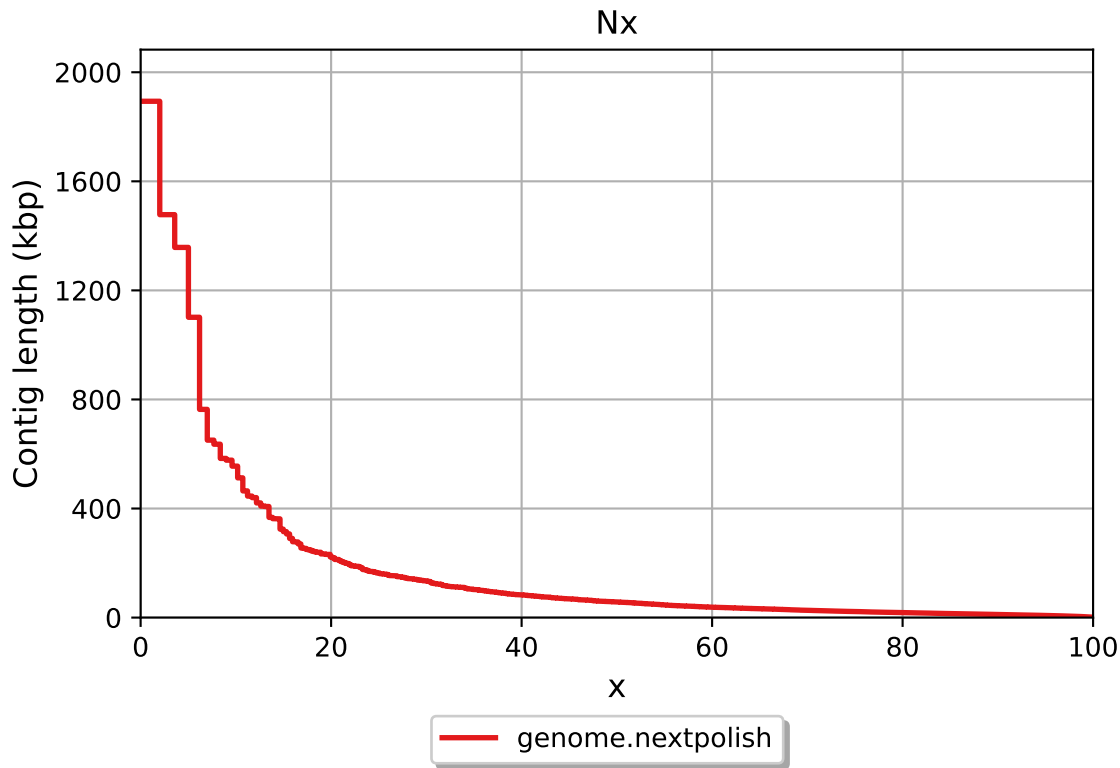
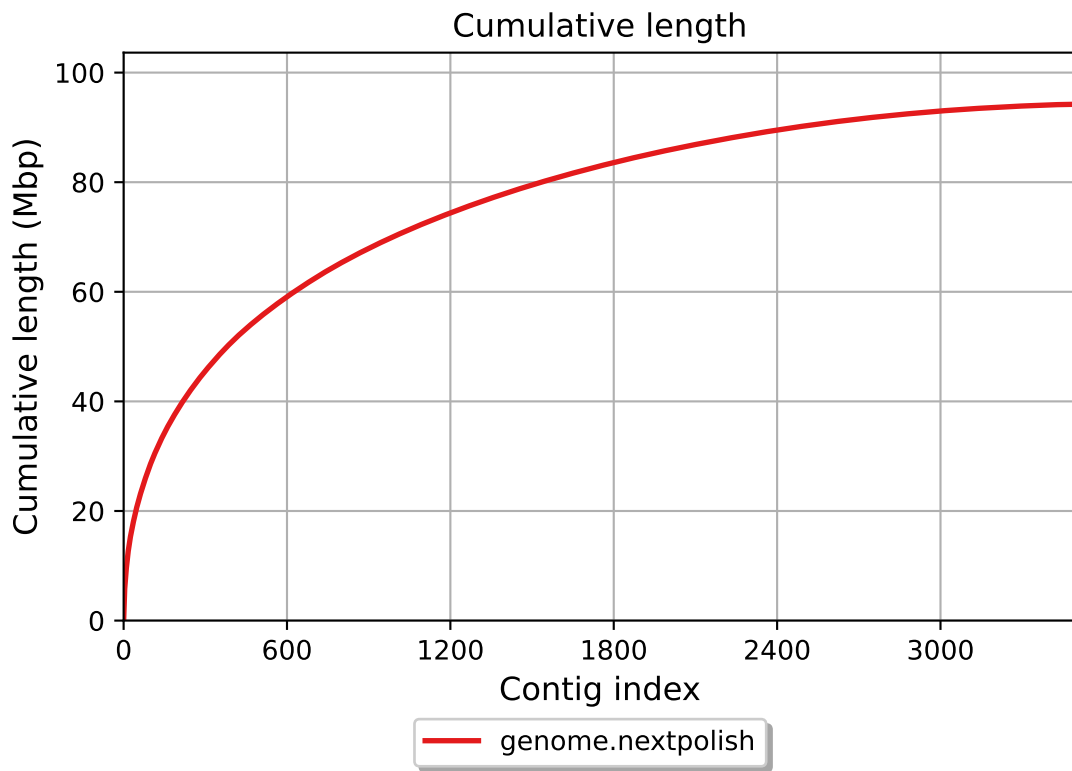


# Report

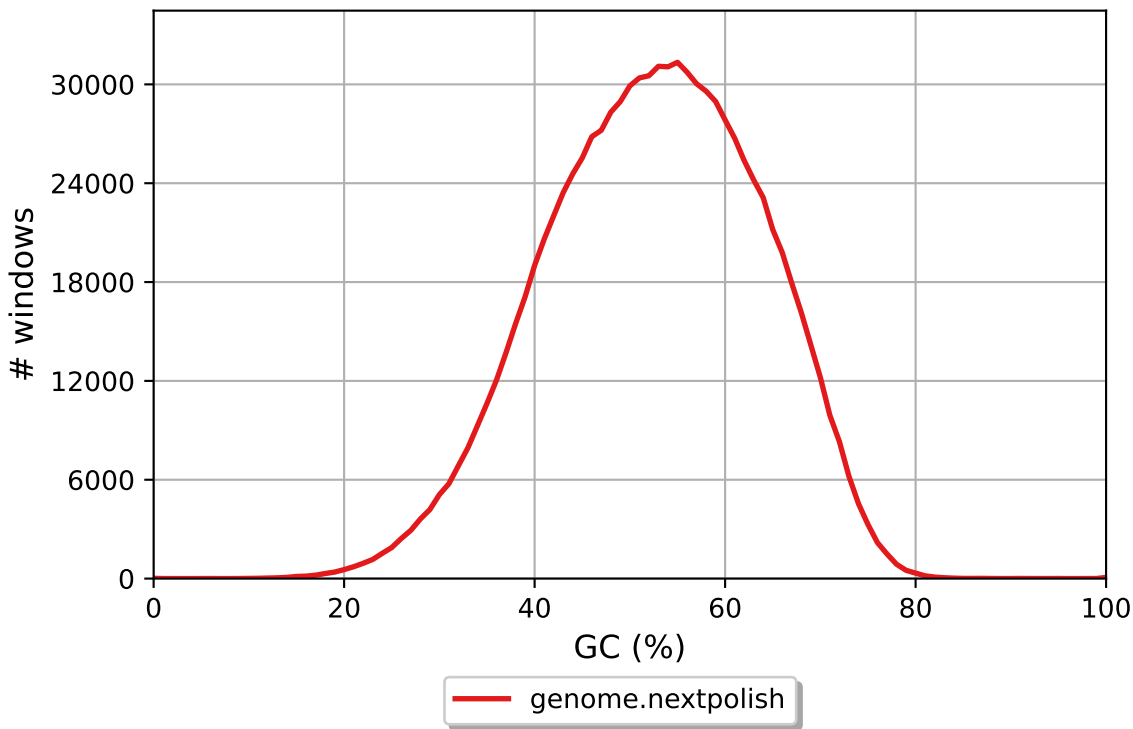
	genome.nextpolish
# contigs (>= 0 bp)	3498
# contigs (>= 1000 bp)	3465
# contigs (>= 5000 bp)	2815
# contigs (>= 10000 bp)	2067
# contigs (>= 25000 bp)	889
# contigs (>= 50000 bp)	387
Total length (>= 0 bp)	94220447
Total length (>= 1000 bp)	94196390
Total length (>= 5000 bp)	92164490
Total length (>= 10000 bp)	86548519
Total length (>= 25000 bp)	67645439
Total length (>= 50000 bp)	50357897
# contigs	3498
Largest contig	1893588
Total length	94220447
GC (%)	52.36
N50	57503
N90	11339
auN	198699.5
L50	327
L90	1903
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	114 + 8 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

