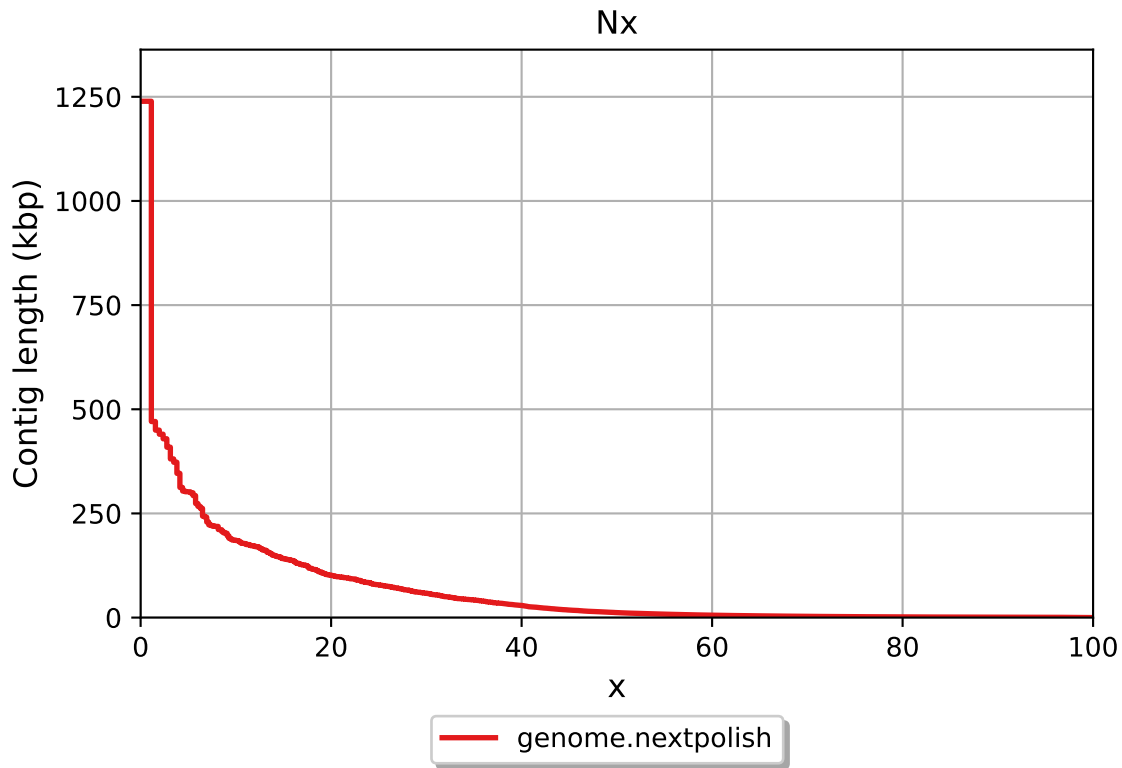


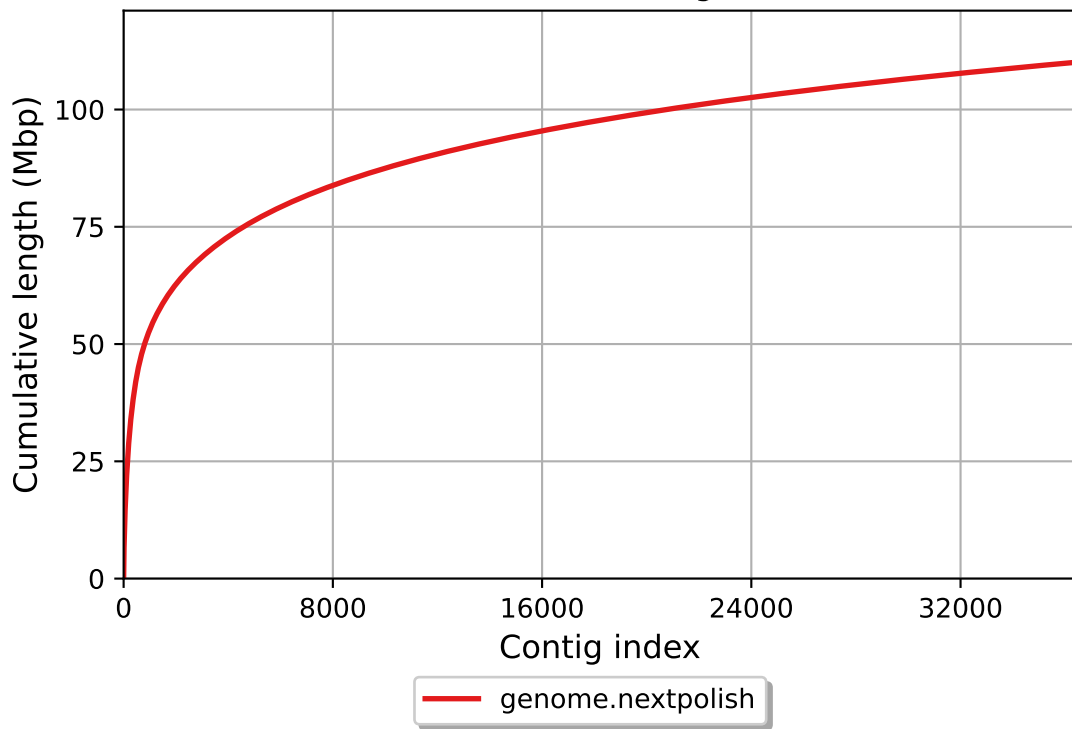
# Report

	genome.nextpolish
# contigs (>= 0 bp)	127496
# contigs (>= 1000 bp)	17445
# contigs (>= 5000 bp)	2938
# contigs (>= 10000 bp)	1392
# contigs (>= 25000 bp)	590
# contigs (>= 50000 bp)	301
Total length (>= 0 bp)	136175898
Total length (>= 1000 bp)	96952153
Total length (>= 5000 bp)	68283838
