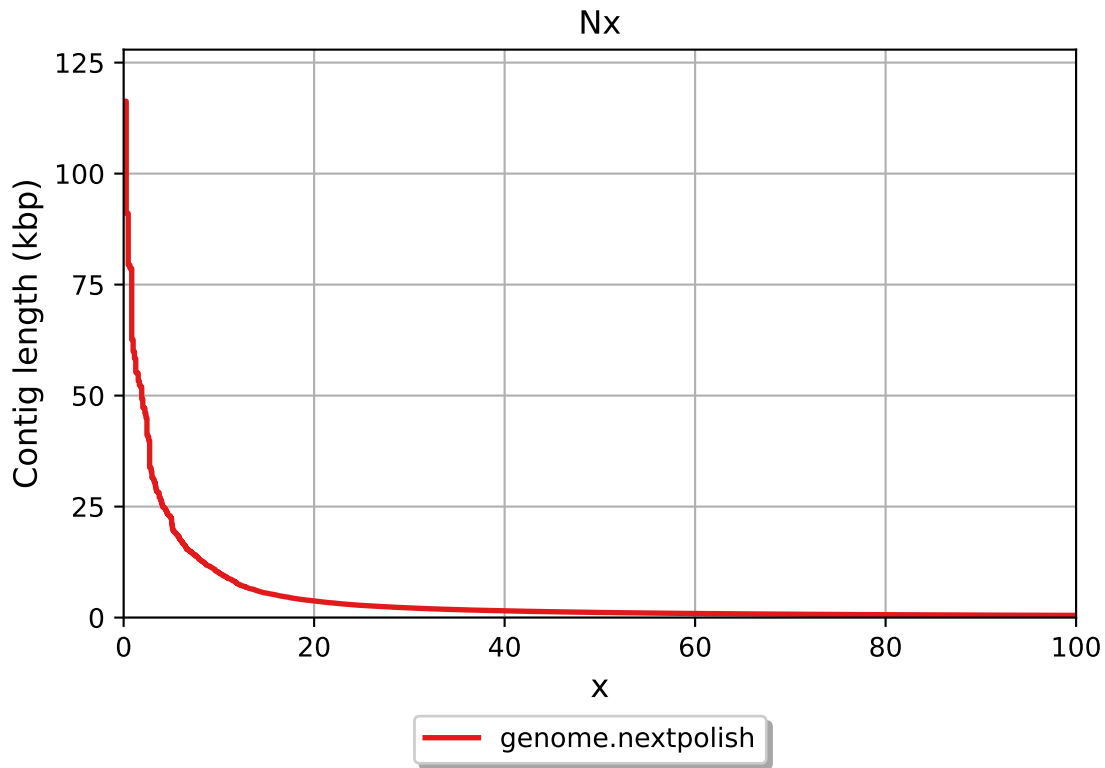


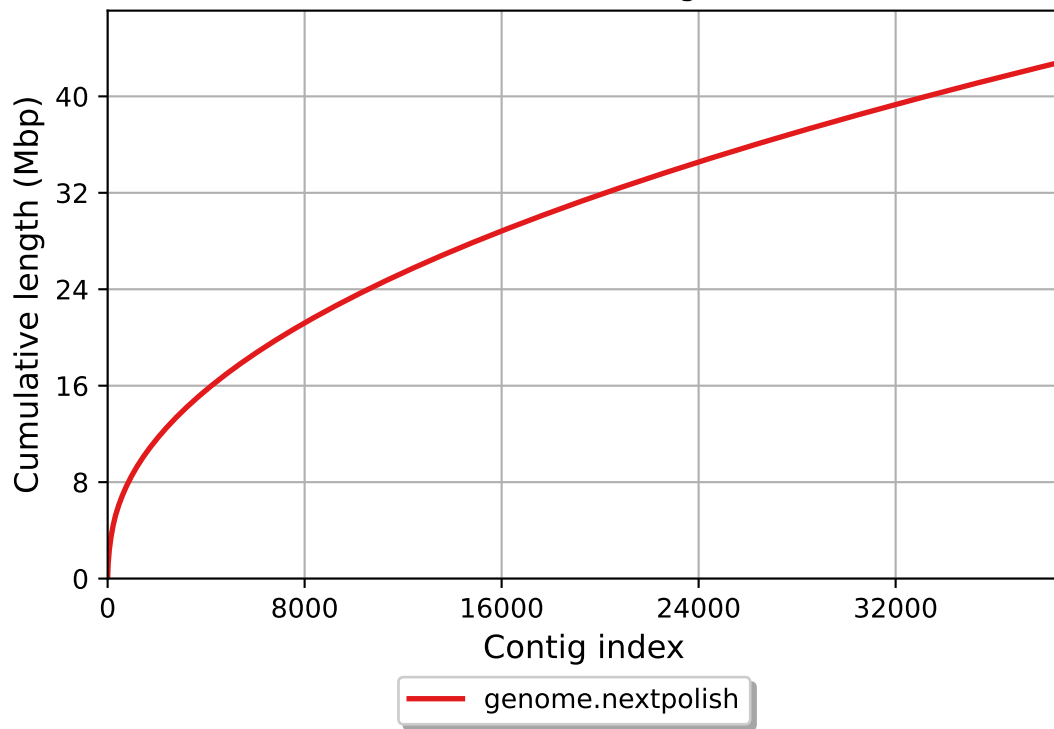
Report

	genome.nextpolish
# contigs (>= 0 bp)	205686
# contigs (>= 1000 bp)	10678
# contigs (>= 5000 bp)	608
# contigs (>= 10000 bp)	214
# contigs (>= 25000 bp)	40
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	91290585
Total length (>= 1000 bp)	24096958
Total length (>= 5000 bp)	6941997
