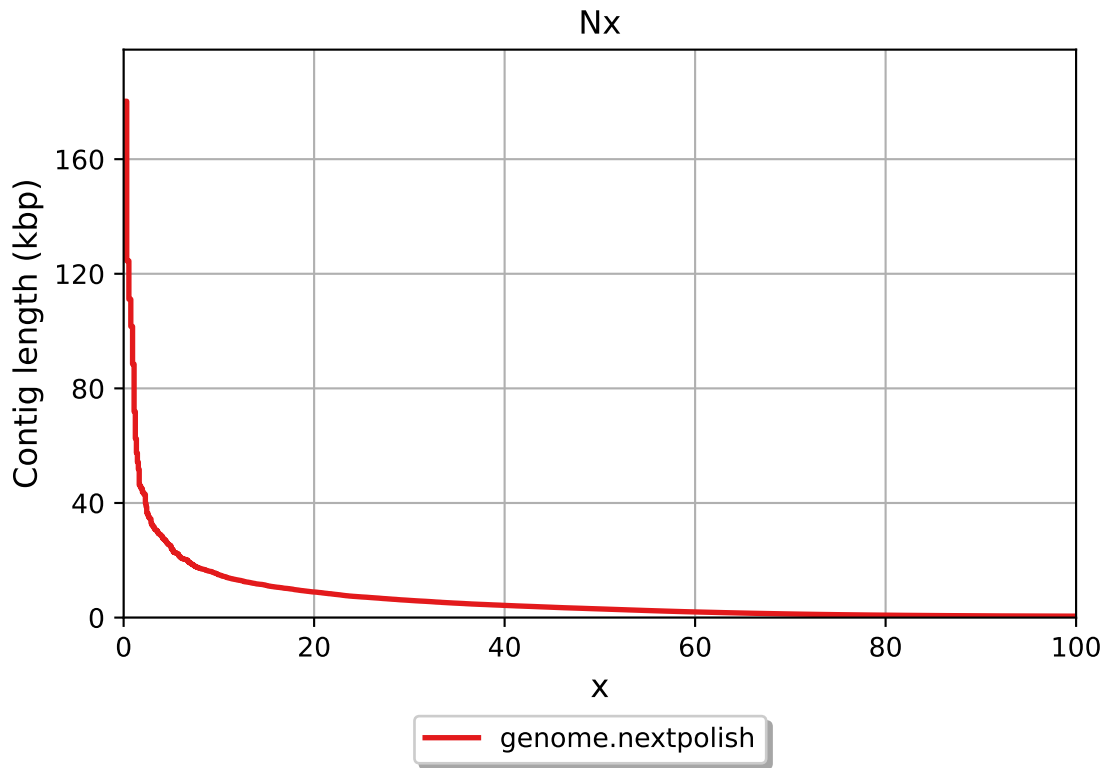


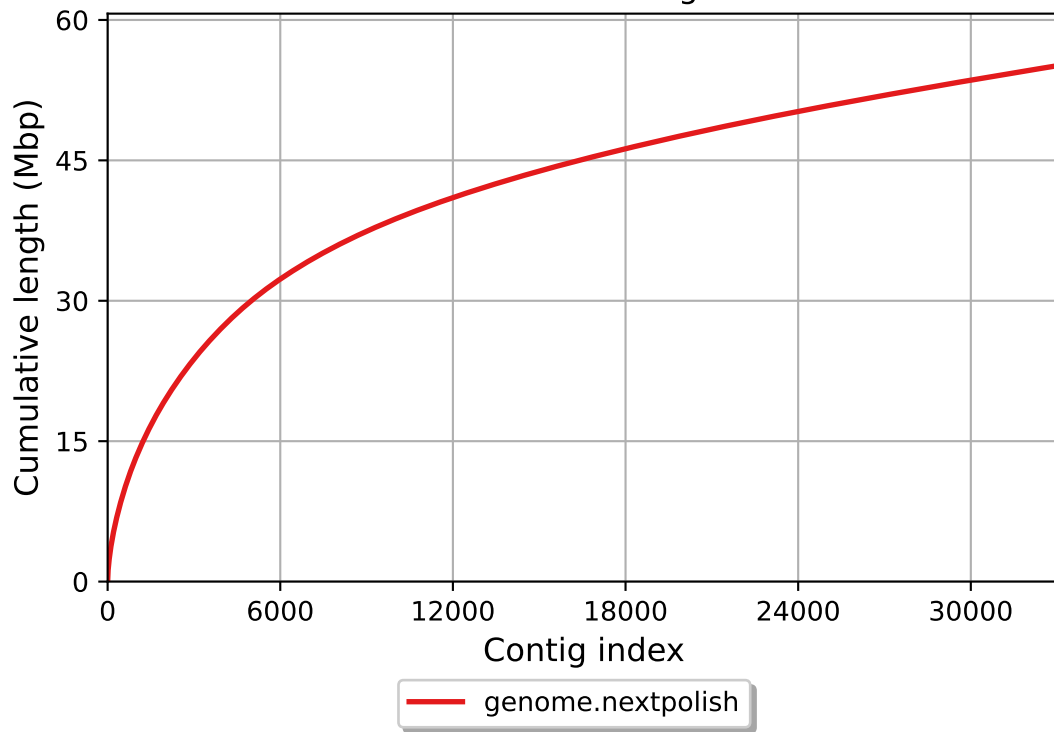
Report

	genome.nextpolish
# contigs (>= 0 bp)	60937
# contigs (>= 1000 bp)	12580
# contigs (>= 5000 bp)	1996
# contigs (>= 10000 bp)	571
# contigs (>= 25000 bp)	67
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	66047335
Total length (>= 1000 bp)	41616515
Total length (>= 5000 bp)	19389819
