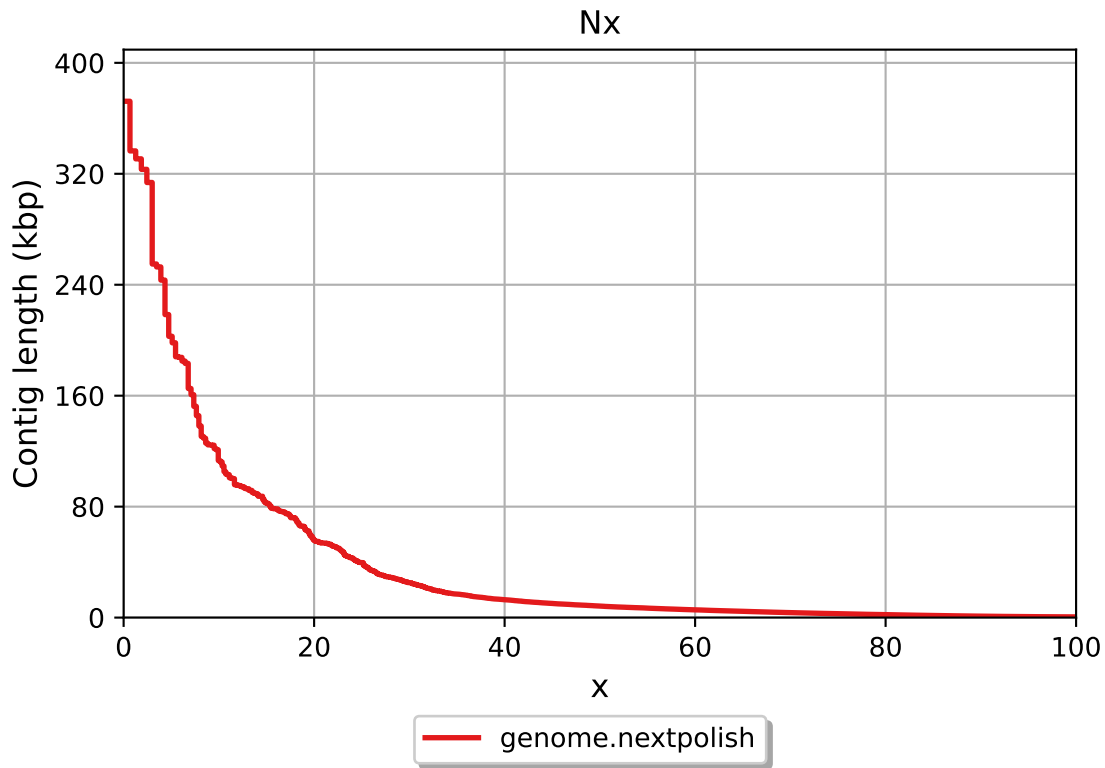


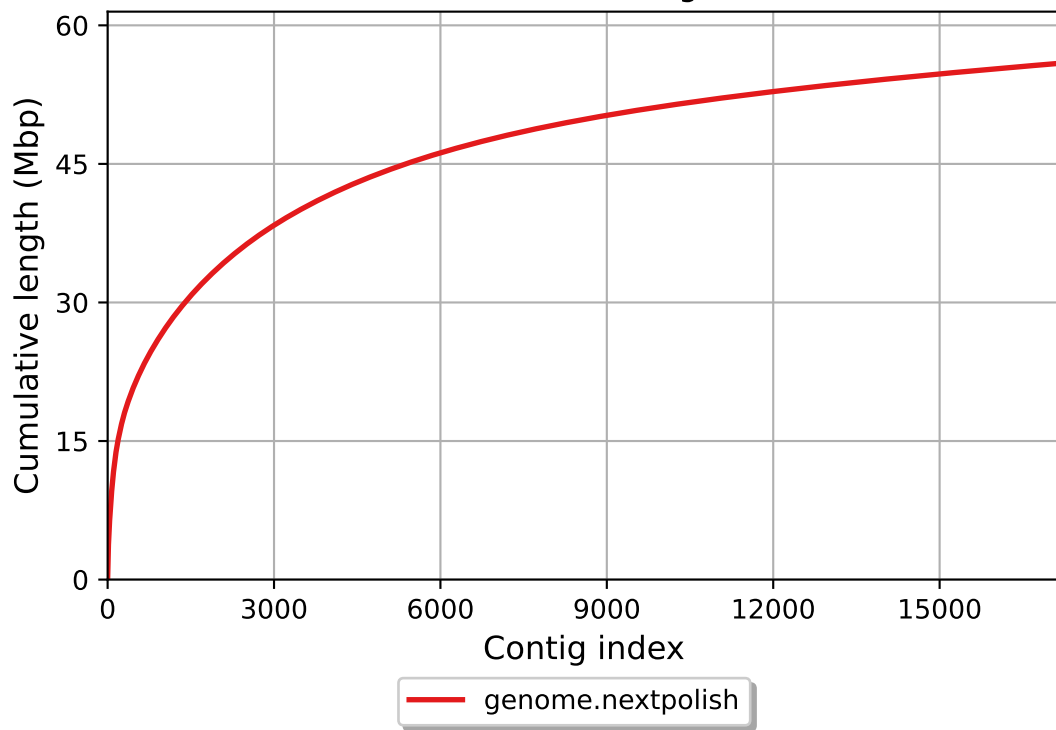
Report

	genome.nextpolish
# contigs (>= 0 bp)	72608
# contigs (>= 1000 bp)	9238
# contigs (>= 5000 bp)	2228
# contigs (>= 10000 bp)	843
# contigs (>= 25000 bp)	252
# contigs (>= 50000 bp)	125
Total length (>= 0 bp)	71316155
Total length (>= 1000 bp)	50502976
Total length (>= 5000 bp)	34983924
