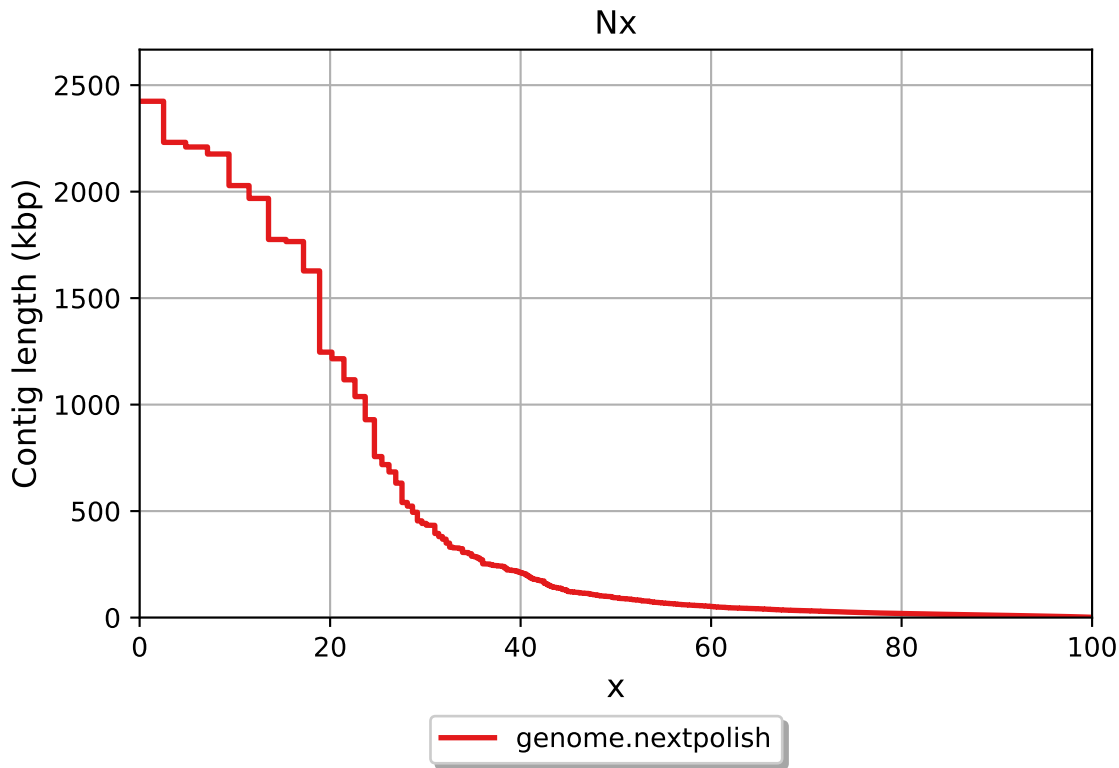
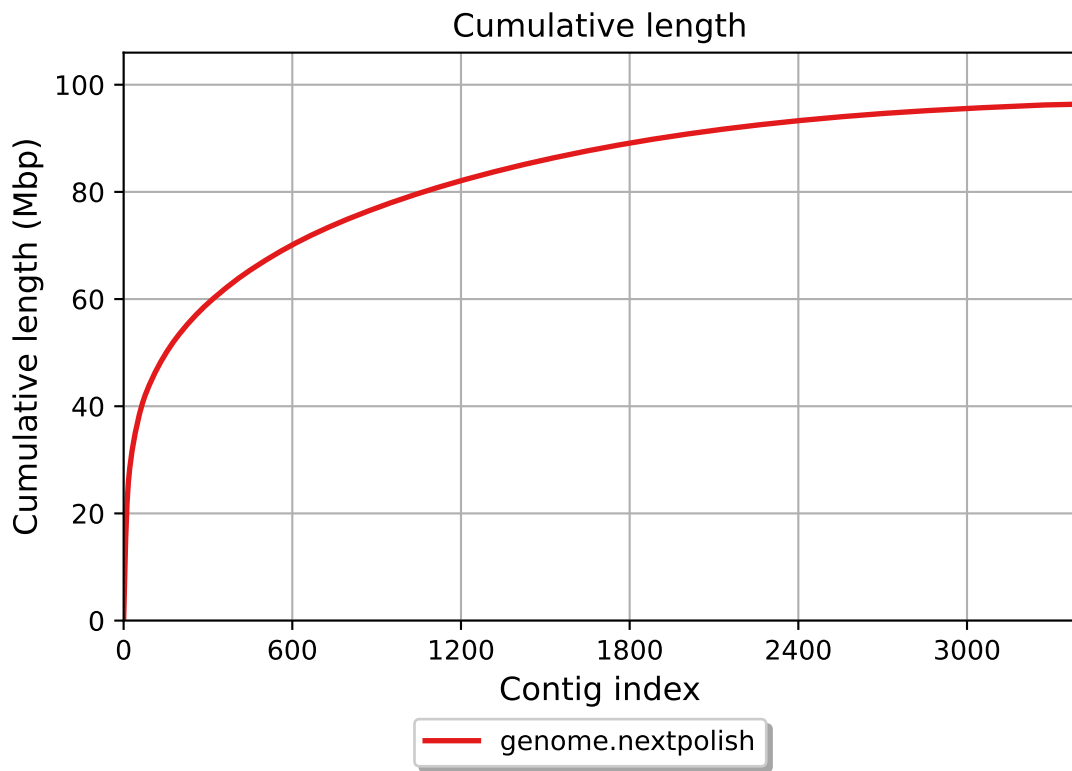


Report

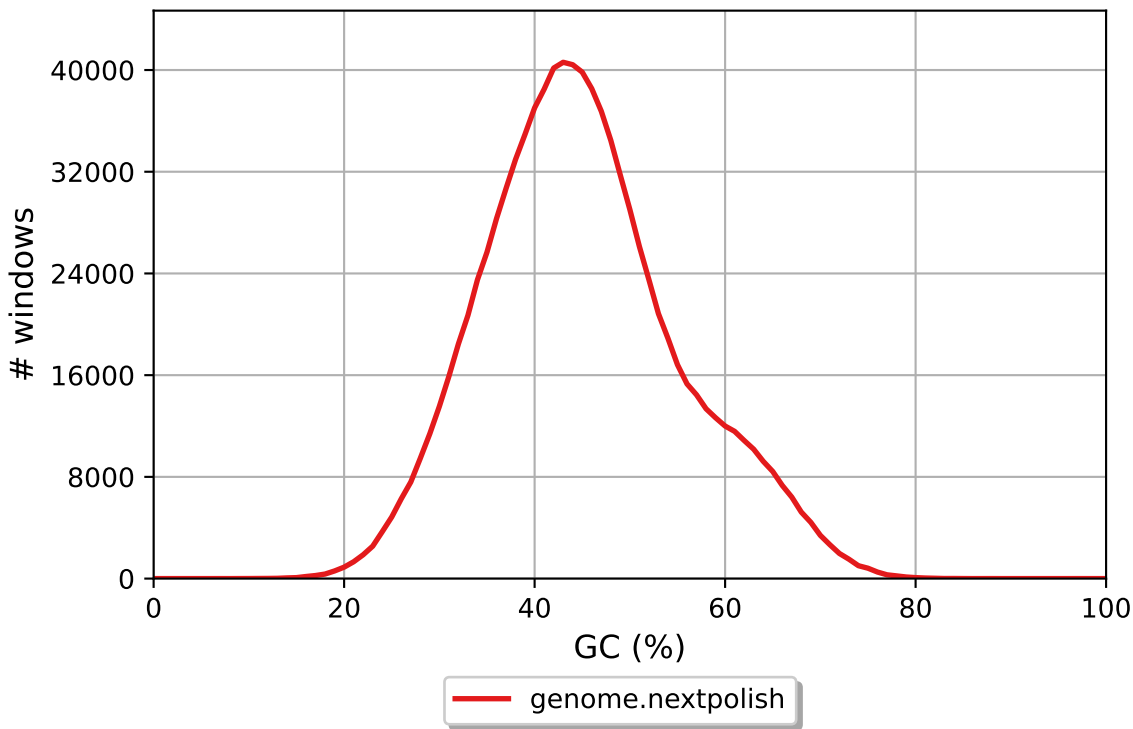
	genome.nextpolish
# contigs (>= 0 bp)	3392
# contigs (>= 1000 bp)	3349
# contigs (>= 5000 bp)	2436
# contigs (>= 10000 bp)	1686
# contigs (>= 25000 bp)	674
# contigs (>= 50000 bp)	283
Total length (>= 0 bp)	96354481
Total length (>= 1000 bp)	96325232
Total length (>= 5000 bp)	93472011
Total length (>= 10000 bp)	88026146
Total length (>= 25000 bp)	72080310
Total length (>= 50000 bp)	58388728
# contigs	3388
Largest contig	2424310
Total length	96353215
GC (%)	45.03
N50	93330
N90	11079
auN	544553.6
L50	131
L90	1562
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	244 + 15 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