Total length (>= 10000 bp)	4316982
Total length (>= 25000 bp)	1755431
Total length (>= 50000 bp)	814006
# contigs	38679
Largest contig	116292
Total length	42828741
GC (%)	44.80
N50	1155
N90	567
auN	4667.7
L50	8176
L90	30610
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 16 part

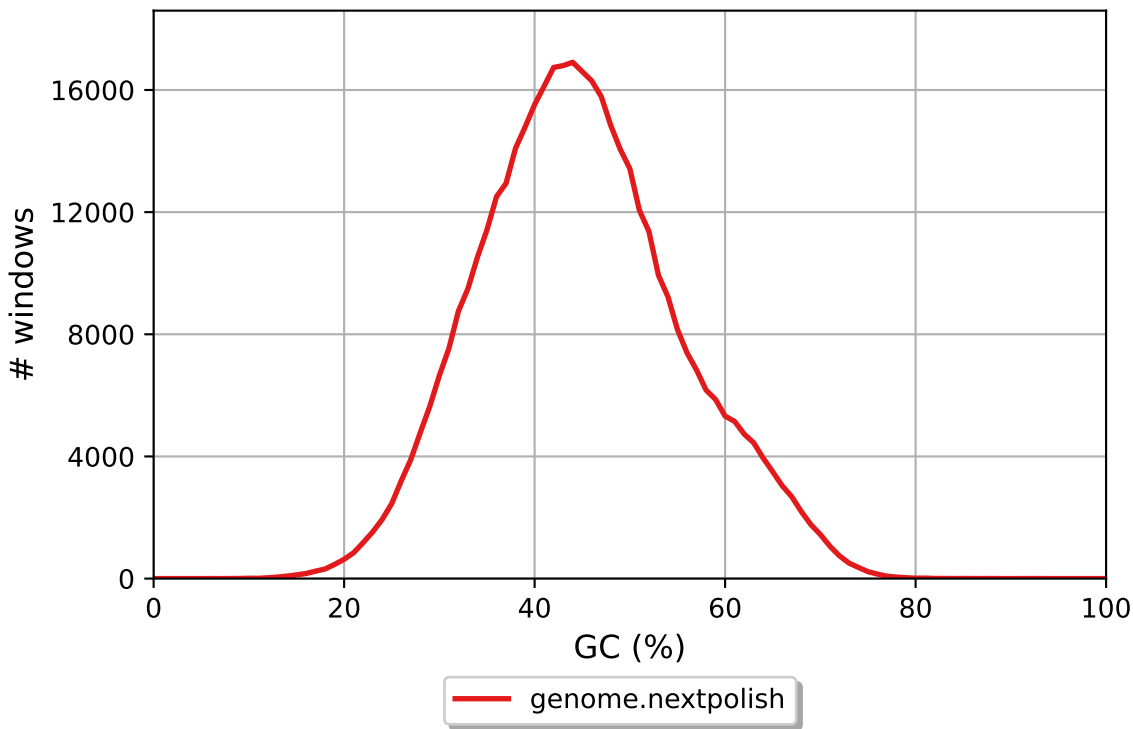
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