Total length (>= 10000 bp)	9702934
Total length (>= 25000 bp)	2726115
Total length (>= 50000 bp)	903775
# contigs	33103
Largest contig	180209
Total length	55159361
GC (%)	47.54
N50	3026
N90	617
auN	7166.9
L50	4117
L90	23038
# N's per 100 kbp	0.02
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 13 part

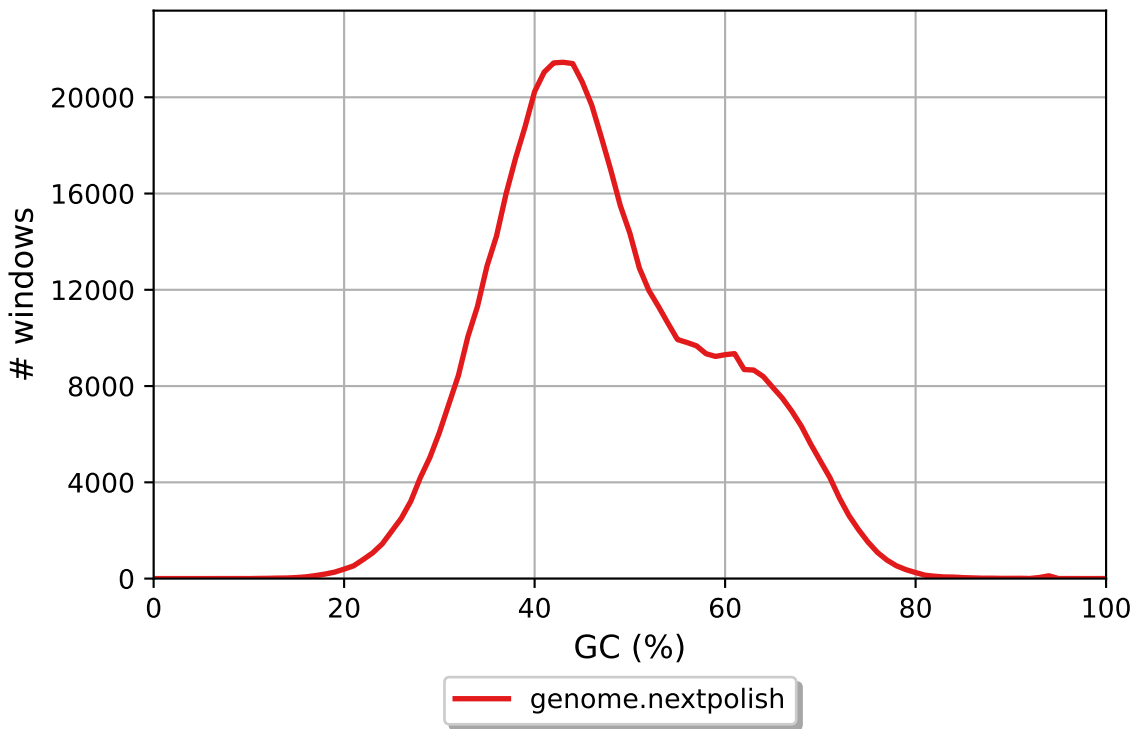
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