Total length (>= 10000 bp)	25356024
Total length (>= 25000 bp)	16872727
Total length (>= 50000 bp)	12654992
# contigs	17172
Largest contig	372290
Total length	55894188
GC (%)	45.37
N50	8302
N90	1028
auN	39394.4
L50	1127
L90	9043
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 9 part

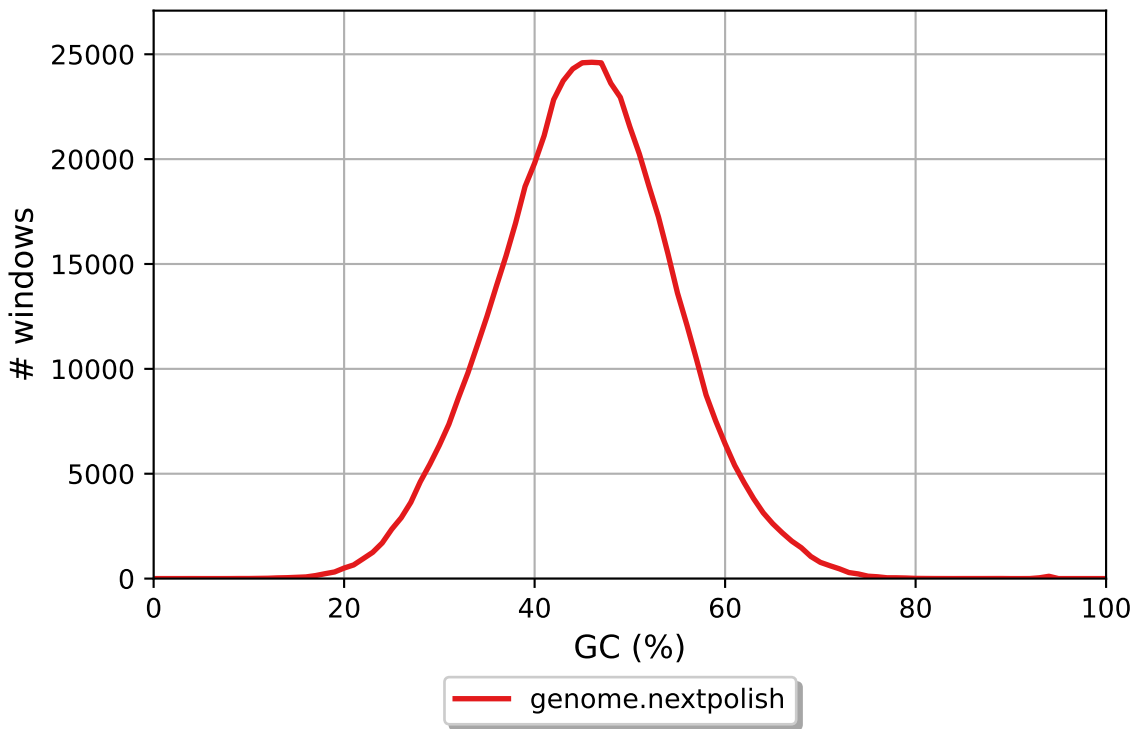
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



