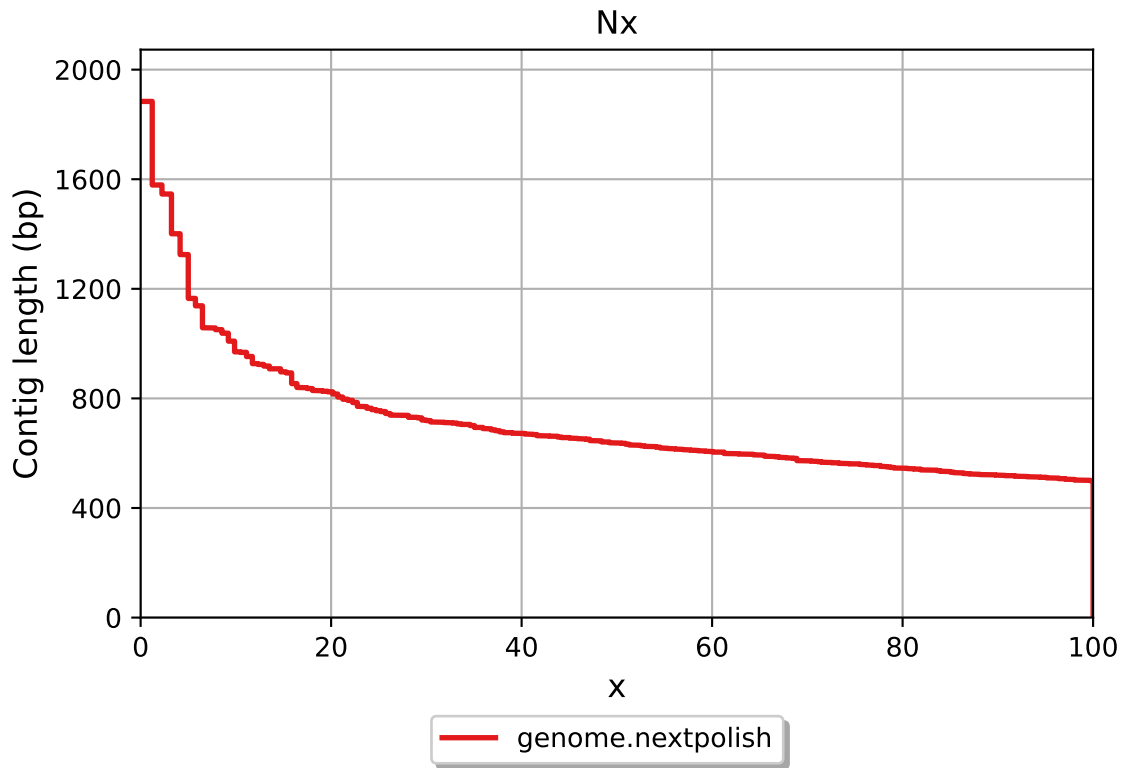
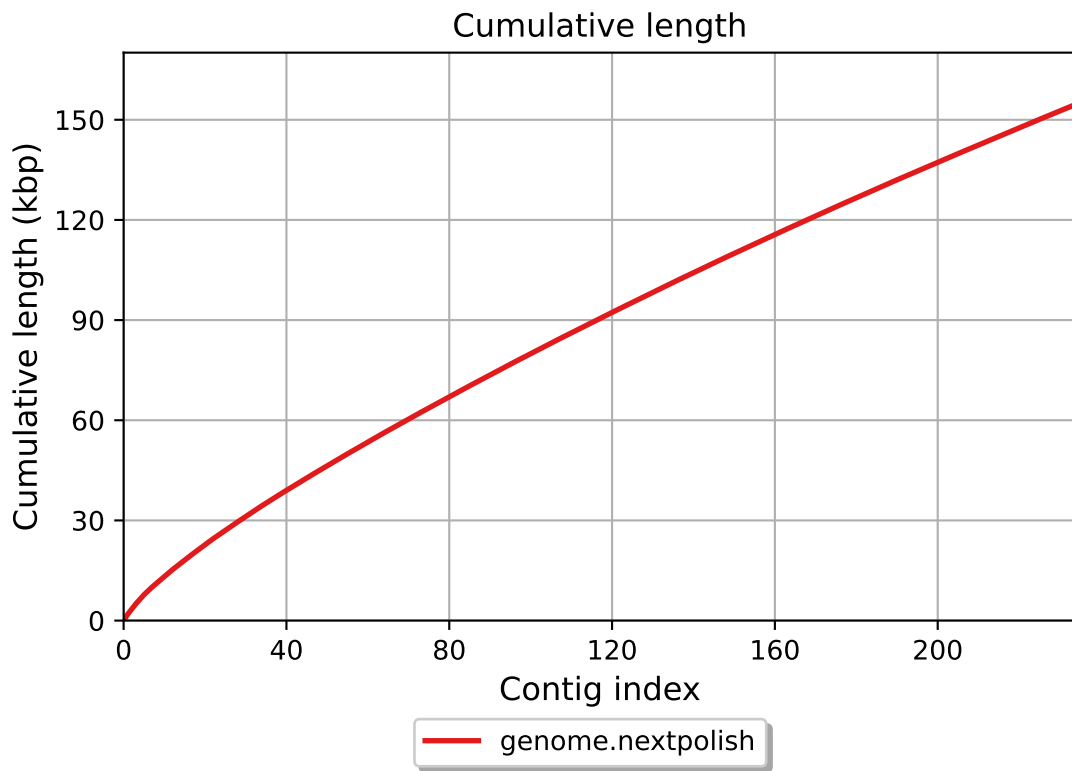


