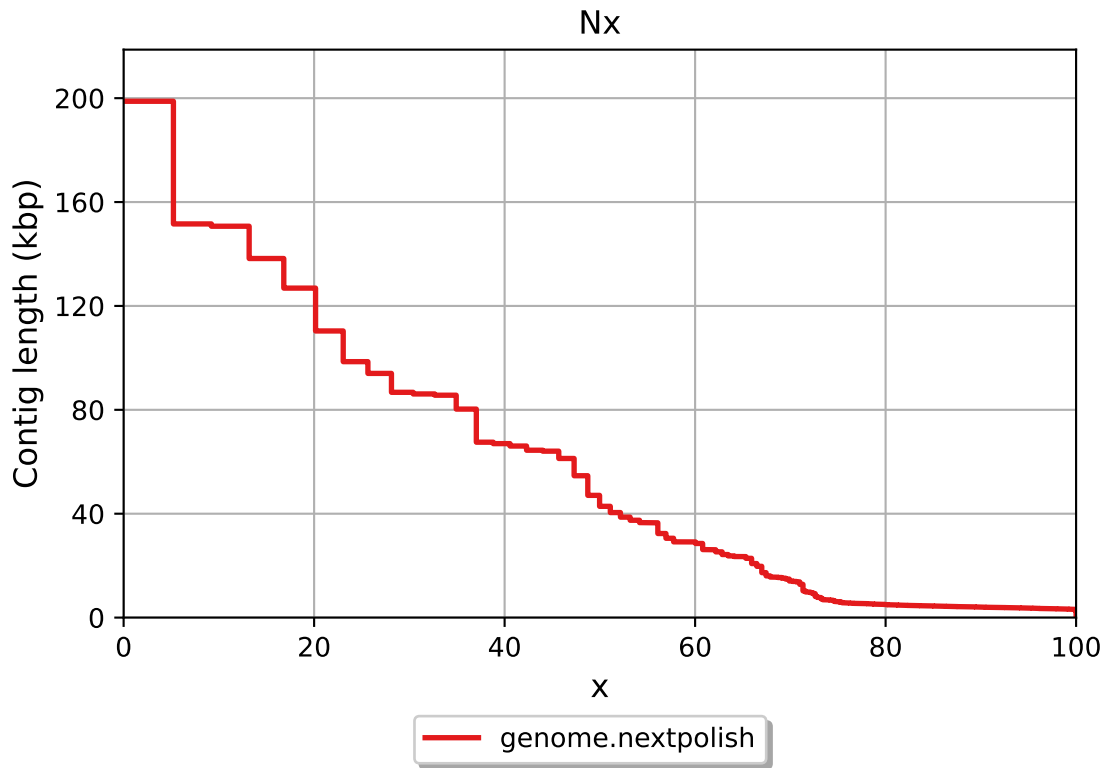
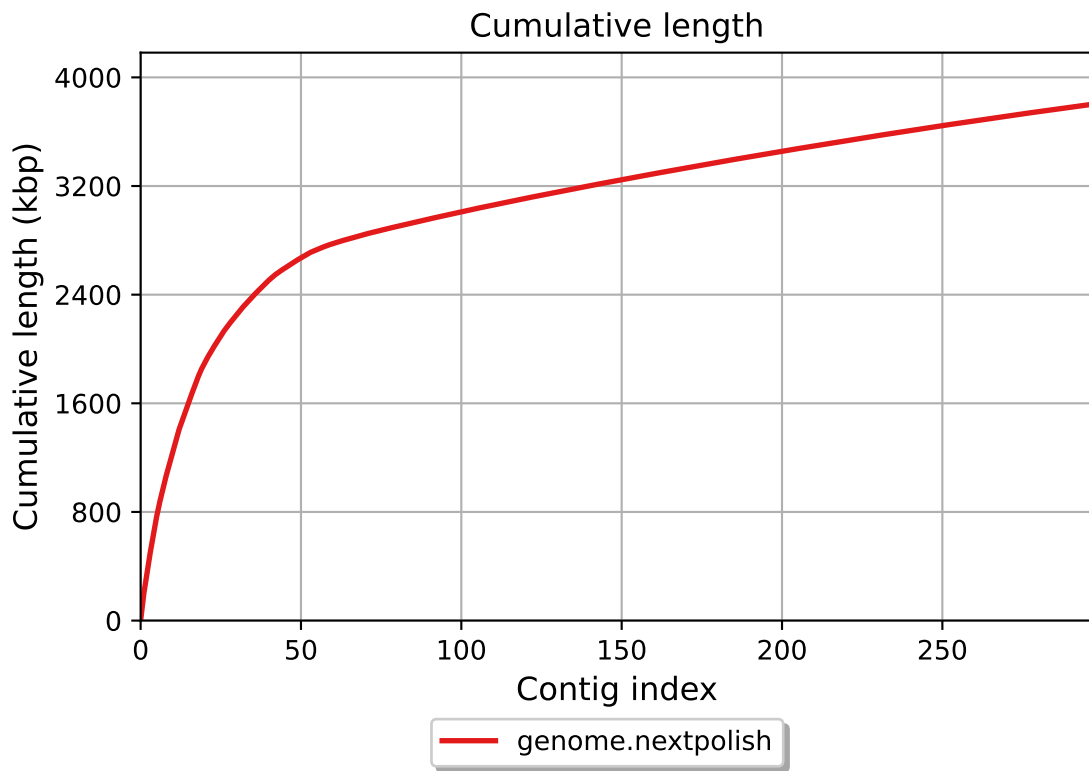


# Report

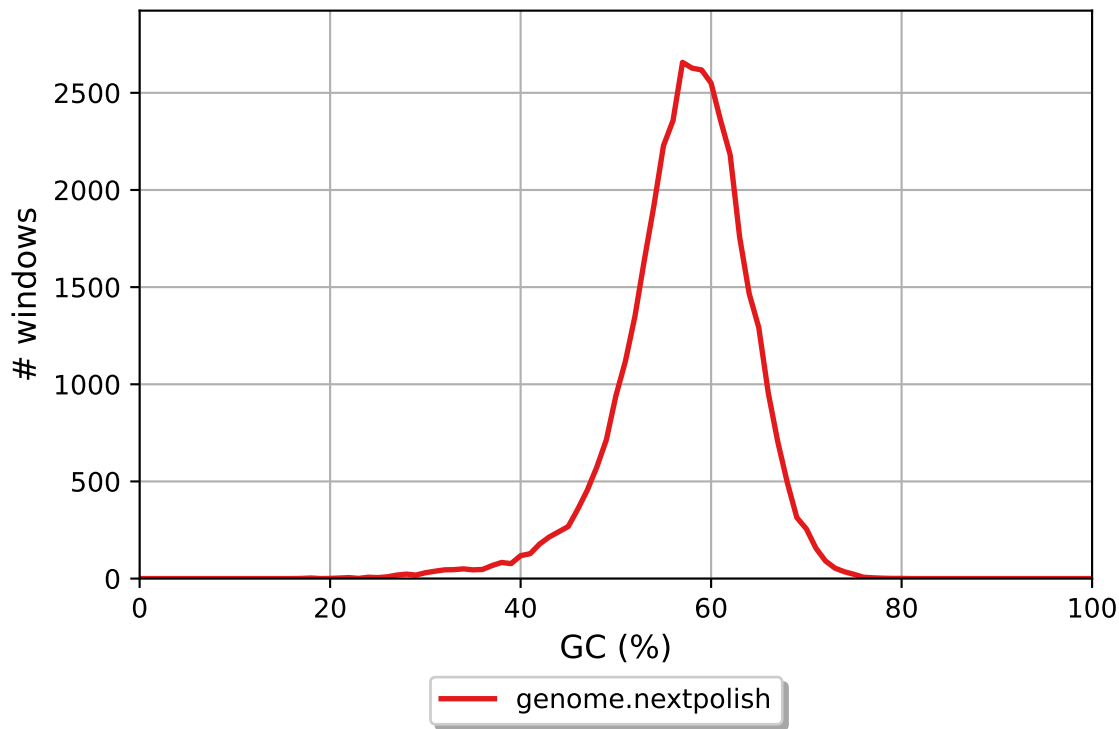
	genome.nextpolish
# contigs (>= 0 bp)	297
# contigs (>= 1000 bp)	297
# contigs (>= 5000 bp)	107
# contigs (>= 10000 bp)	54
# contigs (>= 25000 bp)	35
# contigs (>= 50000 bp)	19
Total length (>= 0 bp)	3802015
Total length (>= 1000 bp)	3802015
Total length (>= 5000 bp)	3045611
Total length (>= 10000 bp)	2722073
Total length (>= 25000 bp)	2388934
Total length (>= 50000 bp)	1852857
# contigs	297
Largest contig	198807
Total length	3802015
GC (%)	57.31
N50	42840
N90	4069
auN	62644.8
L50	21
L90	192
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 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

