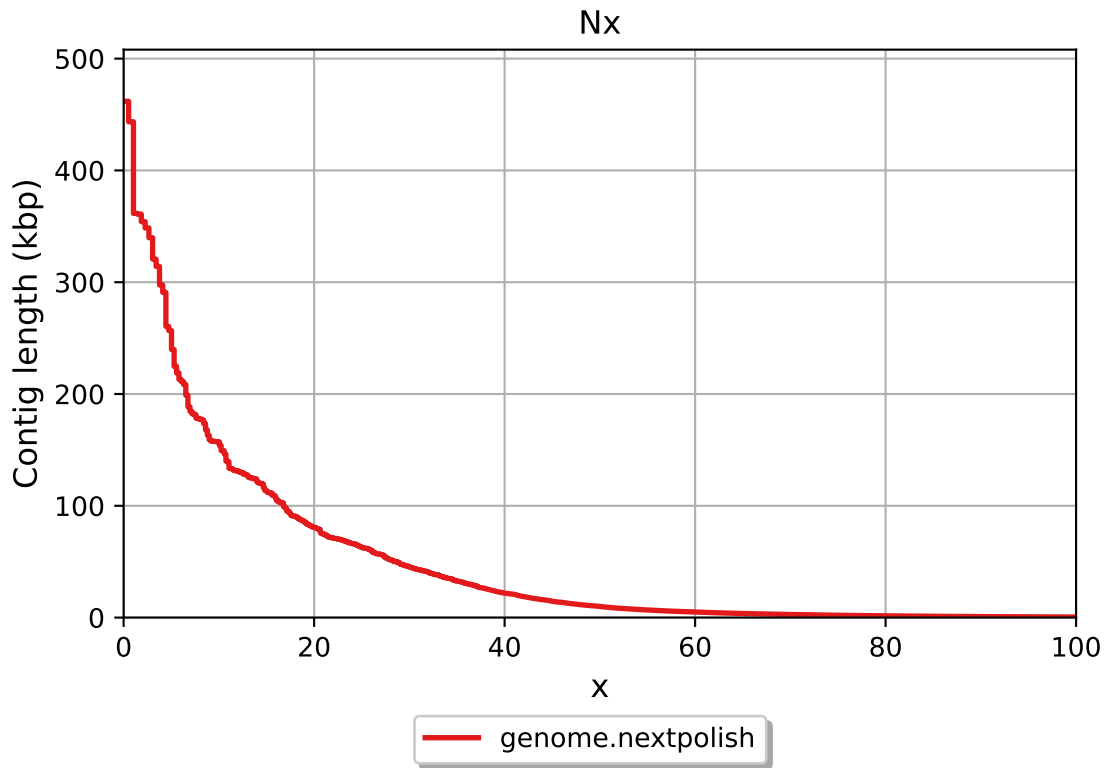


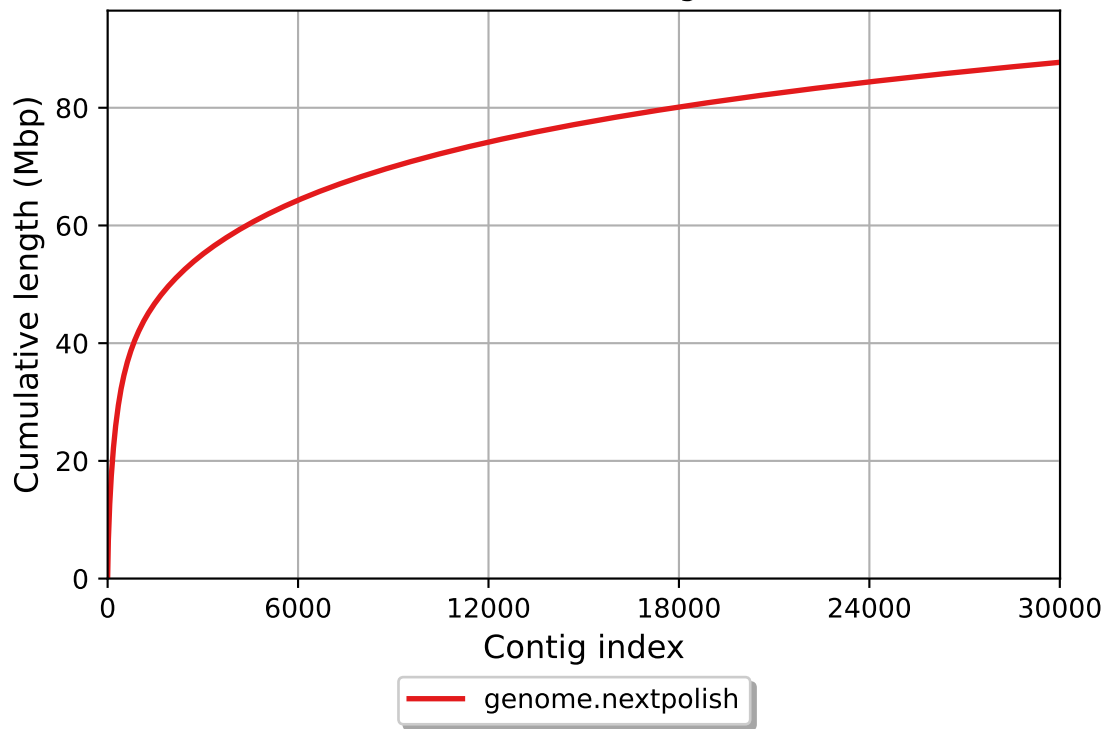
Report

	genome.nextpolish
# contigs (>= 0 bp)	96788
# contigs (>= 1000 bp)	14688
# contigs (>= 5000 bp)	2446
# contigs (>= 10000 bp)	1144
# contigs (>= 25000 bp)	479
# contigs (>= 50000 bp)	233
Total length (>= 0 bp)	106875838
Total length (>= 1000 bp)	77130120
Total length (>= 5000 bp)	52616722