Report

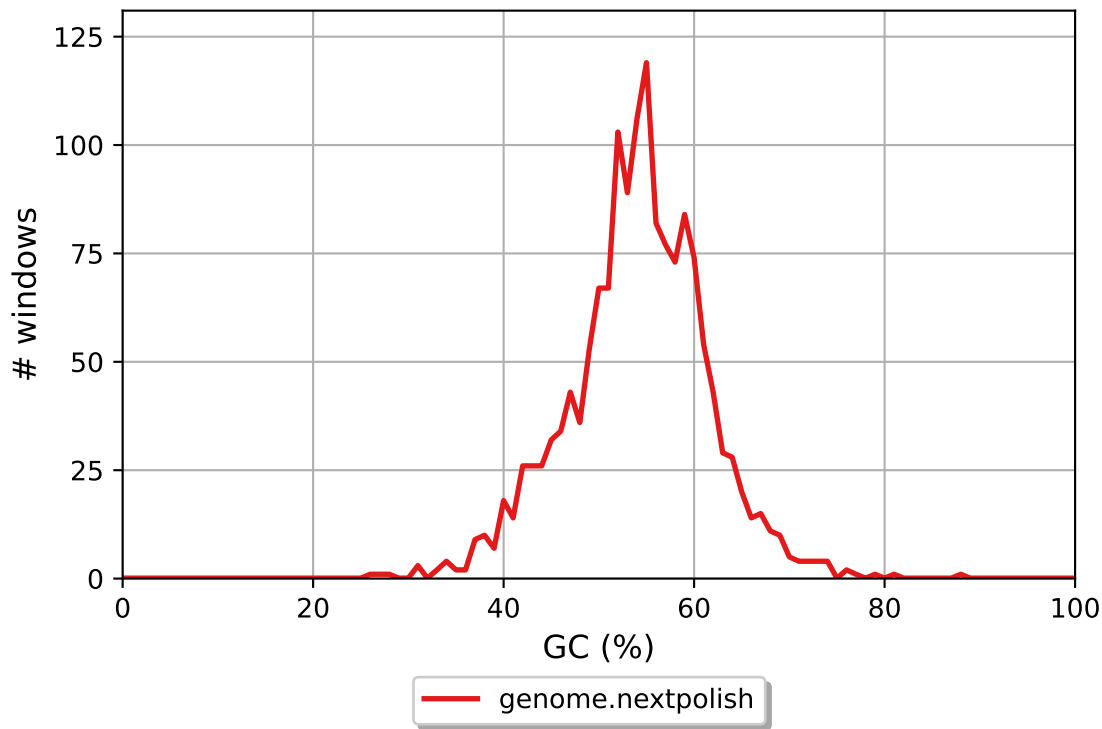
	genome.nextpolish
# contigs (>= 0 bp)	6465
# contigs (>= 1000 bp)	12
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1310097
Total length (>= 1000 bp)	15251
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	234
Largest contig	1884
Total length	154641
GC (%)	54.16
N50	637
N90	520
auN	713.6
L50	96
L90	204
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 1 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).

