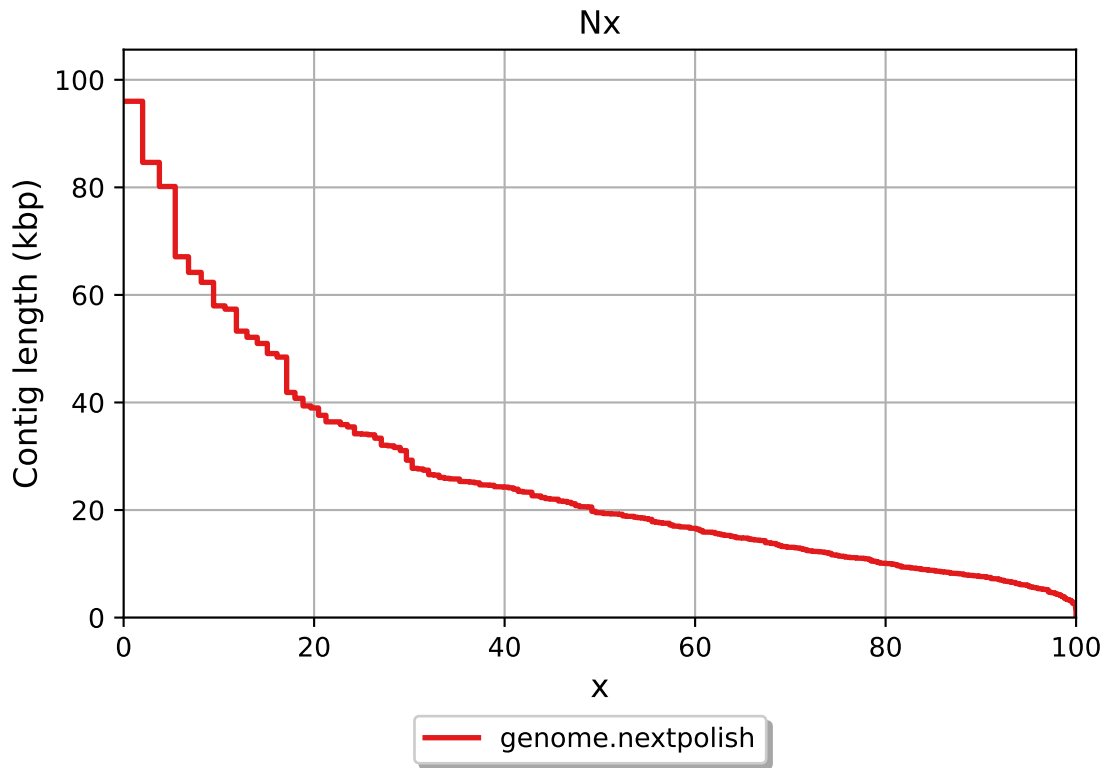
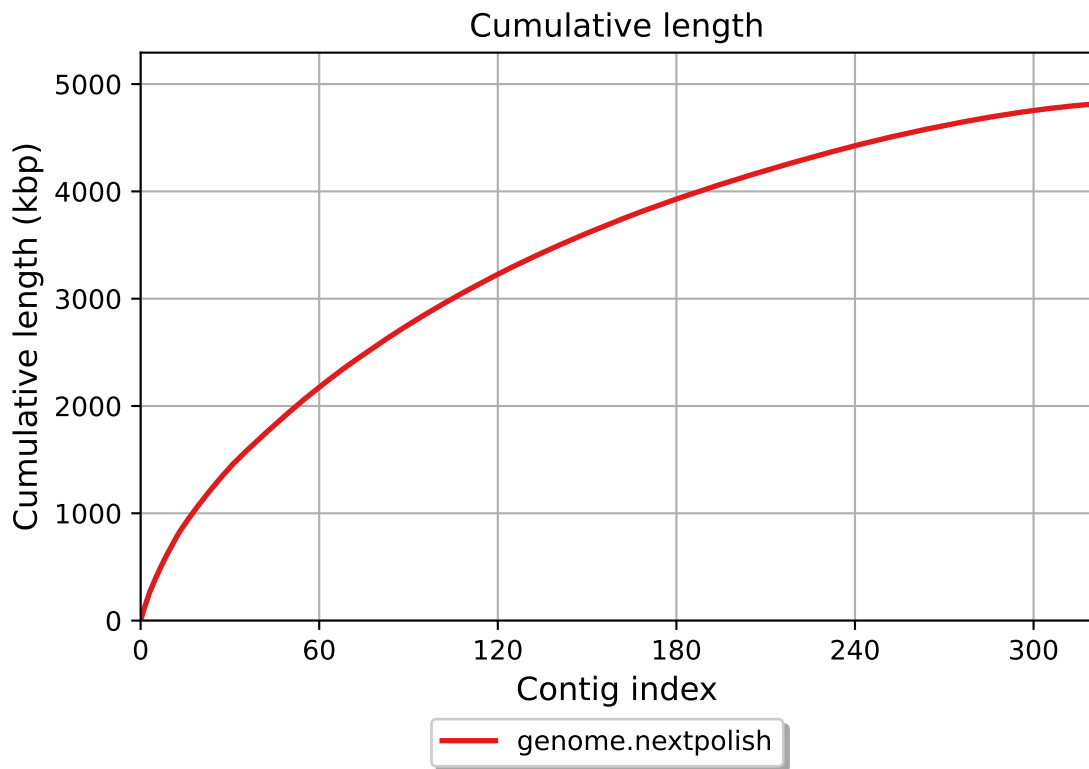


# Report

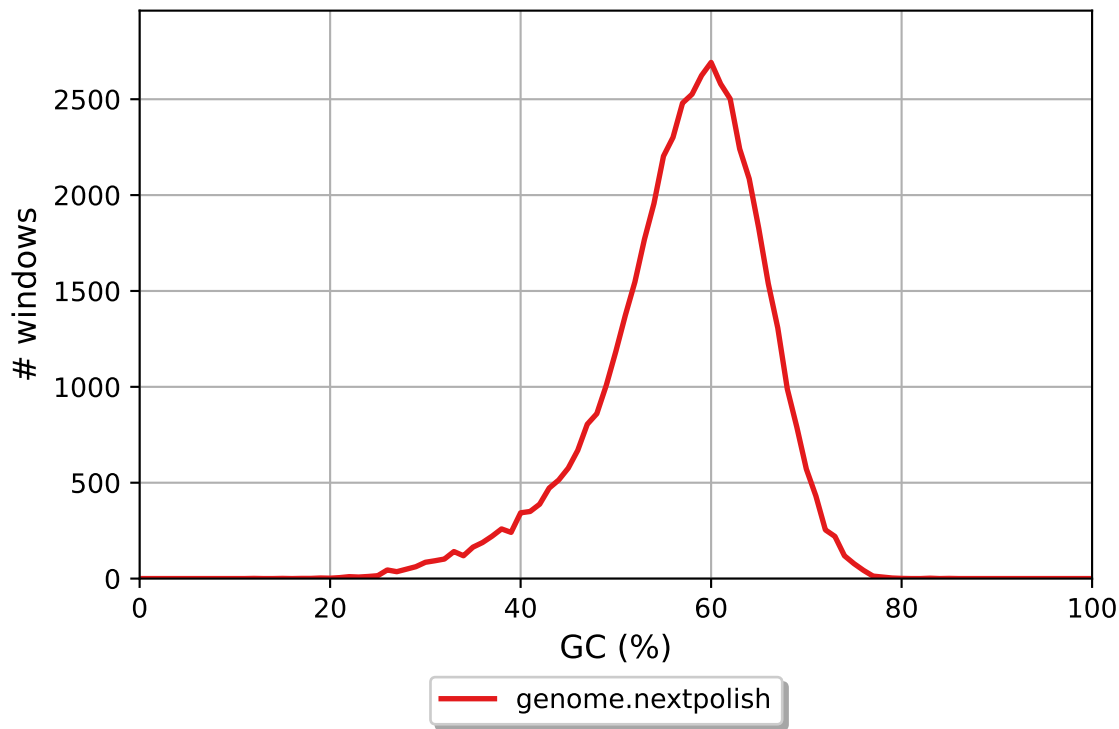
	genome.nextpolish
# contigs (>= 0 bp)	320
# contigs (>= 1000 bp)	319
# contigs (>= 5000 bp)	281
# contigs (>= 10000 bp)	175
# contigs (>= 25000 bp)	44
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	4812026
Total length (>= 1000 bp)	4811211
Total length (>= 5000 bp)	4671969
Total length (>= 10000 bp)	3880072
Total length (>= 25000 bp)	1798131
Total length (>= 50000 bp)	726100
# contigs	320
Largest contig	96009
Total length	4812026
GC (%)	56.99
N50	19468
N90	7647
auN	26694.0
L50	72
L90	228
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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