Total length (>= 10000 bp)	43812217
Total length (>= 25000 bp)	33710895
Total length (>= 50000 bp)	24979271
# contigs	30009
Largest contig	461856
Total length	87730452
GC (%)	45.90
N50	9953
N90	885
auN	50732.4
L50	1150
L90	16632
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 14 part

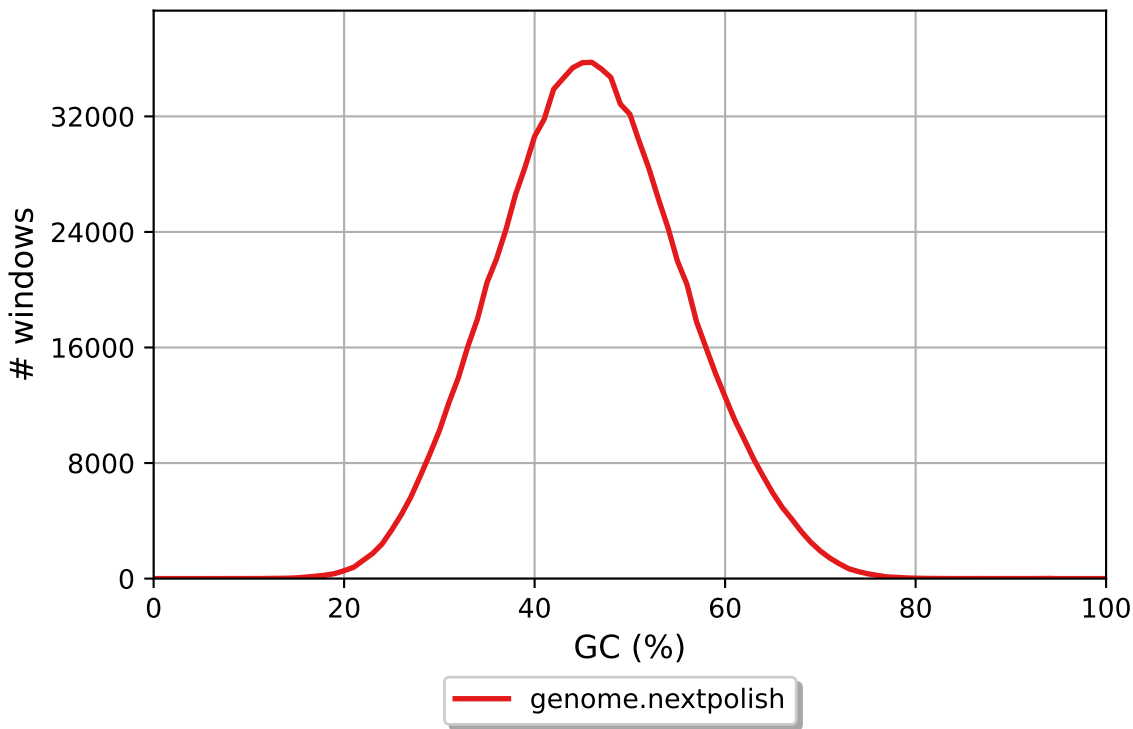
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