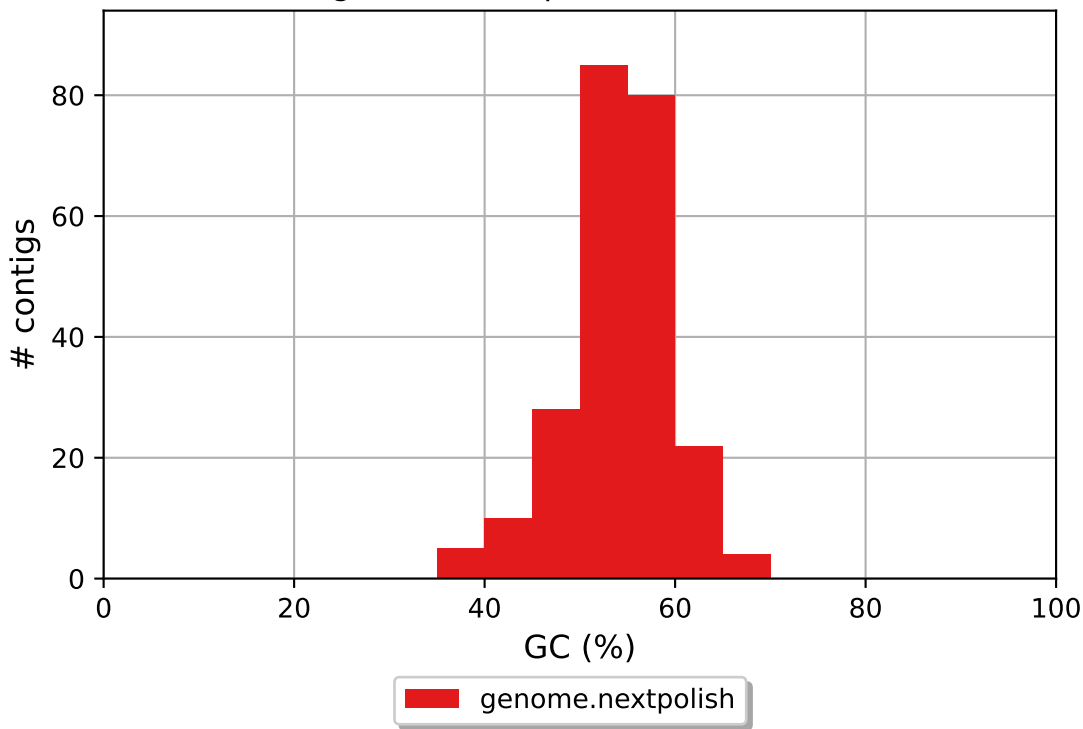




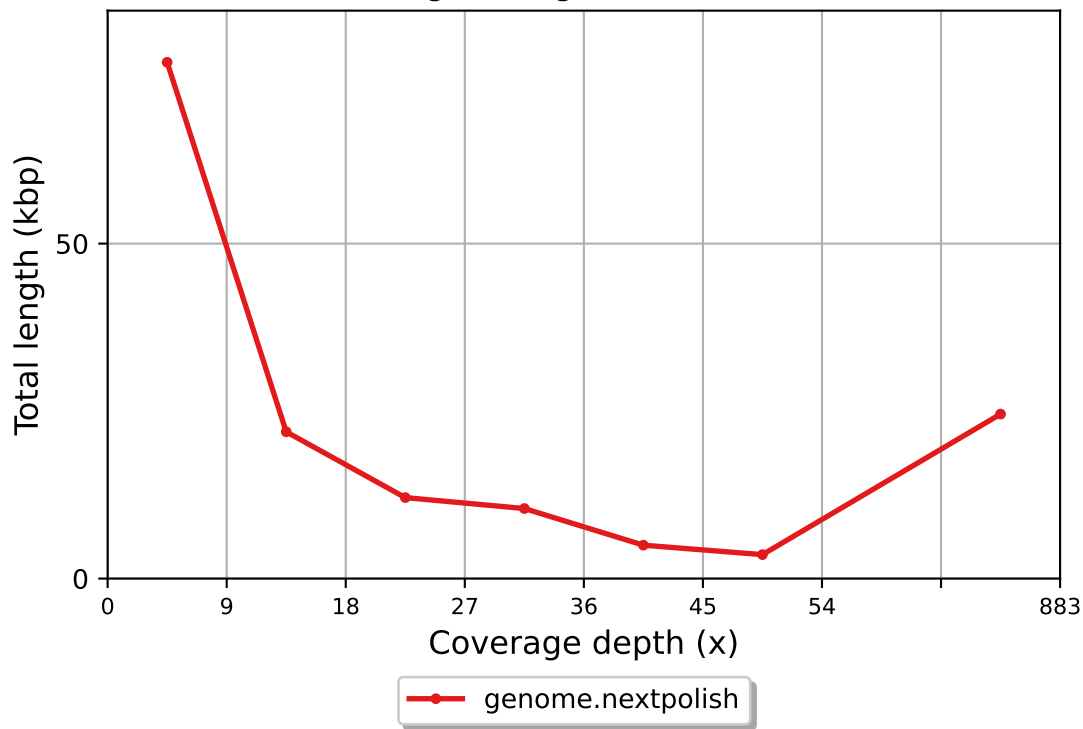
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 9x)



genome.nextpolish coverage histogram (bin size: 9x)

