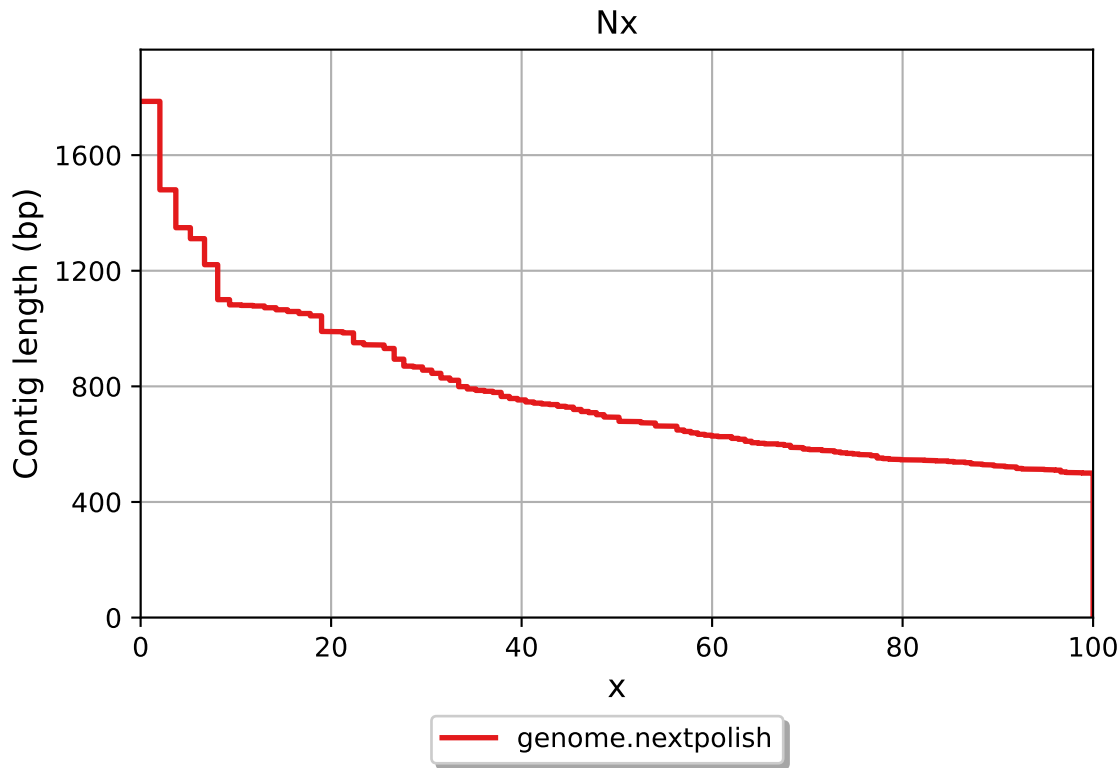


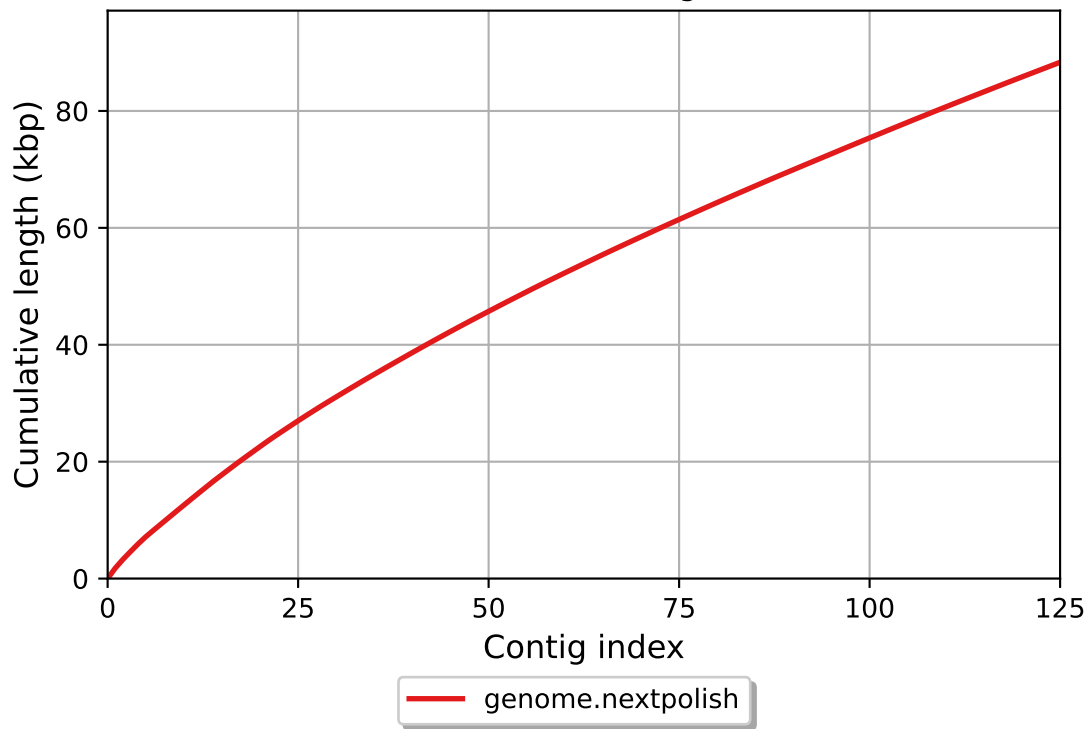
Report

	genome.nextpolish
# contigs (>= 0 bp)	3887
# contigs (>= 1000 bp)	14
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	783209
Total length (>= 1000 bp)	16779
