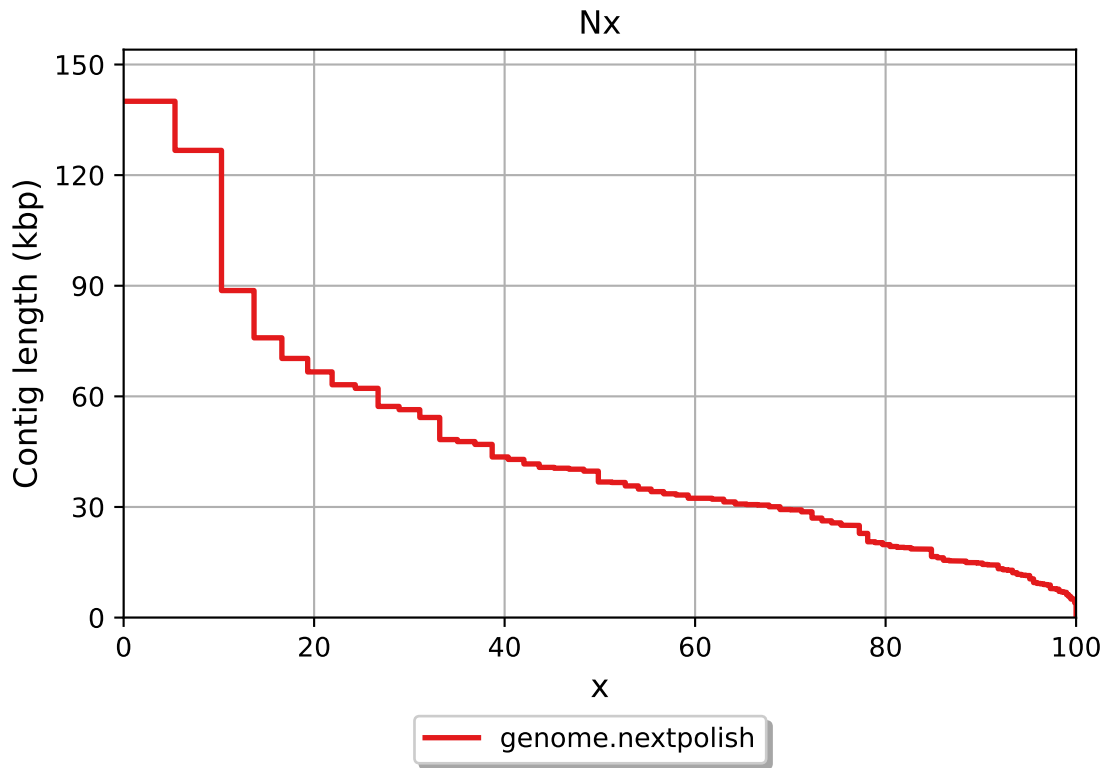
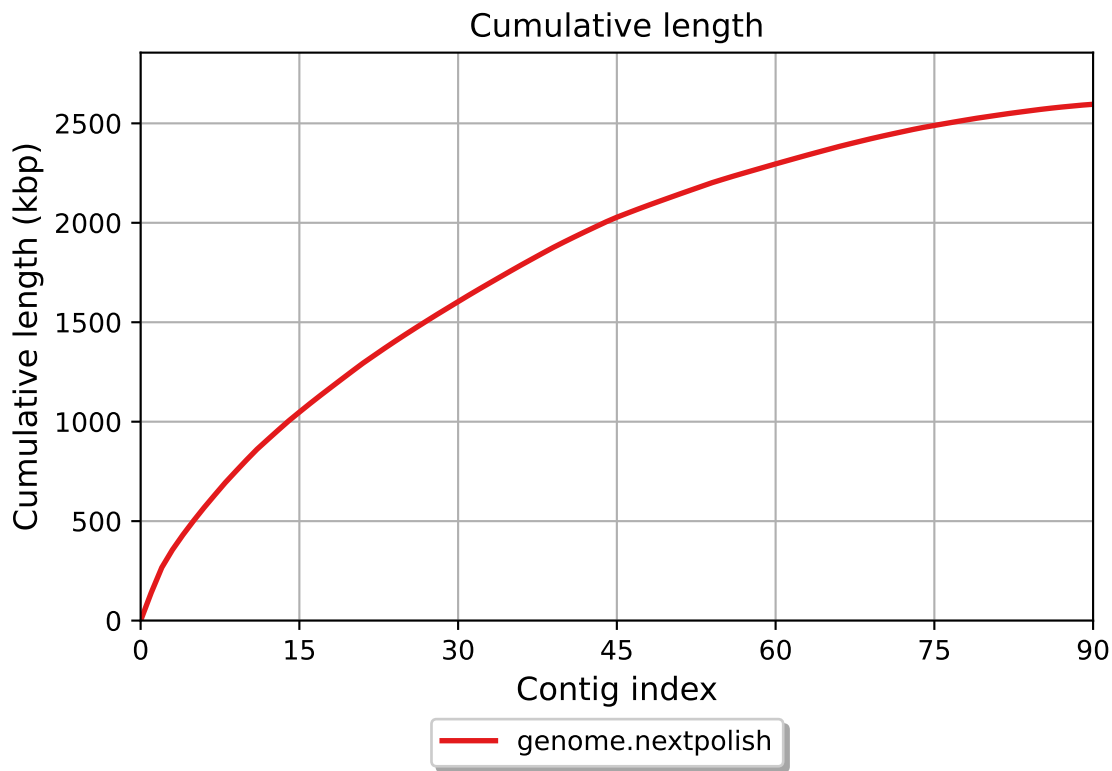


Report

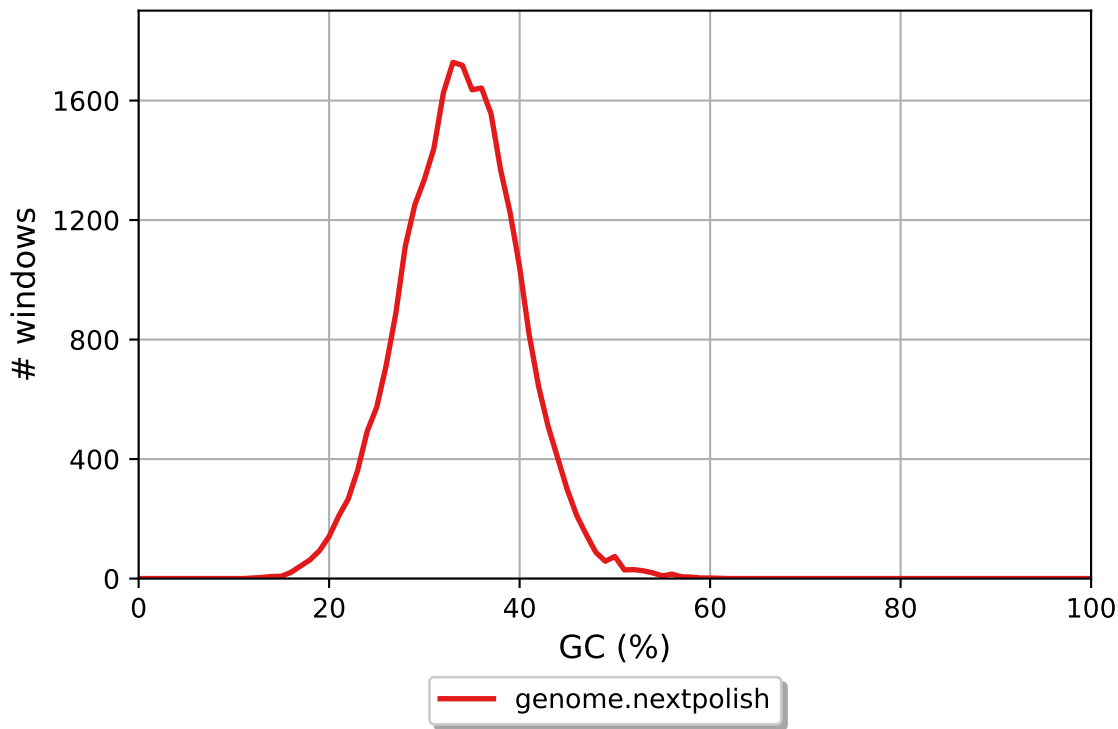
	genome.nextpolish
# contigs (>= 0 bp)	90
# contigs (>= 1000 bp)	90
# contigs (>= 5000 bp)	89
# contigs (>= 10000 bp)	74
# contigs (>= 25000 bp)	43
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	2595934
Total length (>= 1000 bp)	2595934
Total length (>= 5000 bp)	2591880
Total length (>= 10000 bp)	2479983
Total length (>= 25000 bp)	1980160
Total length (>= 50000 bp)	861493
# contigs	90
Largest contig	140032
Total length	2595934
GC (%)	33.82
N50	36776
N90	14787
auN	48417.5
L50	22
L90	63
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 0 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

