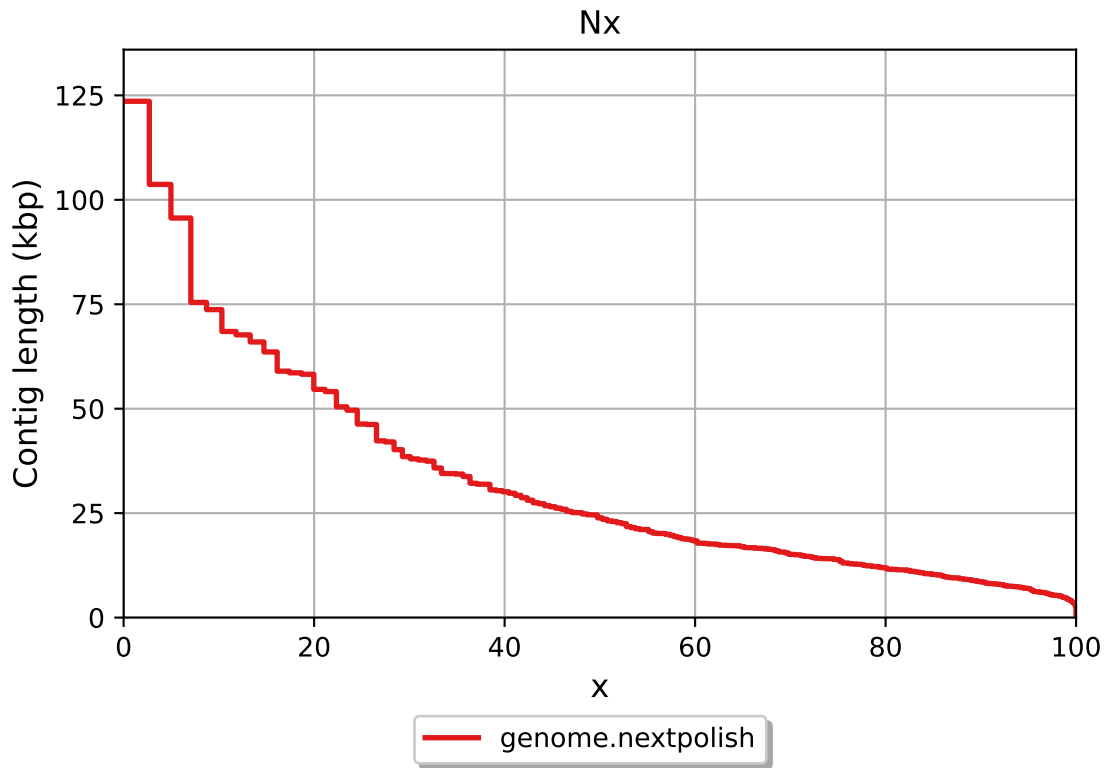
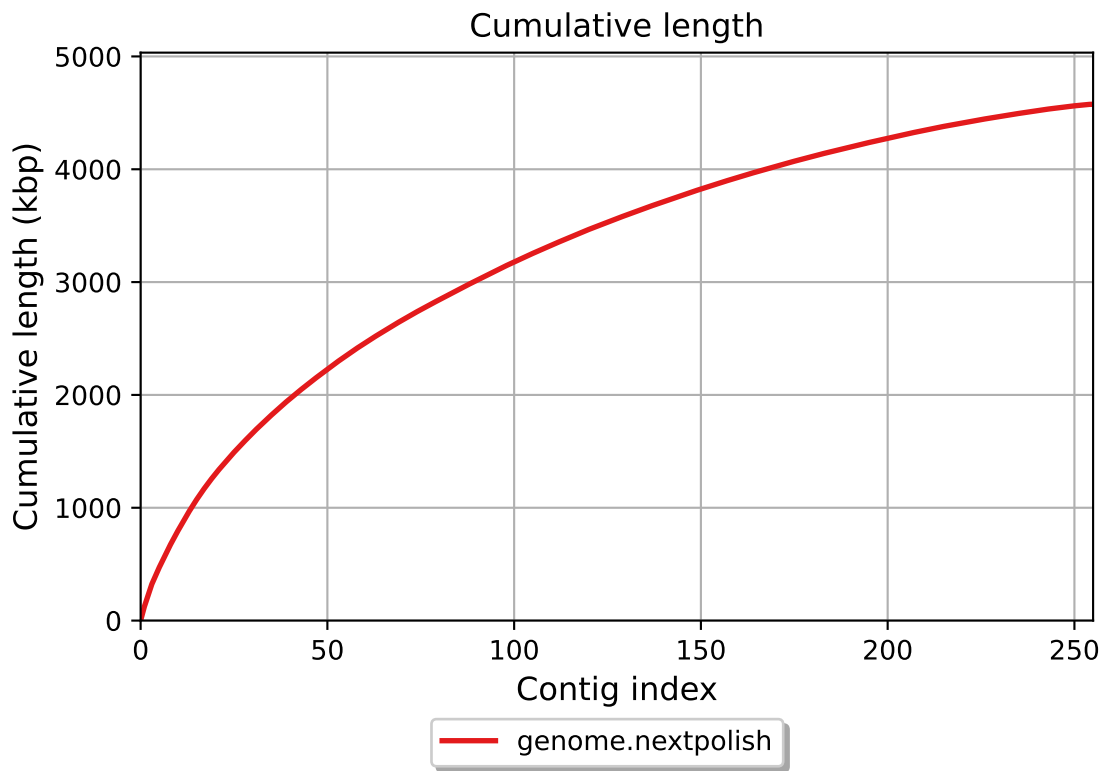


Report

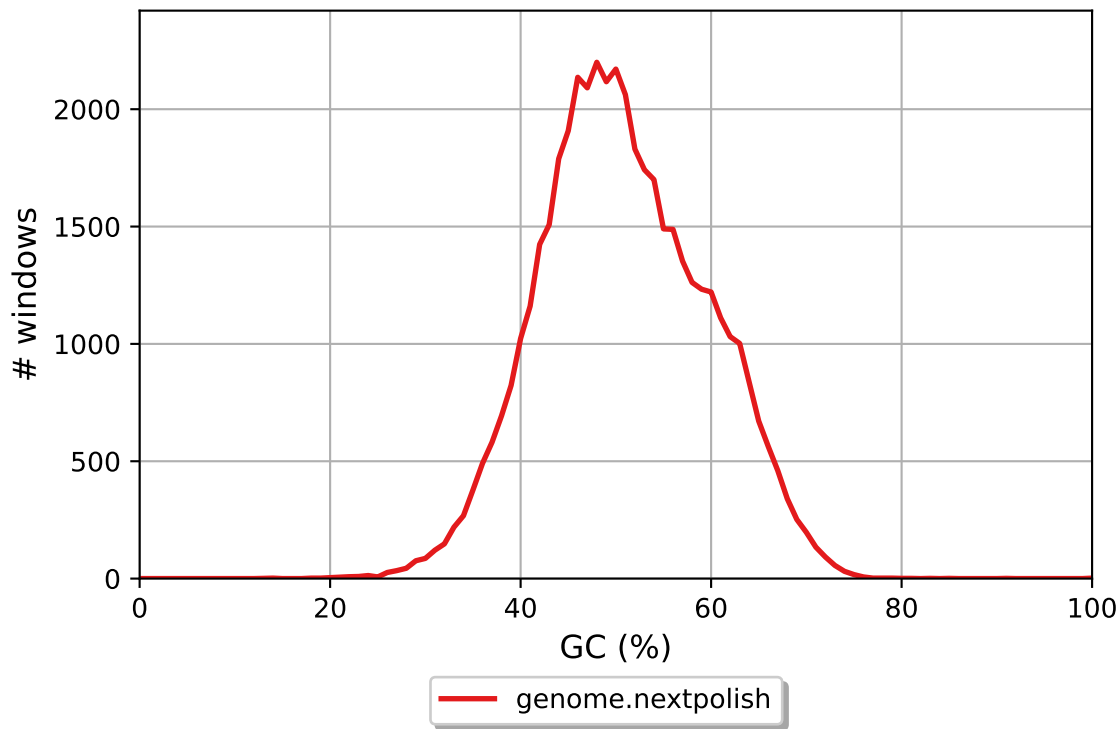
	genome.nextpolish
# contigs (>= 0 bp)	255
# contigs (>= 1000 bp)	254
# contigs (>= 5000 bp)	238
# contigs (>= 10000 bp)	160
# contigs (>= 25000 bp)	49
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	4576018
Total length (>= 1000 bp)	4575075
Total length (>= 5000 bp)	4509702
Total length (>= 10000 bp)	3929329
Total length (>= 25000 bp)	2202778
Total length (>= 50000 bp)	1072767
# contigs	255
Largest contig	123587
Total length	4576018
GC (%)	50.74
N50	23864
N90	8588
auN	33886.9
L50	53
L90	181
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 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