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	125
Largest contig	1786
Total length	88328
GC (%)	53.93
N50	693
N90	525
auN	777.8
L50	48
L90	108
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

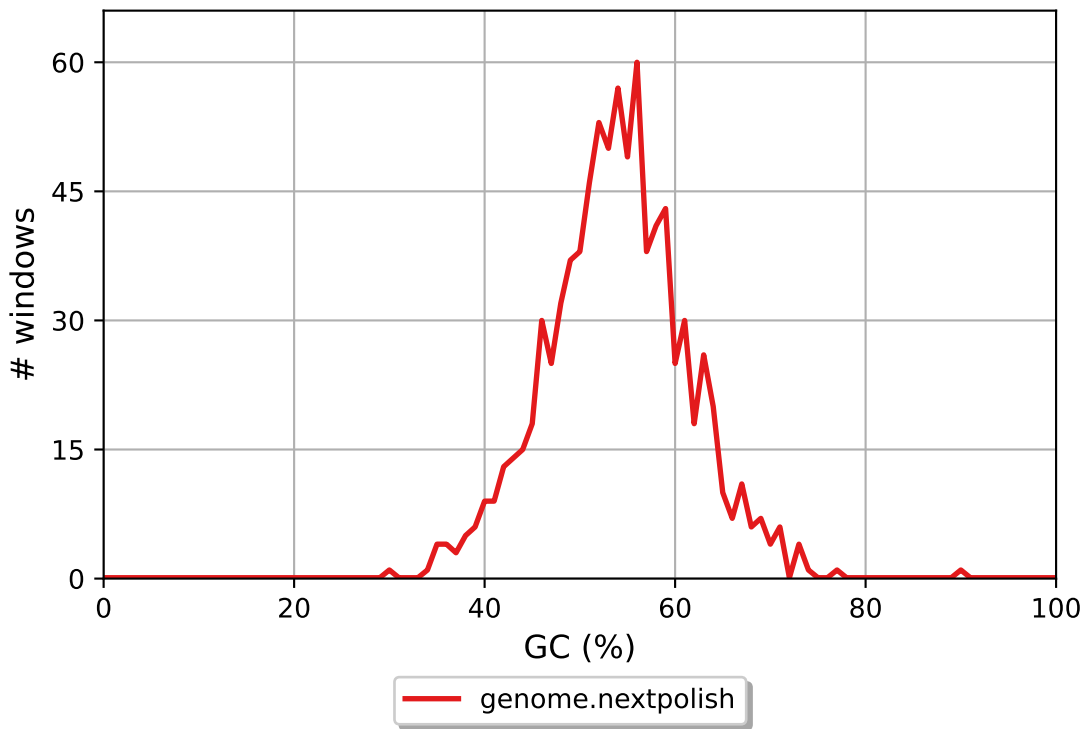
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