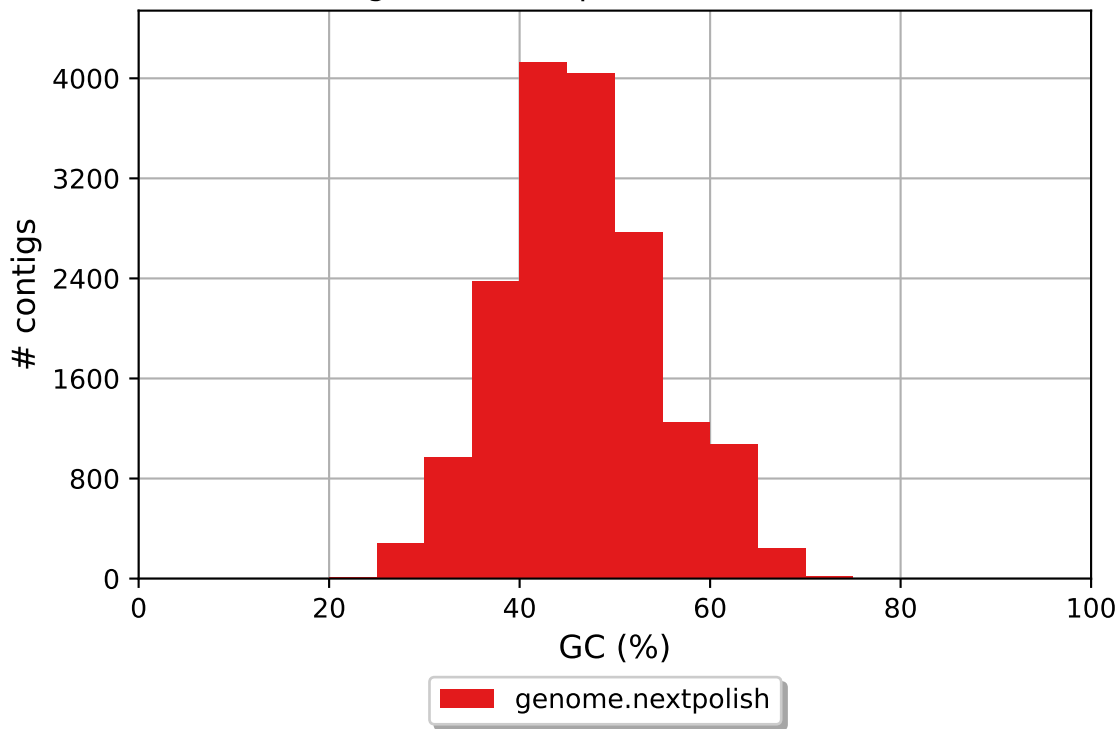
Cumulative length



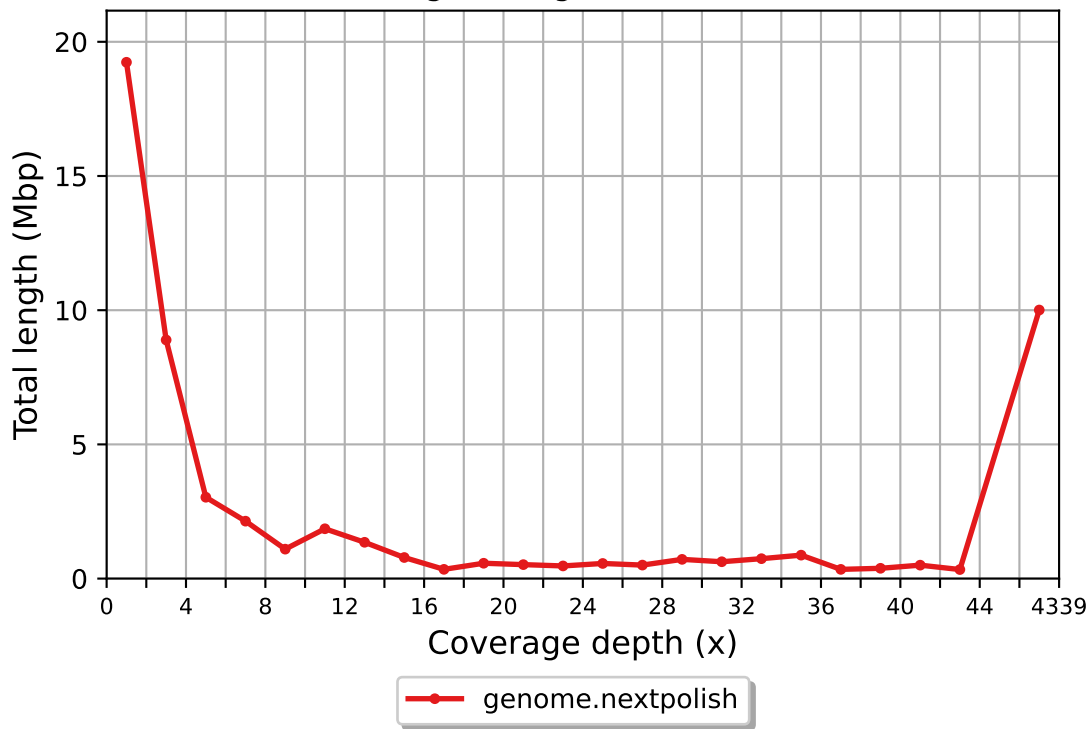
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 2x)



genome.nextpolish coverage histogram (bin size: 2x)