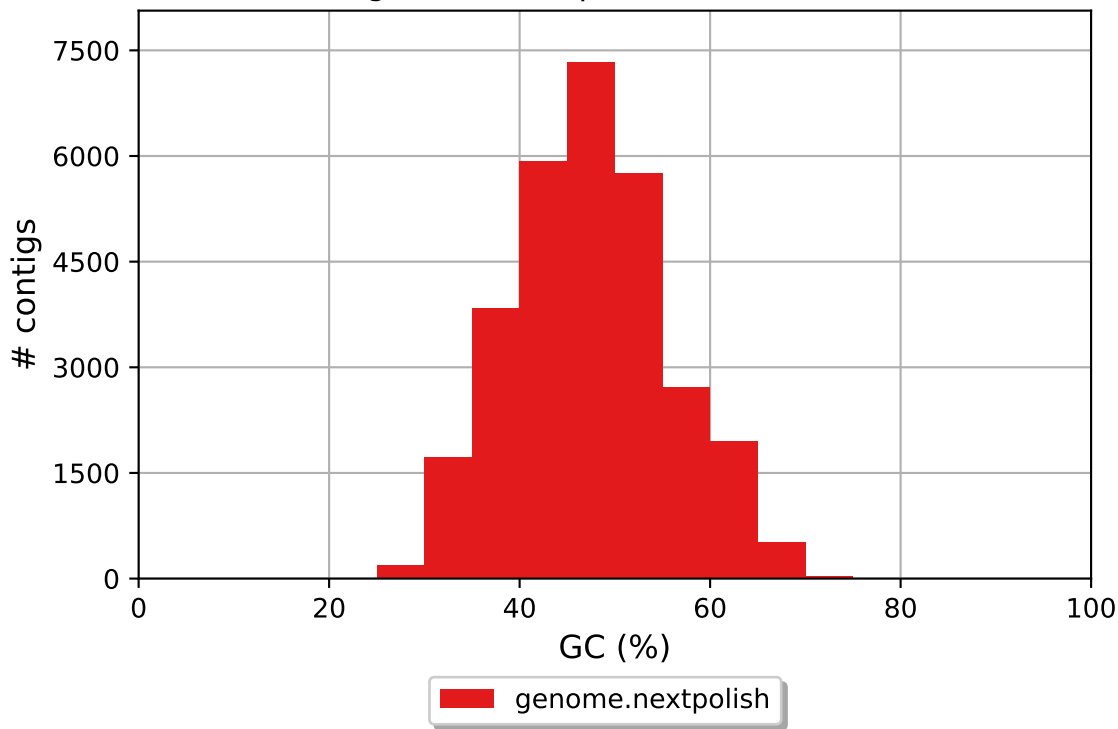
Cumulative length



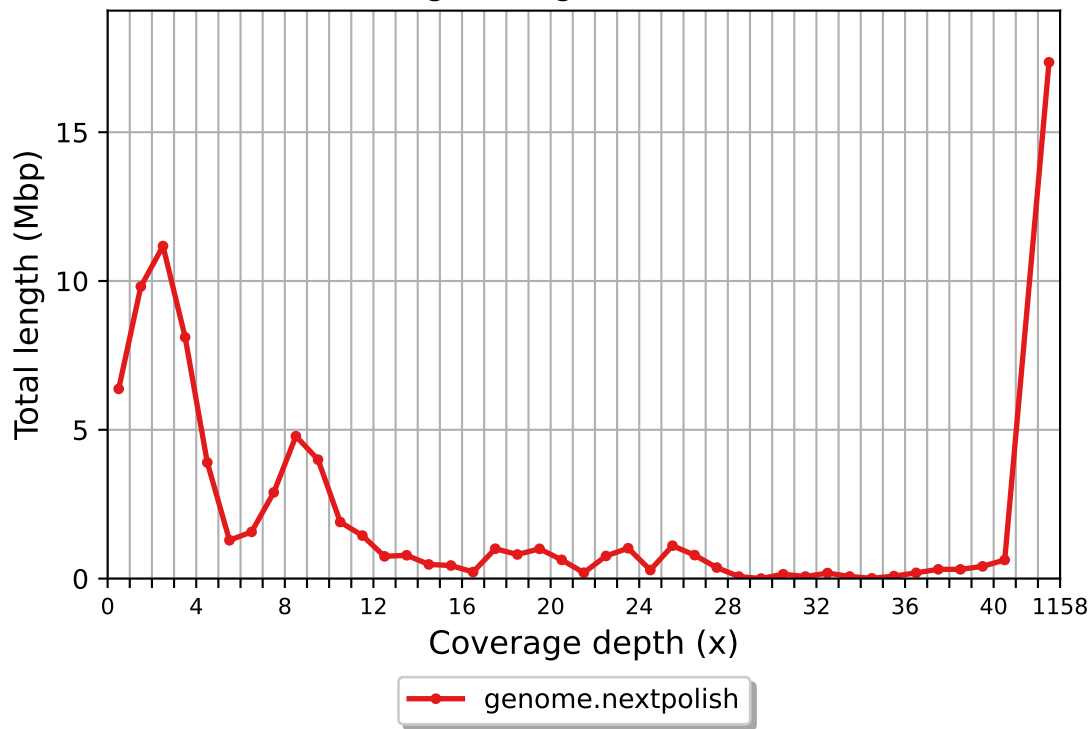
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

