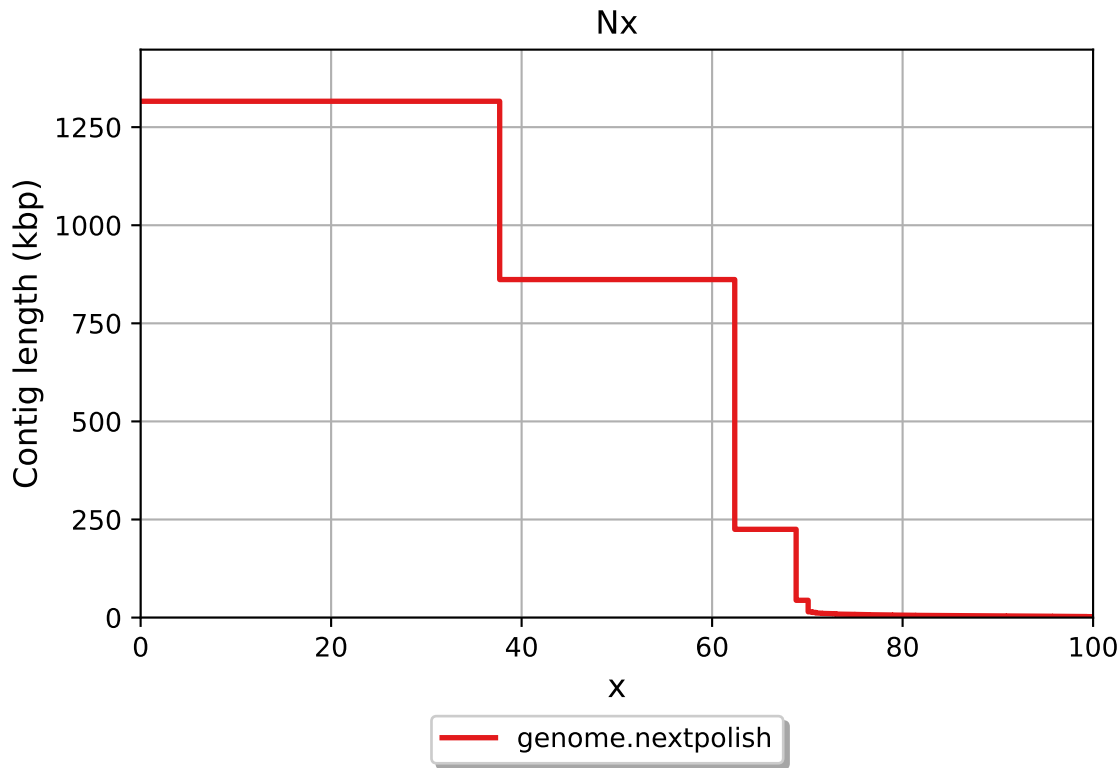
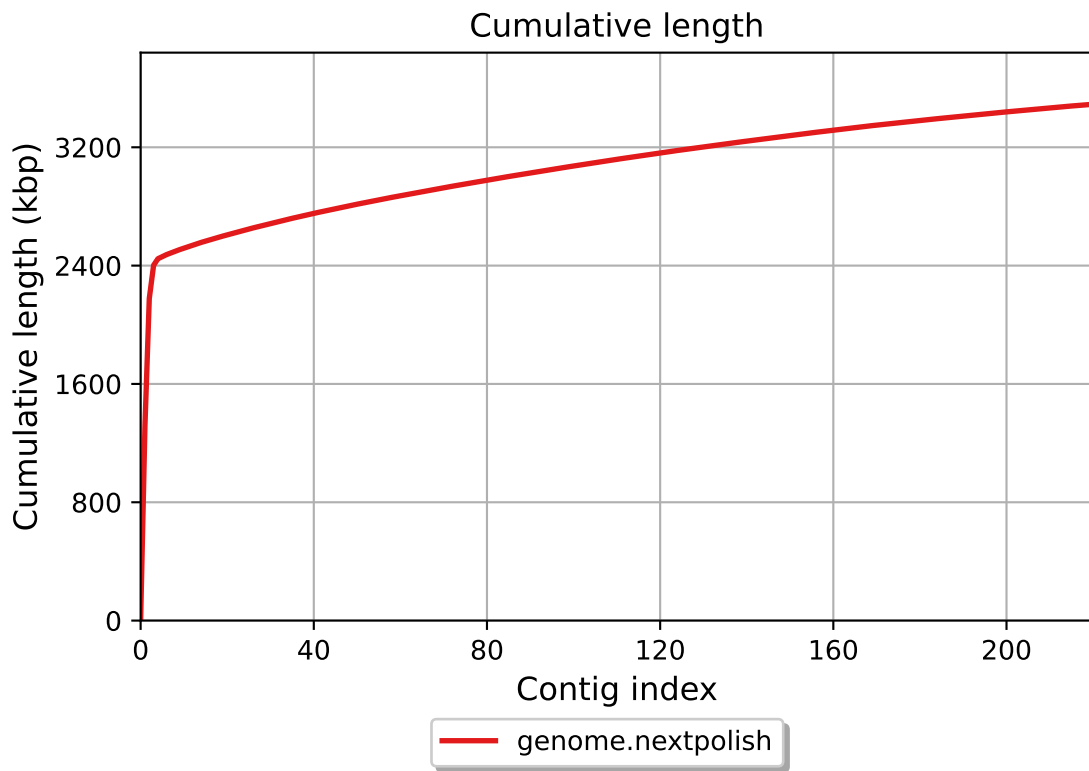


Report

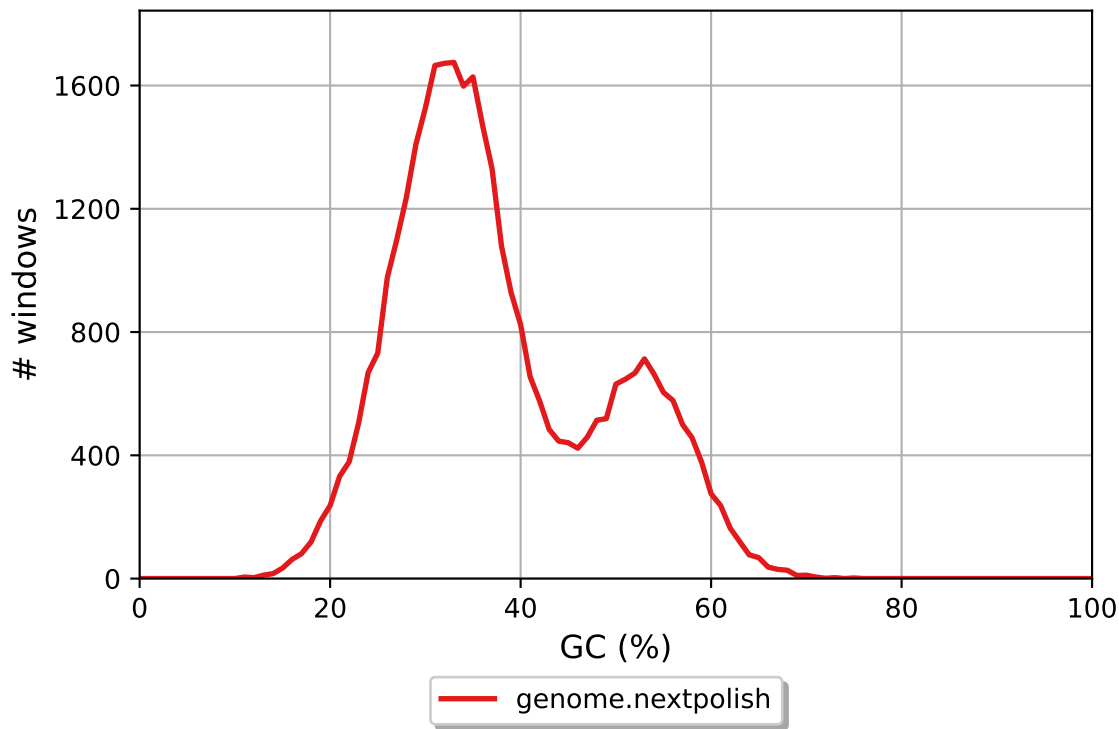
	genome.nextpolish
# contigs (>= 0 bp)	220
# contigs (>= 1000 bp)	220
# contigs (>= 5000 bp)	80
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	3490817
Total length (>= 1000 bp)	3490817
Total length (>= 5000 bp)	2977566
Total length (>= 10000 bp)	2528200
Total length (>= 25000 bp)	2446525
Total length (>= 50000 bp)	2402459
# contigs	220
Largest contig	1315962
Total length	3490817
GC (%)	37.81
N50	861519
N90	4292
auN	725473.4
L50	2
L90	116
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 2 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

