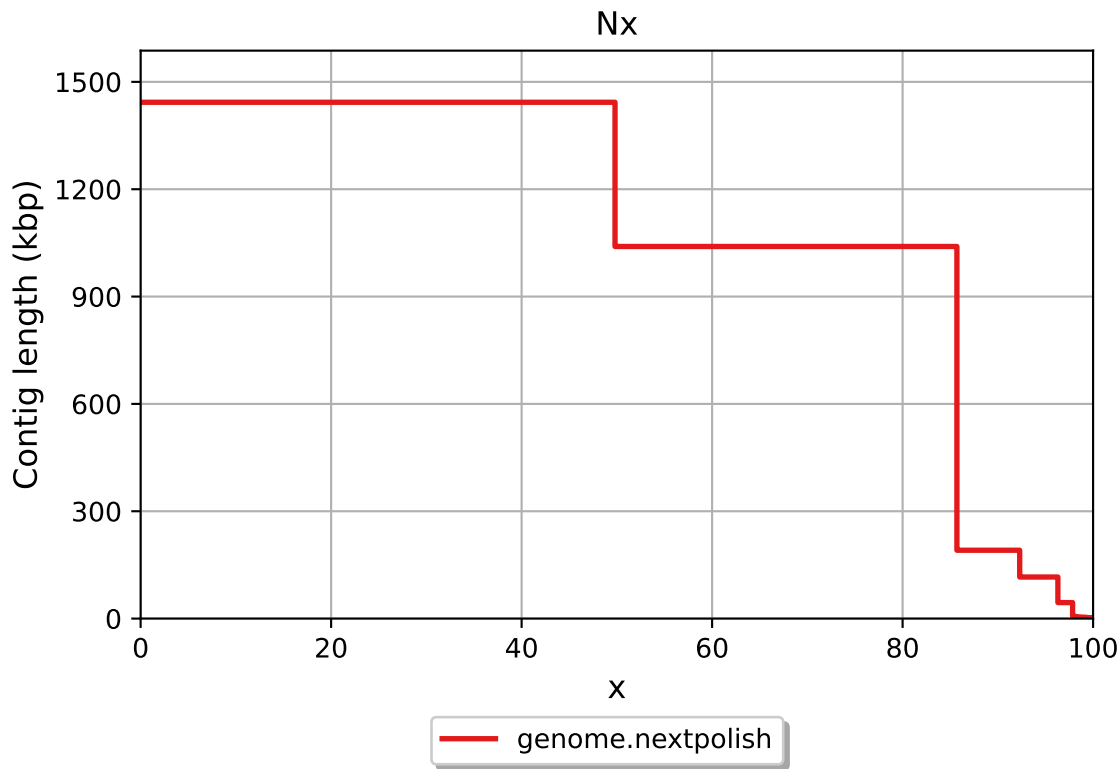
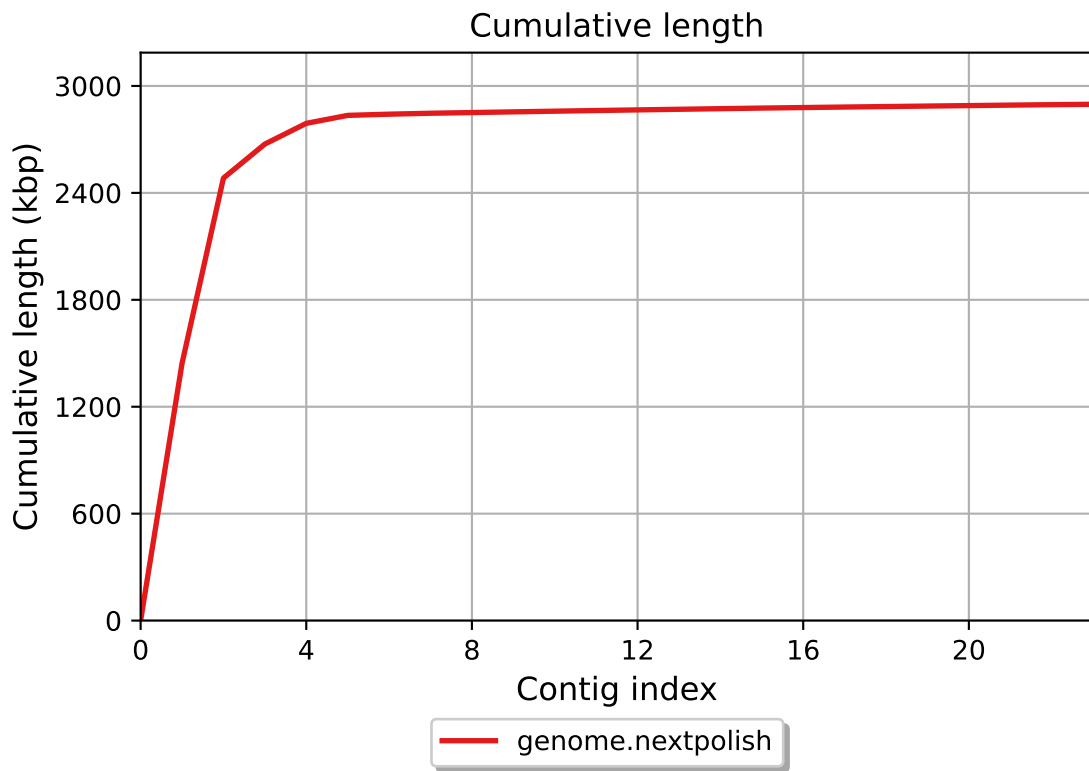


Report

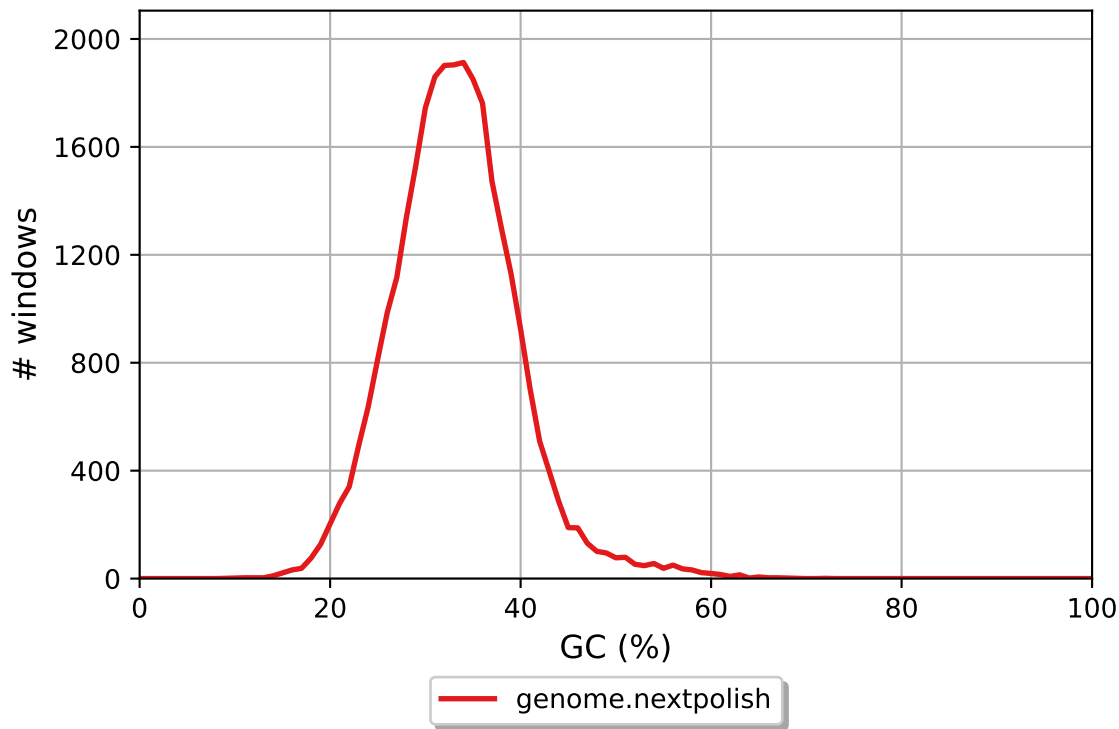
	genome.nextpolish
# contigs (>= 0 bp)	23
# contigs (>= 1000 bp)	23
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2897242
Total length (>= 1000 bp)	2897242
Total length (>= 5000 bp)	2846822
Total length (>= 10000 bp)	2834891
Total length (>= 25000 bp)	2834891
Total length (>= 50000 bp)	2790162
# contigs	23
Largest contig	1443005
Total length	2897242
GC (%)	33.23
N50	1039965
N90	191008
auN	1110024.6
L50	2
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 6 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