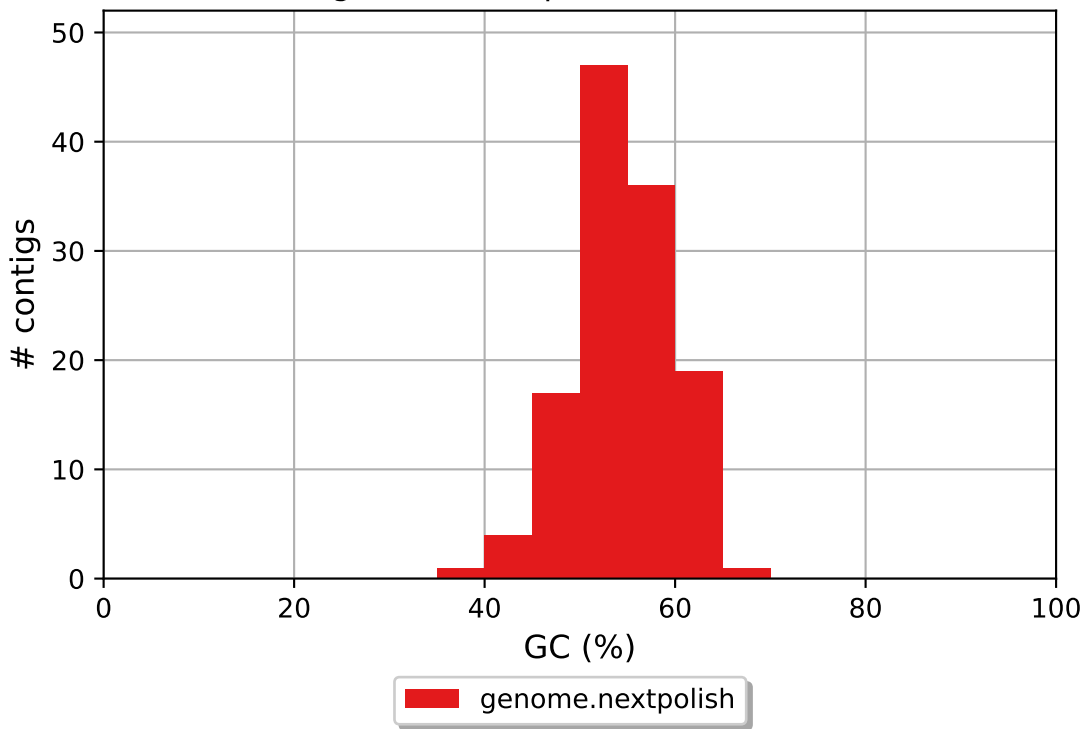
Cumulative length



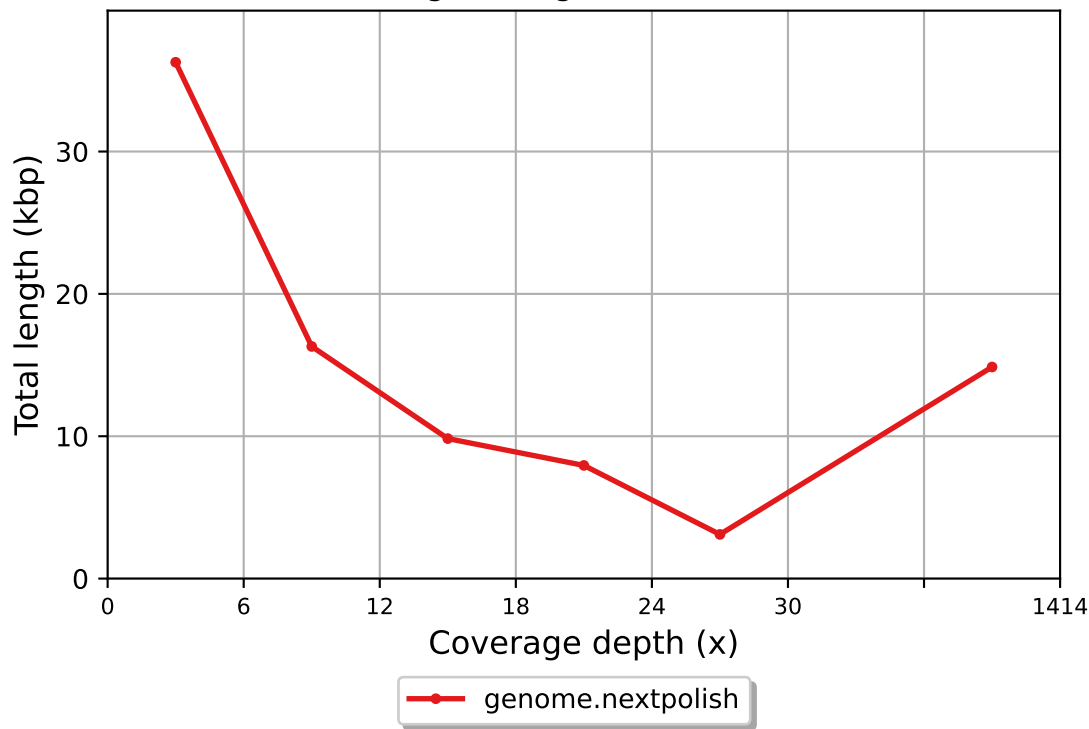
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 6x)



genome.nextpolish coverage histogram (bin size: 6x)