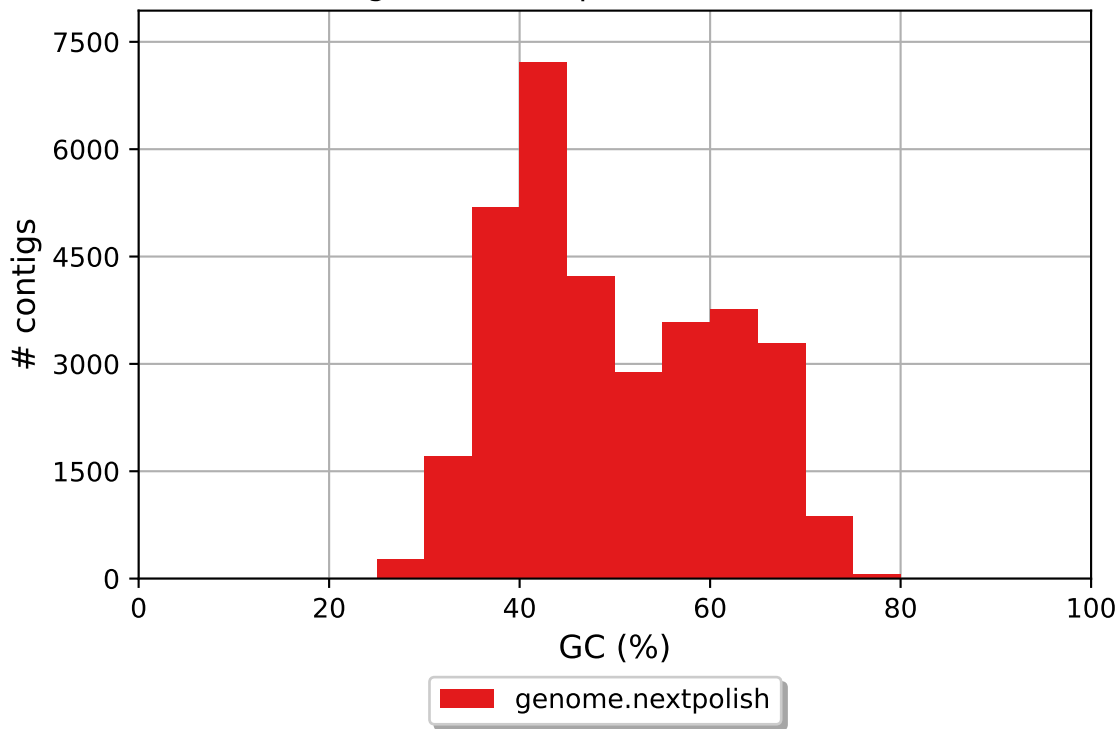
Cumulative length



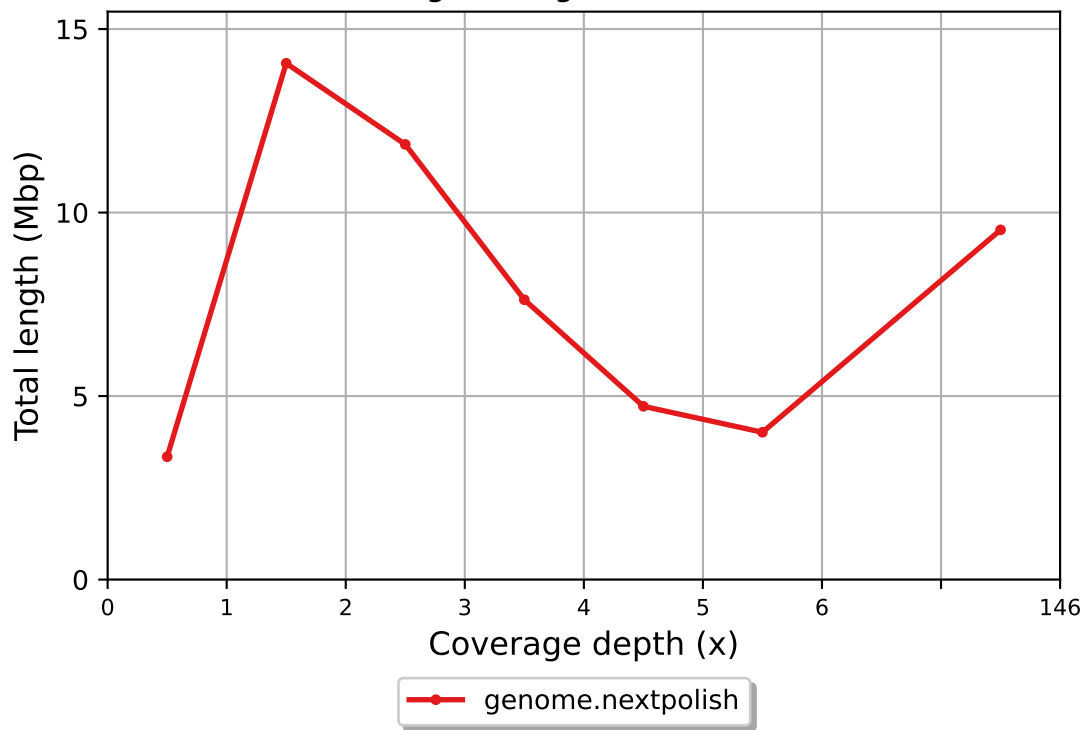
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