Total length (>= 10000 bp)	57653824
Total length (>= 25000 bp)	45654653
Total length (>= 50000 bp)	35346046
# contigs	36418
Largest contig	1239270
Total length	110074880
GC (%)	46.73
N50	11963
N90	896
auN	70823.3
L50	1151
L90	19681
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	49 + 17 part

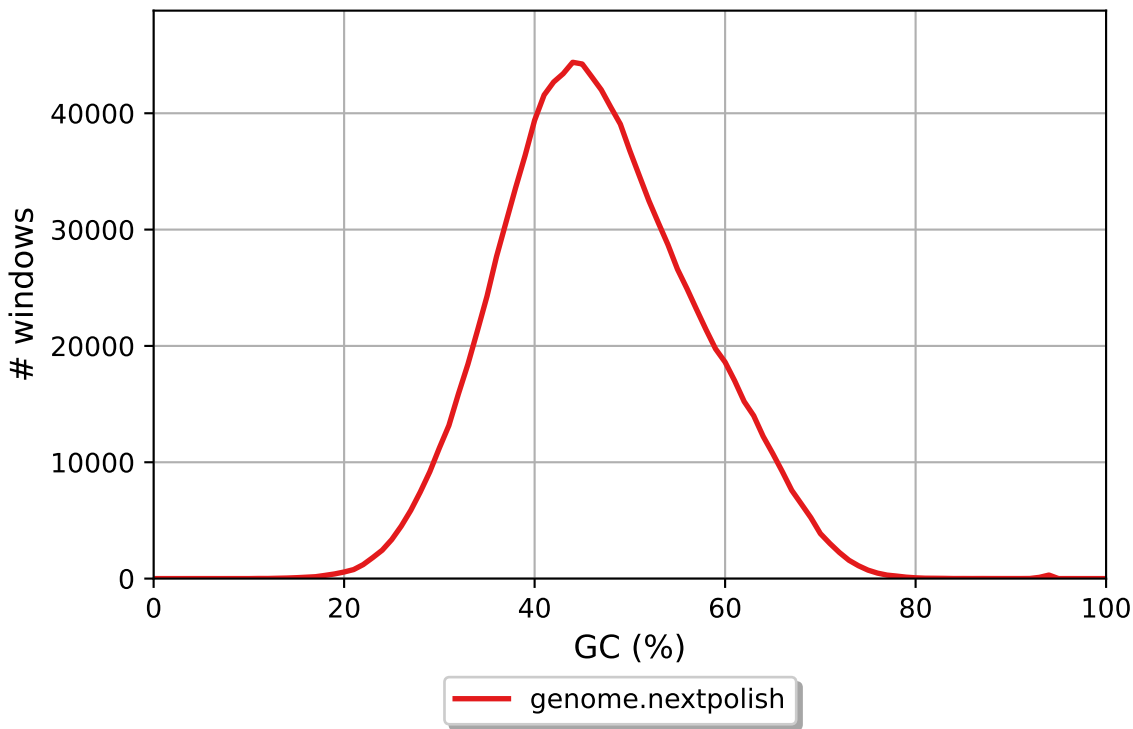
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



