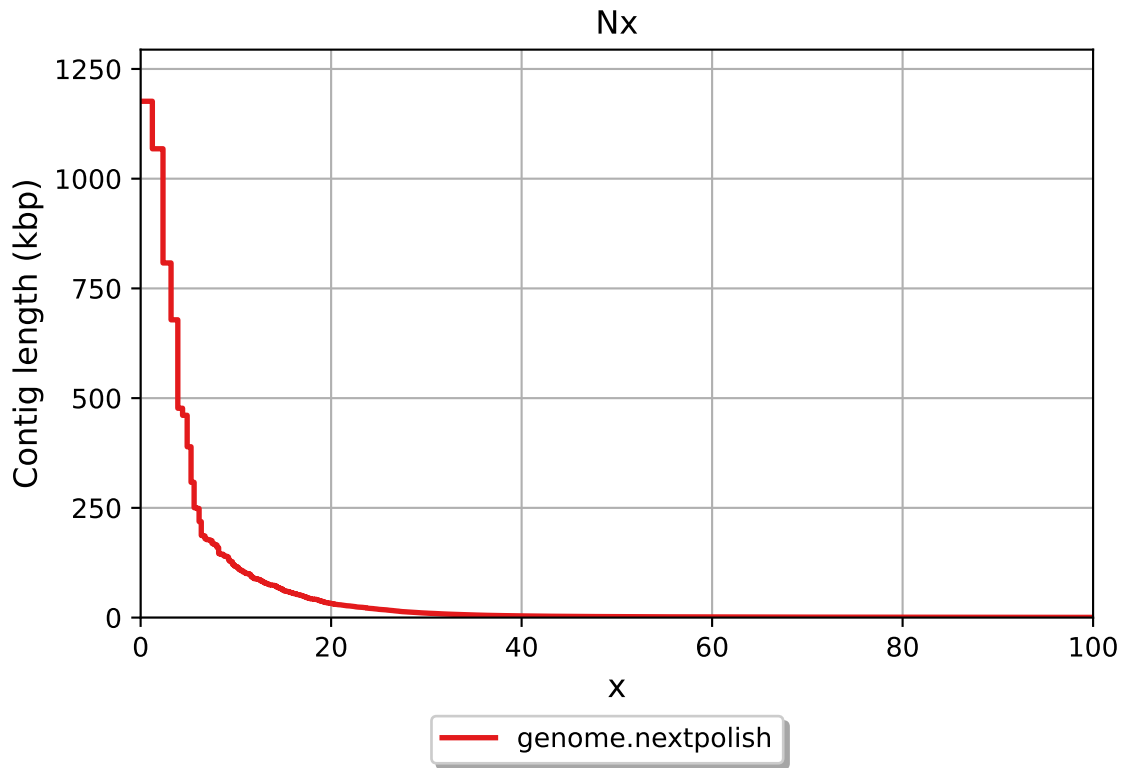
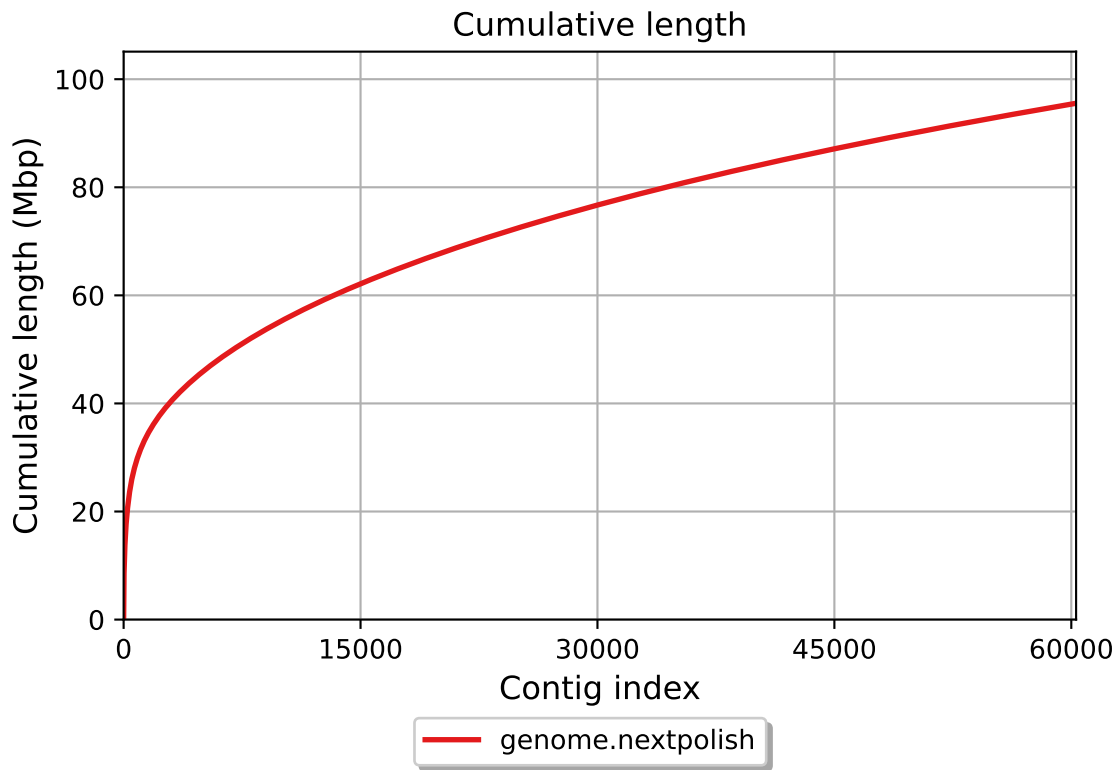


