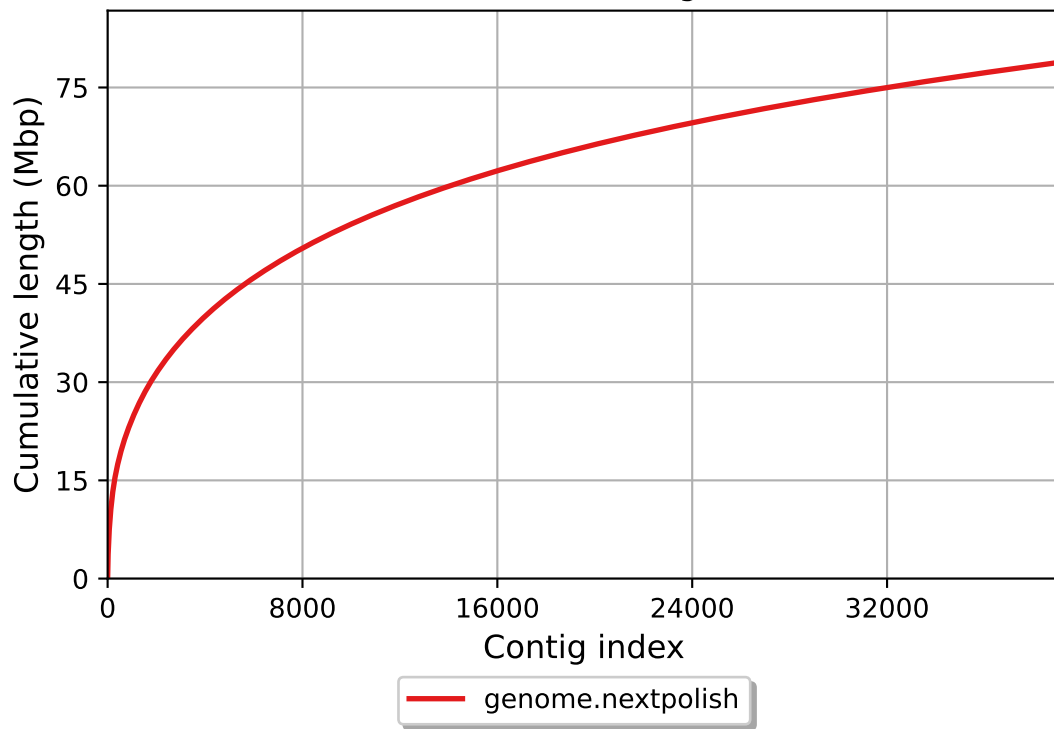
Report

	genome.nextpolish
# contigs (>= 0 bp)	145462
# contigs (>= 1000 bp)	17964
# contigs (>= 5000 bp)	2389
# contigs (>= 10000 bp)	850
# contigs (>= 25000 bp)	245
# contigs (>= 50000 bp)	91
Total length (>= 0 bp)	109098102
Total length (>= 1000 bp)	64344786
Total length (>= 5000 bp)	33451730
Total length (>= 10000 bp)	22899788
Total length (>= 25000 bp)	14135340
Total length (>= 50000 bp)	8785222
# contigs	39104
Largest contig	392780
Total length	78852150
GC (%)	50.17
N50	3571
N90	720
auN	21208.1
L50	3822
L90	25819
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	22 + 11 part

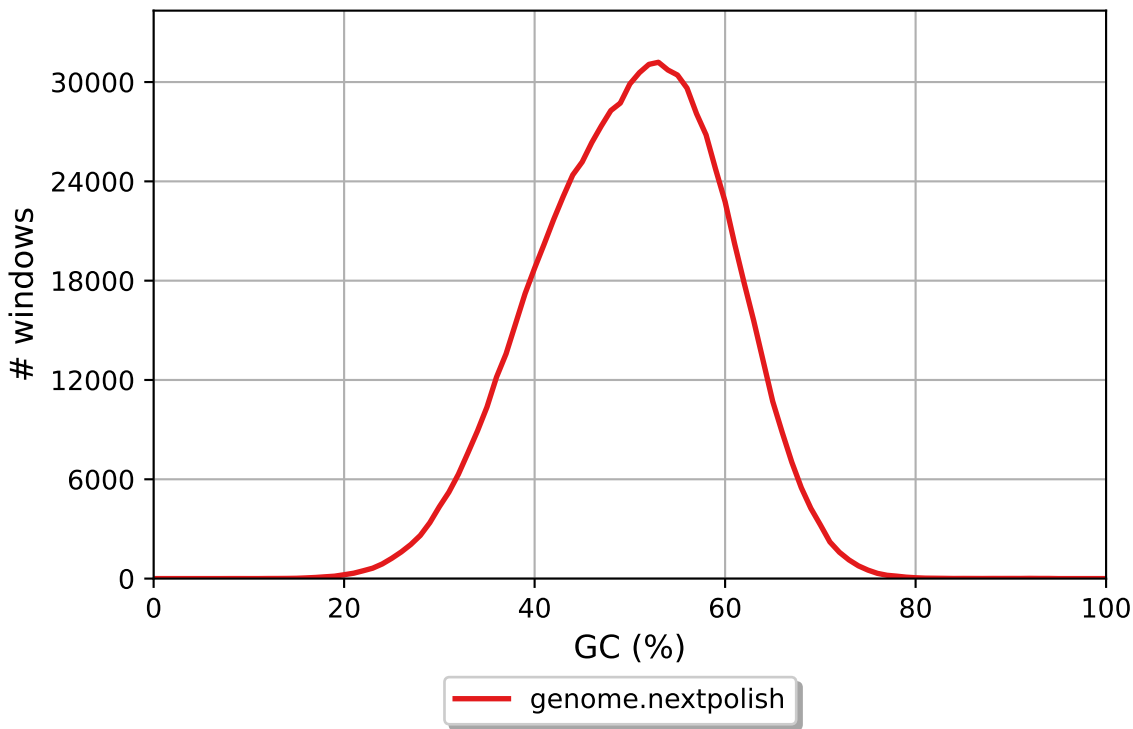
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



