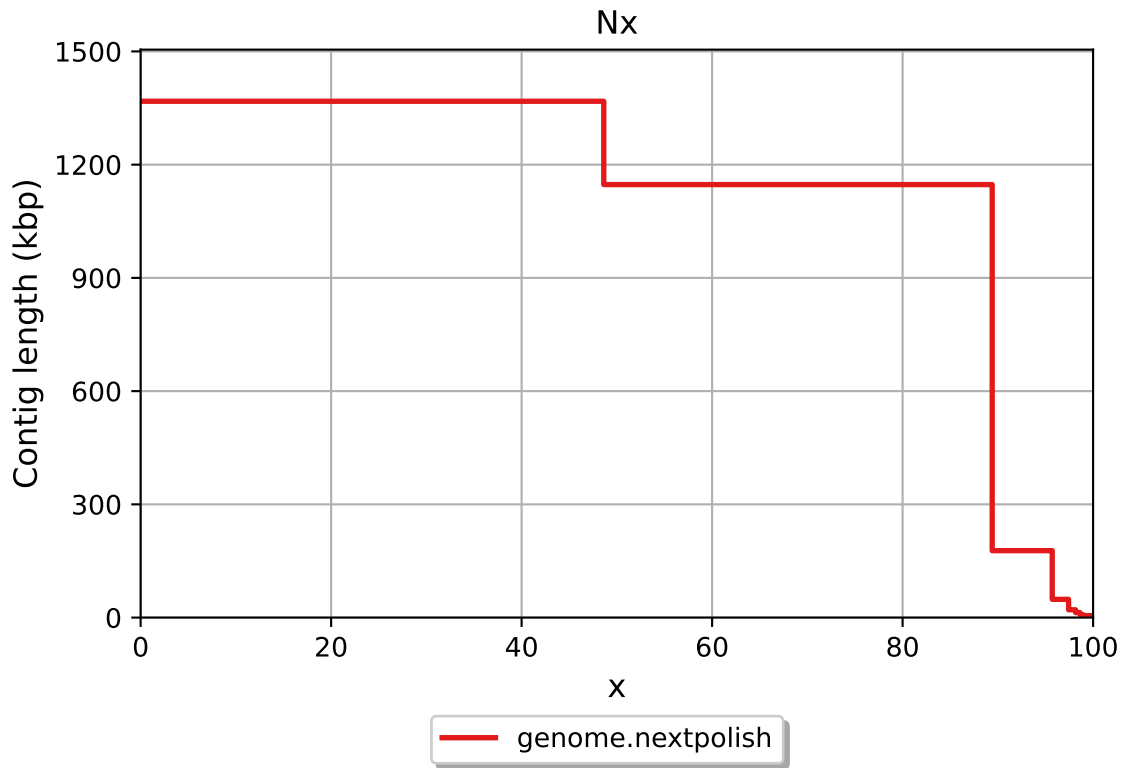
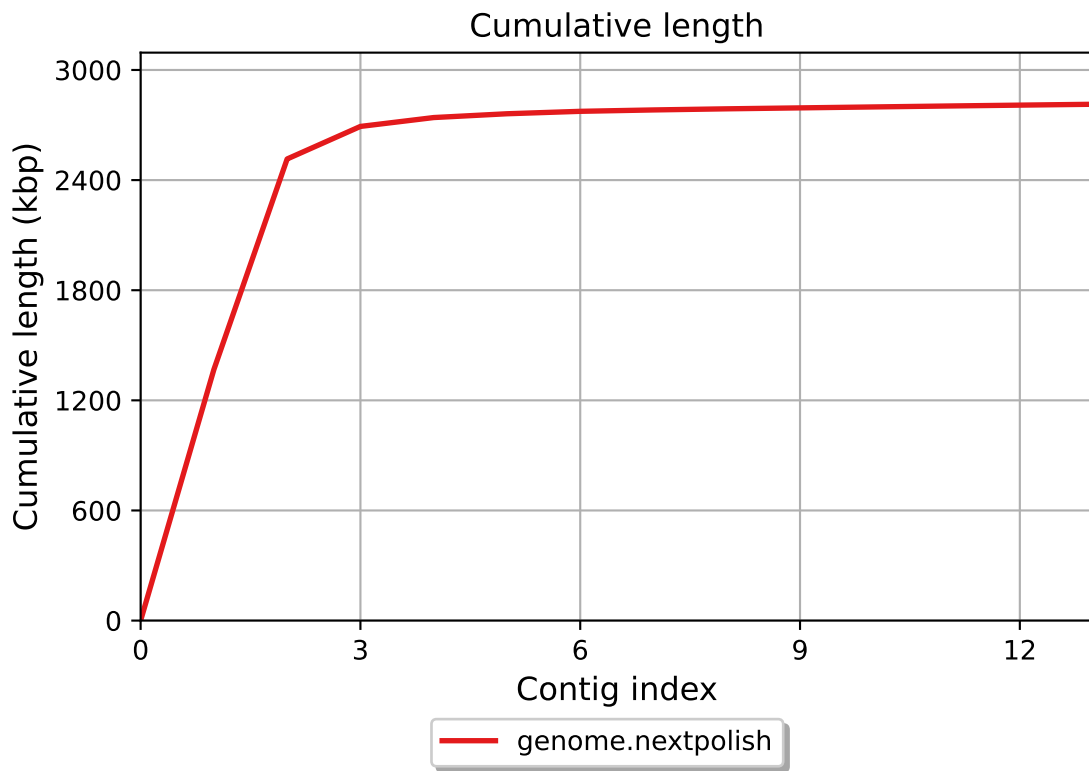


# Report

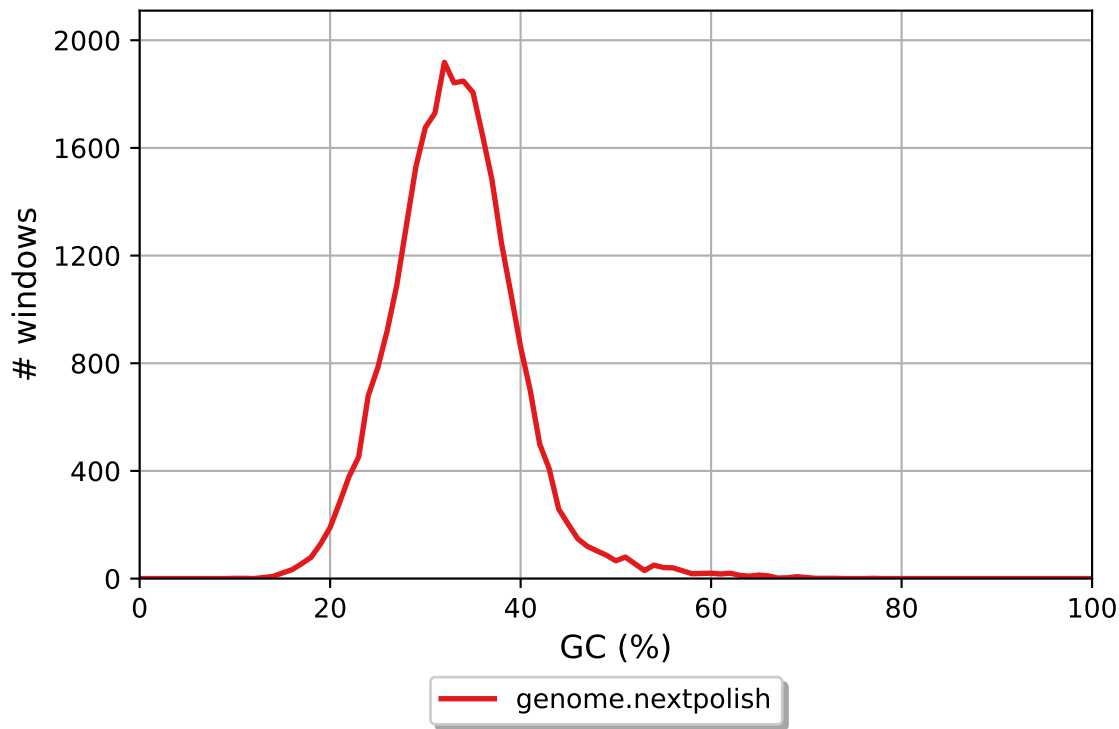
	genome.nextpolish
# contigs (>= 0 bp)	13
# contigs (>= 1000 bp)	13
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	2813137
Total length (>= 1000 bp)	2813137
Total length (>= 5000 bp)	2803617
Total length (>= 10000 bp)	2774695
Total length (>= 25000 bp)	2740747
Total length (>= 50000 bp)	2692420
# contigs	13
Largest contig	1367935
Total length	2813137
GC (%)	33.19
N50	1147193
N90	177292
auN	1145300.4
L50	2
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 10 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