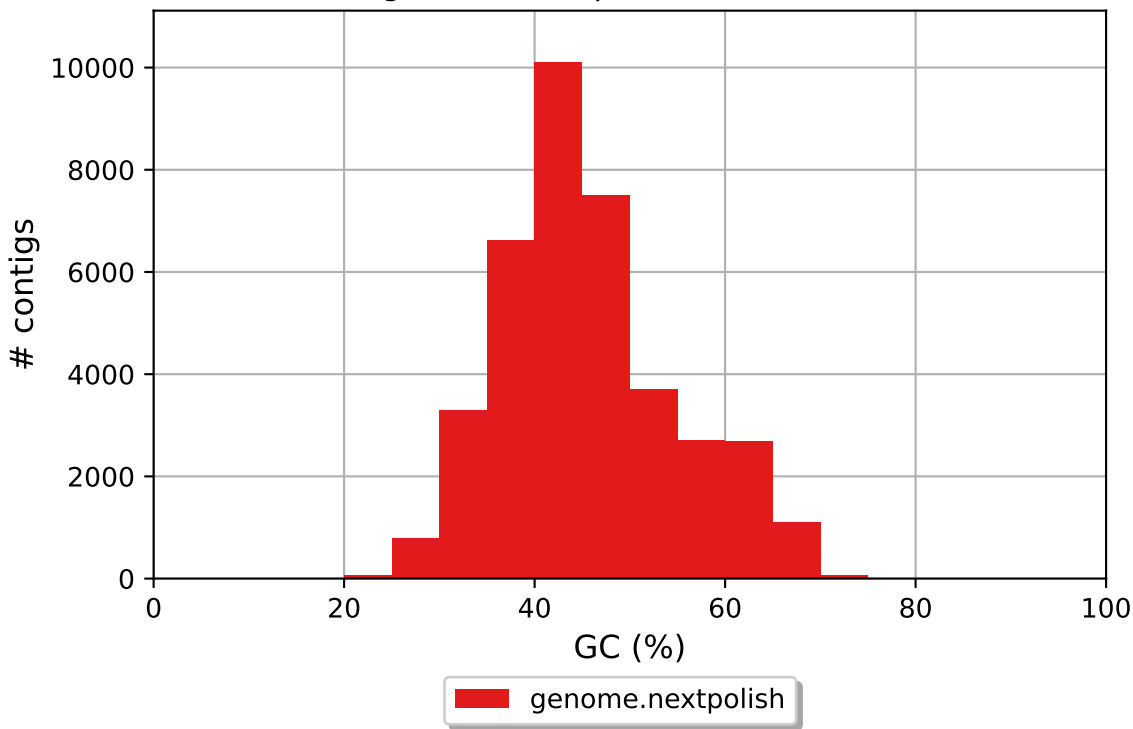
Cumulative length



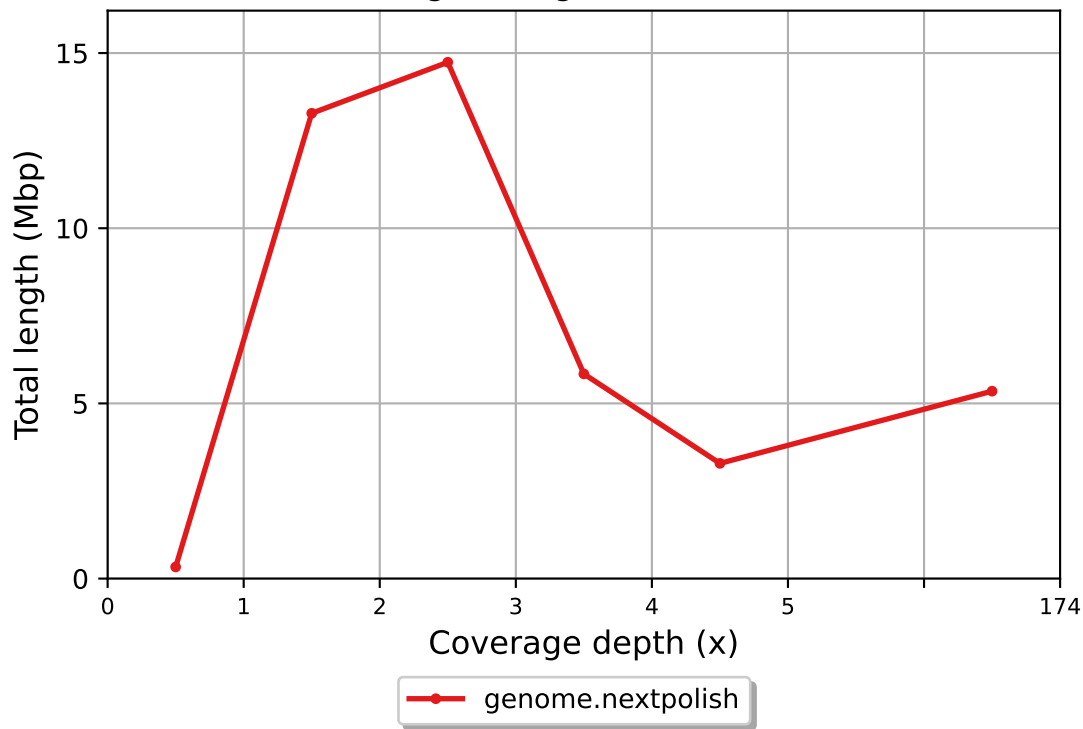
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