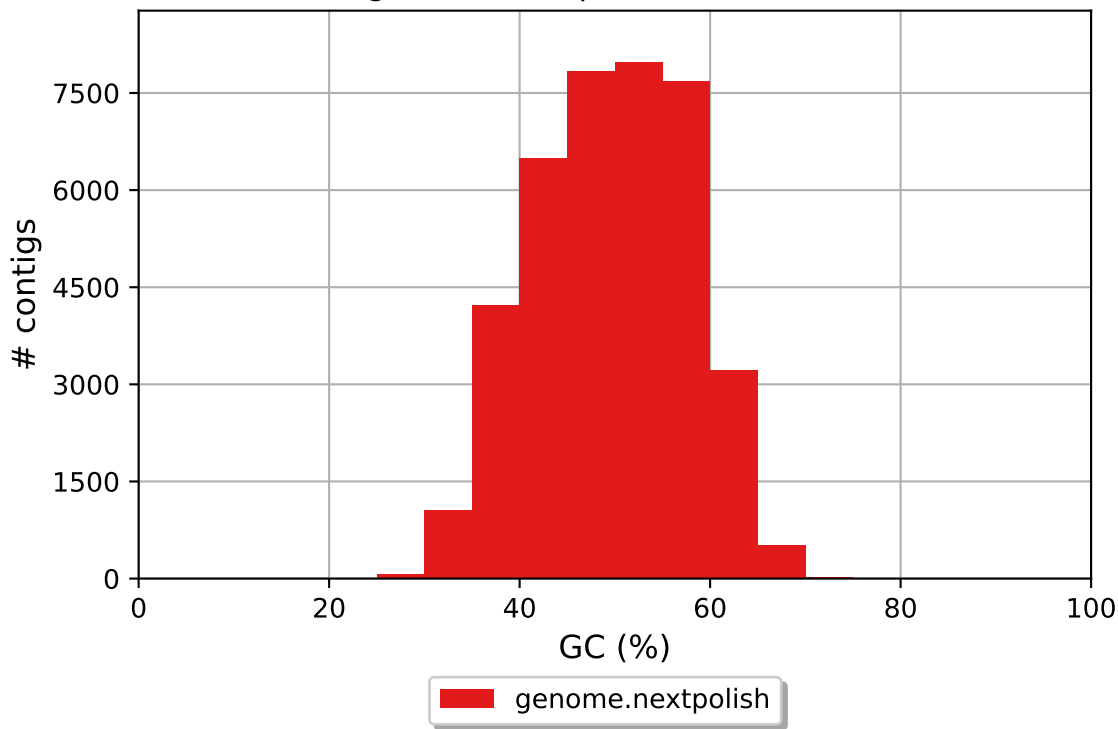
Cumulative length



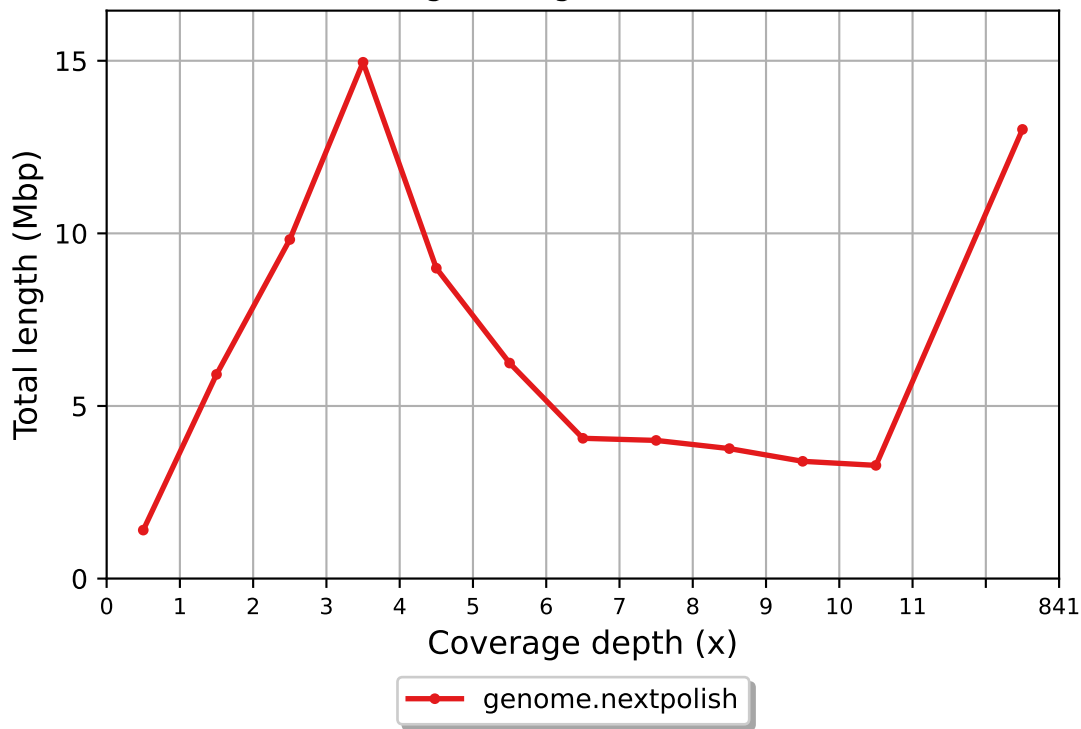
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