Report

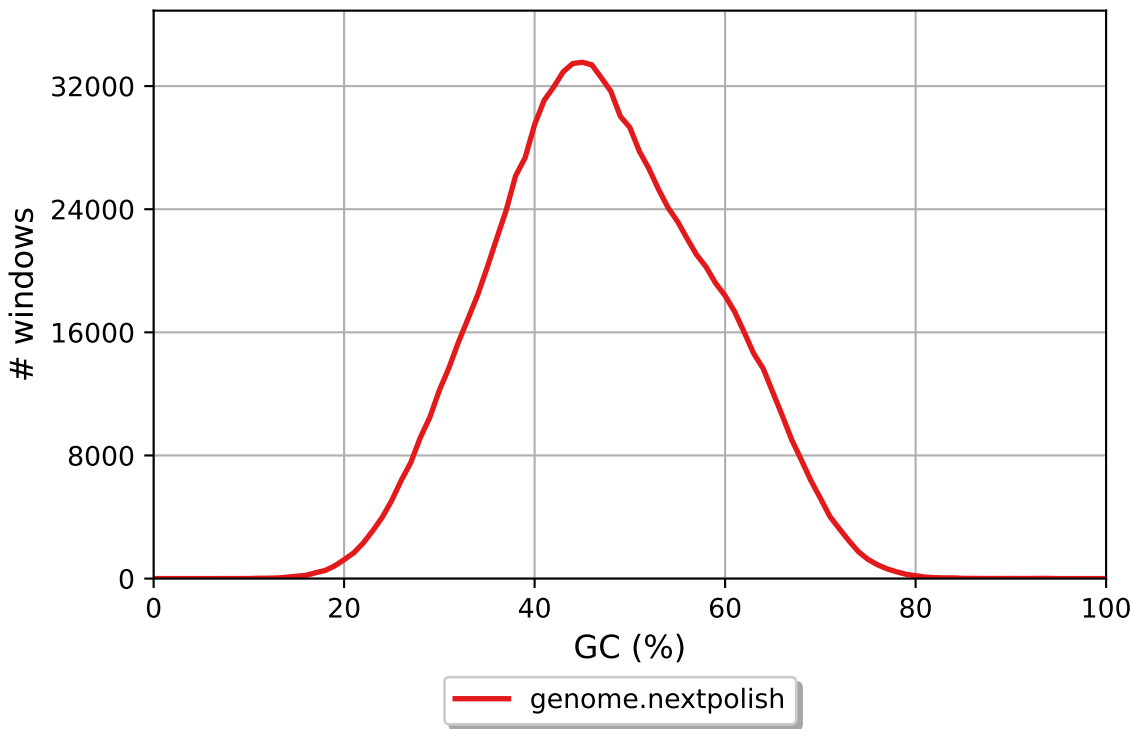
	genome.nextpolish
# contigs (>= 0 bp)	216968
# contigs (>= 1000 bp)	20873
# contigs (>= 5000 bp)	1671
# contigs (>= 10000 bp)	755
# contigs (>= 25000 bp)	282
# contigs (>= 50000 bp)	124
Total length (>= 0 bp)	142136995
Total length (>= 1000 bp)	68609530
Total length (>= 5000 bp)	35095538
Total length (>= 10000 bp)	28793530
Total length (>= 25000 bp)	21492055
Total length (>= 50000 bp)	16196371
# contigs	60304
Largest contig	1176383
Total length	95545825
GC (%)	47.11
N50	2146
N90	627
auN	62739.1
L50	5849
L90	43182
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	49 + 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).