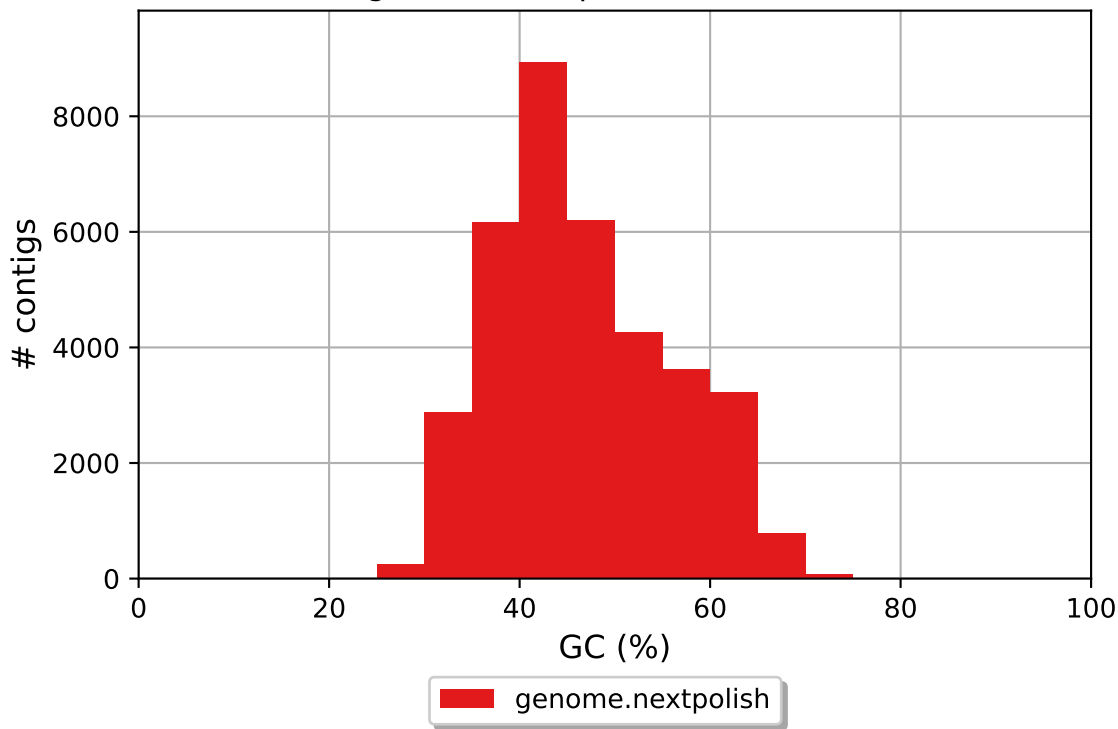
Cumulative length



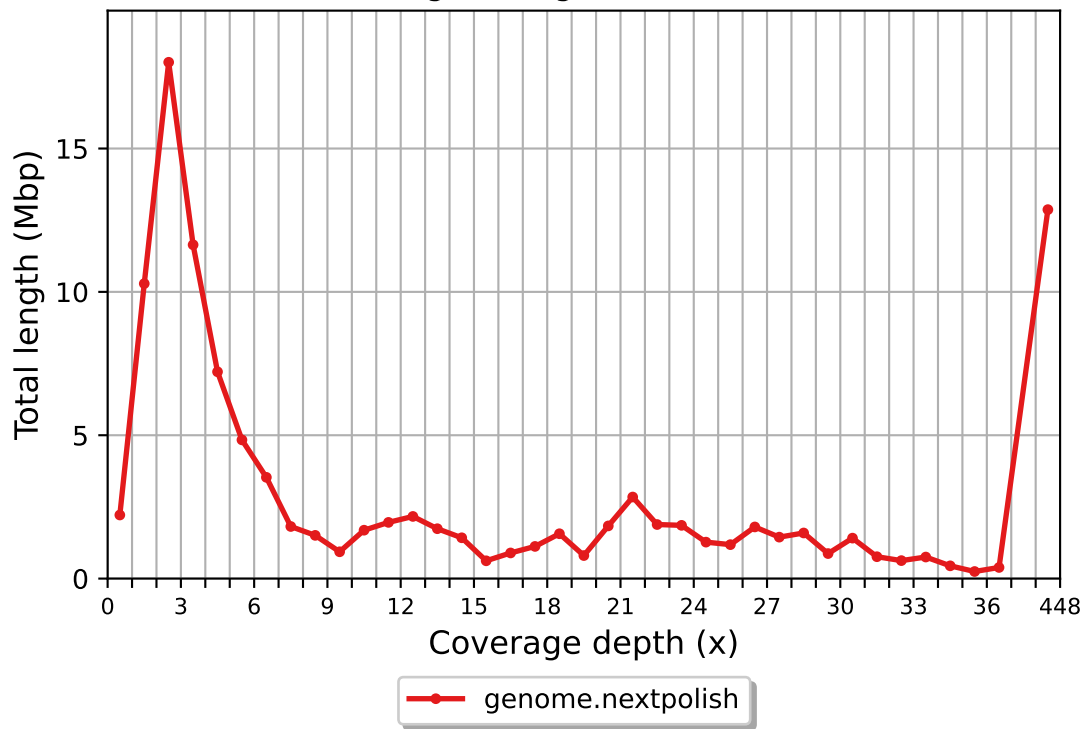
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

