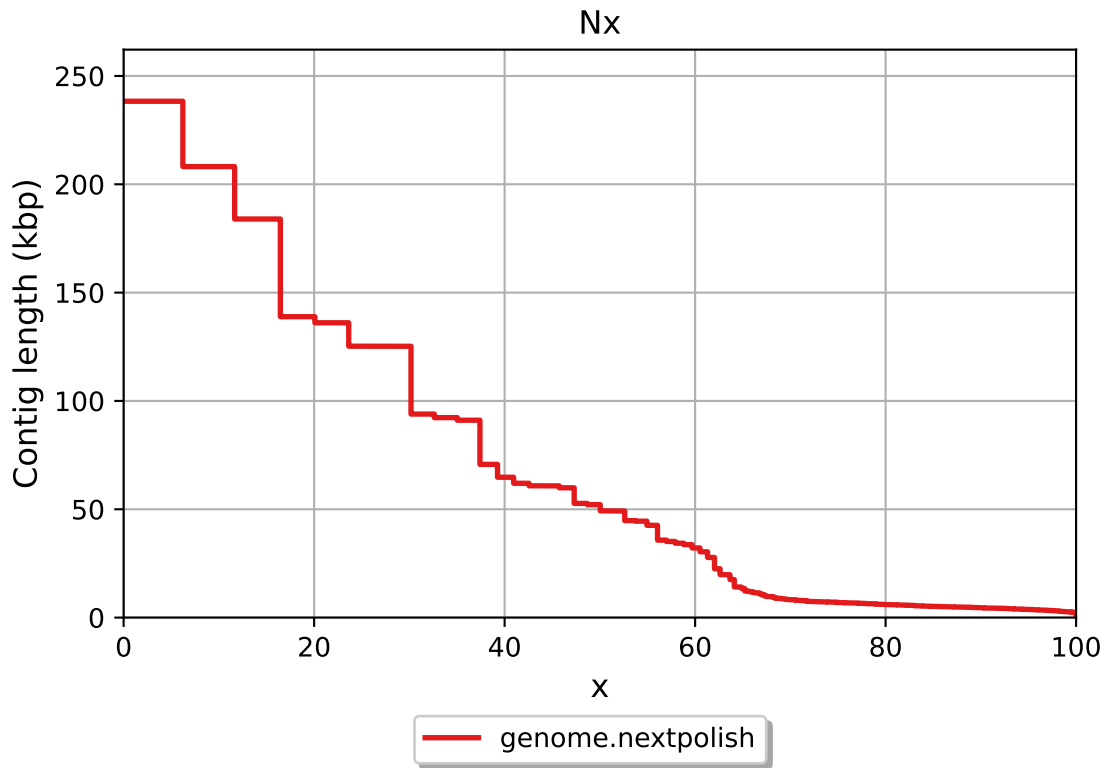
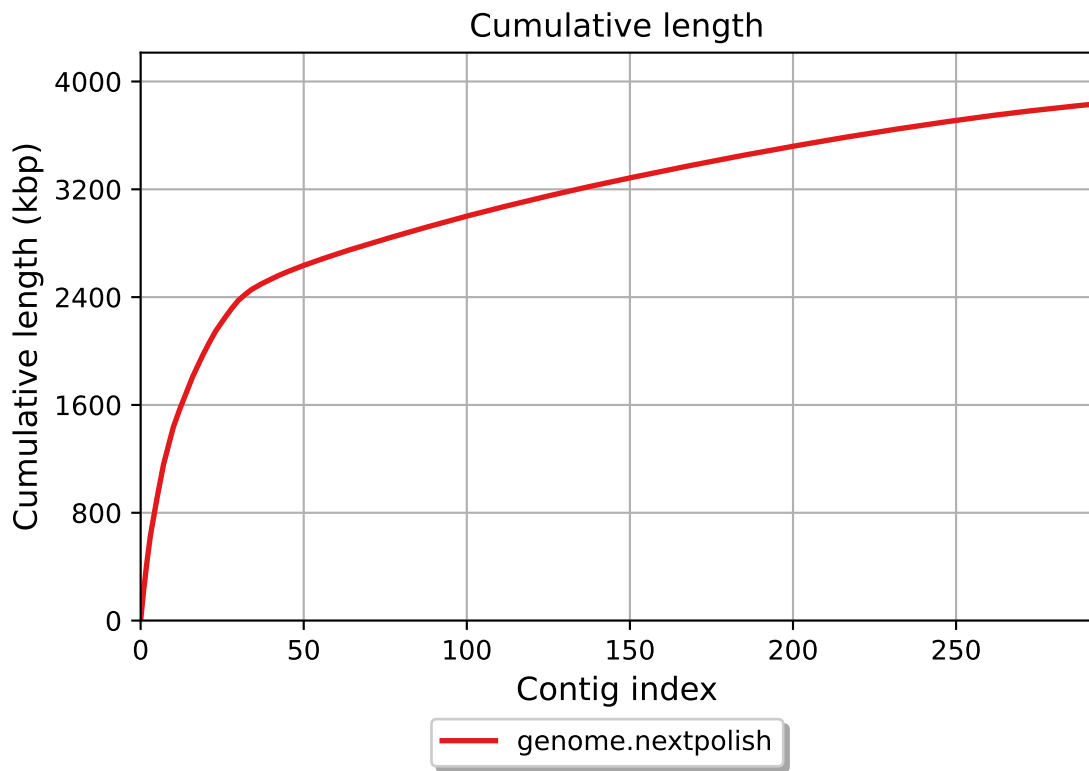


# Report

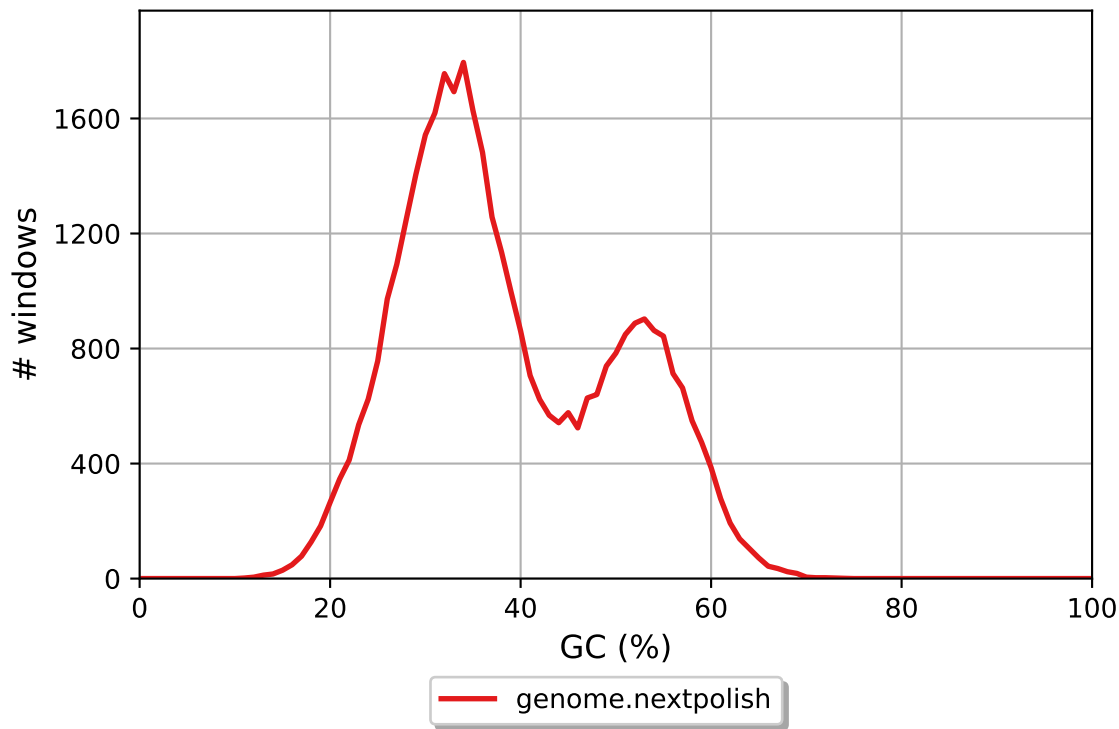
	genome.nextpolish
# contigs (>= 0 bp)	292
# contigs (>= 1000 bp)	292
# contigs (>= 5000 bp)	157
# contigs (>= 10000 bp)	44
# contigs (>= 25000 bp)	30
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	3831018
Total length (>= 1000 bp)	3831018
Total length (>= 5000 bp)	3319856
Total length (>= 10000 bp)	2578504
Total length (>= 25000 bp)	2376534
Total length (>= 50000 bp)	1917060
# contigs	292
Largest contig	238318
Total length	3831018
GC (%)	38.85
N50	52130
N90	4532
auN	74824.4
L50	18
L90	184
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 1 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