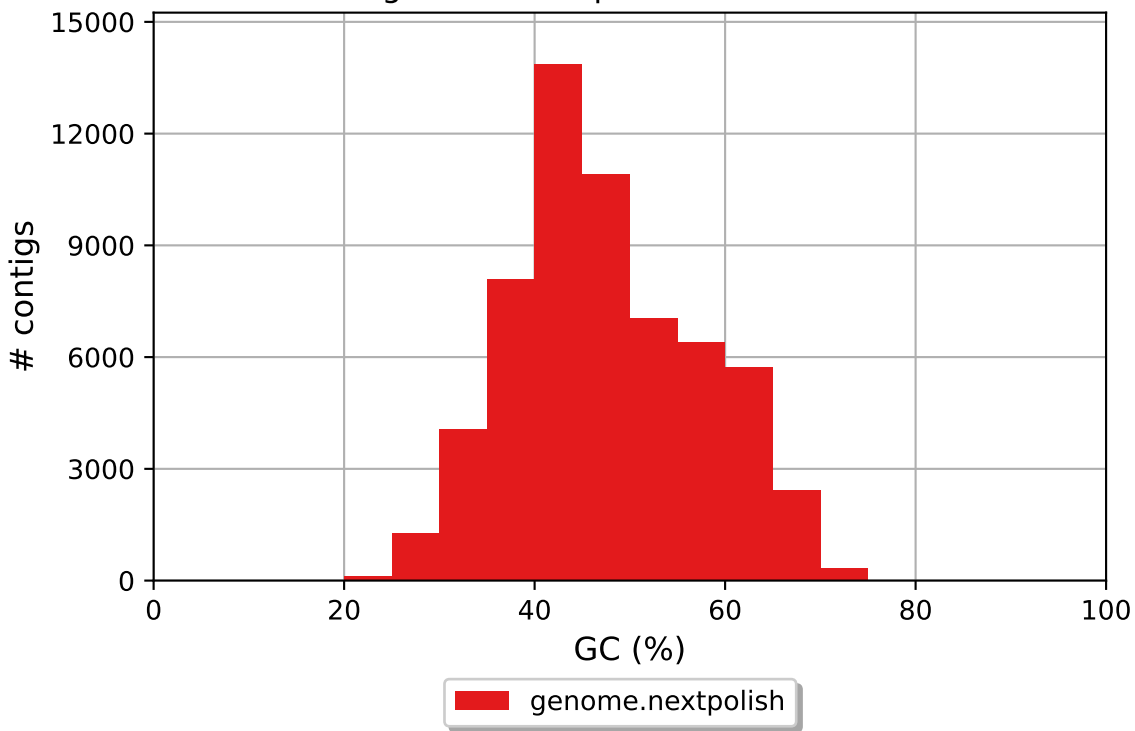




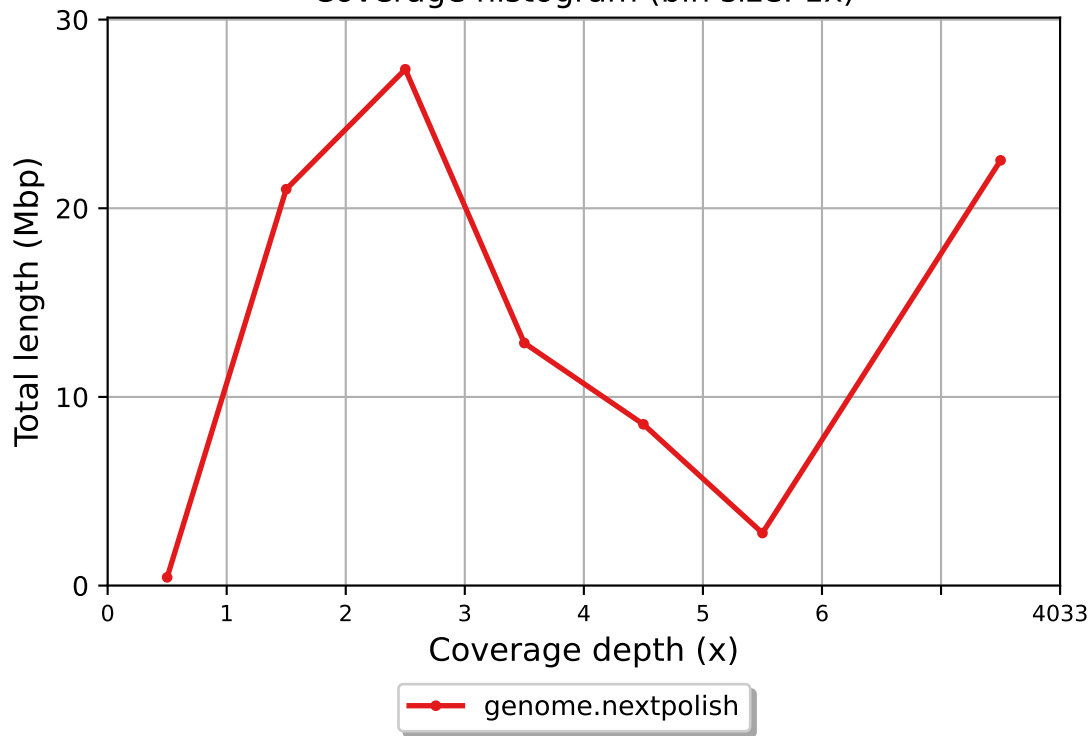
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 1x)



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

