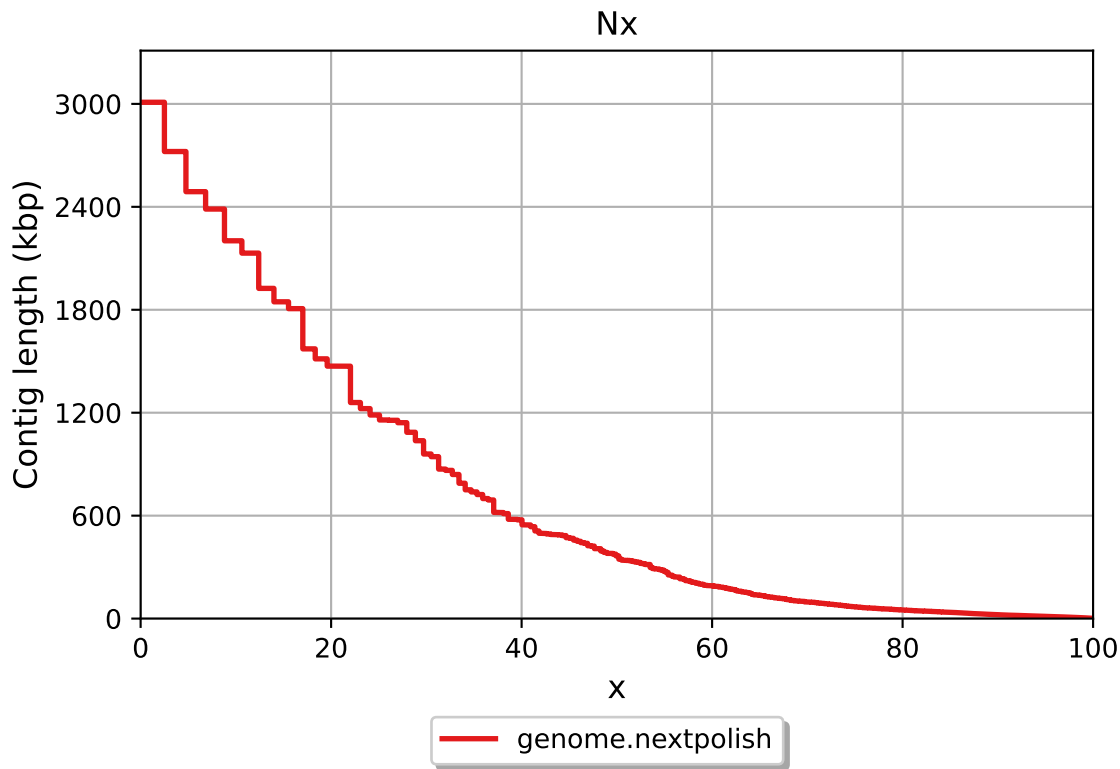
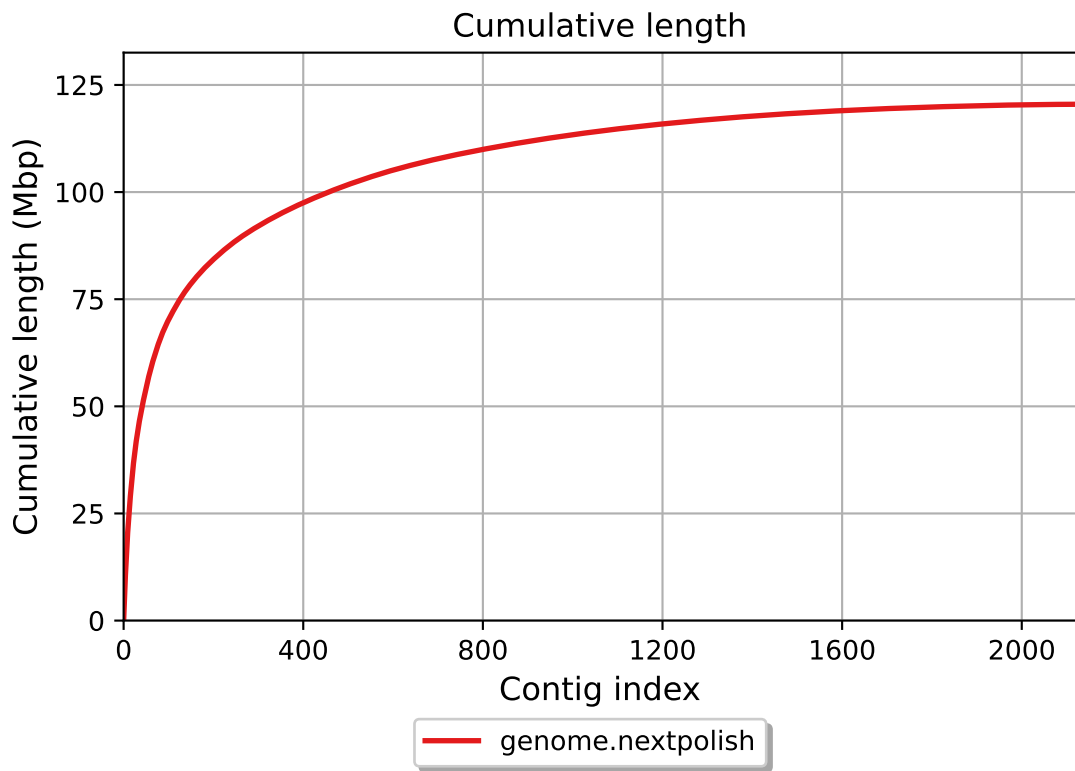


Report

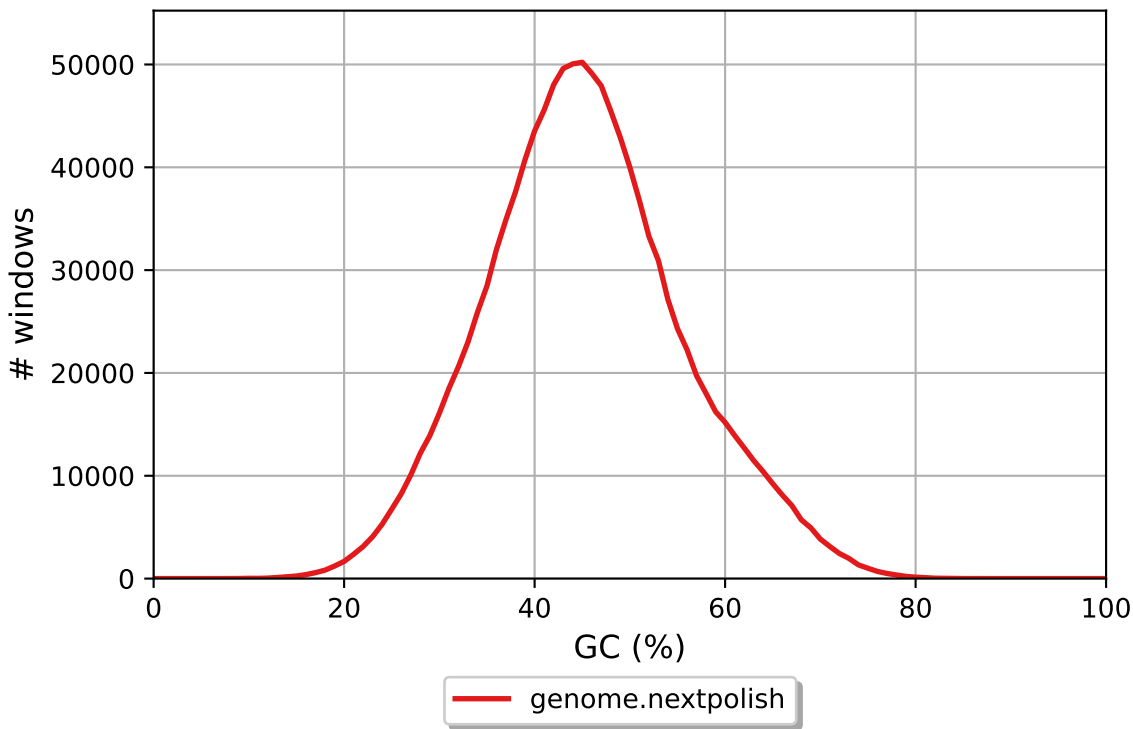
	genome.nextpolish
# contigs (>= 0 bp)	2122
# contigs (>= 1000 bp)	2072
# contigs (>= 5000 bp)	1623
# contigs (>= 10000 bp)	1240
# contigs (>= 25000 bp)	679
# contigs (>= 50000 bp)	381
Total length (>= 0 bp)	120492381
Total length (>= 1000 bp)	120458930
Total length (>= 5000 bp)	119125851
Total length (>= 10000 bp)	116314515
Total length (>= 25000 bp)	107280445
Total length (>= 50000 bp)	96590850
# contigs	2121
Largest contig	3009474
Total length	120491884
GC (%)	45.24
N50	365384
N90	22520
auN	740835.4
L50	65
L90	728
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	321 + 13 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

