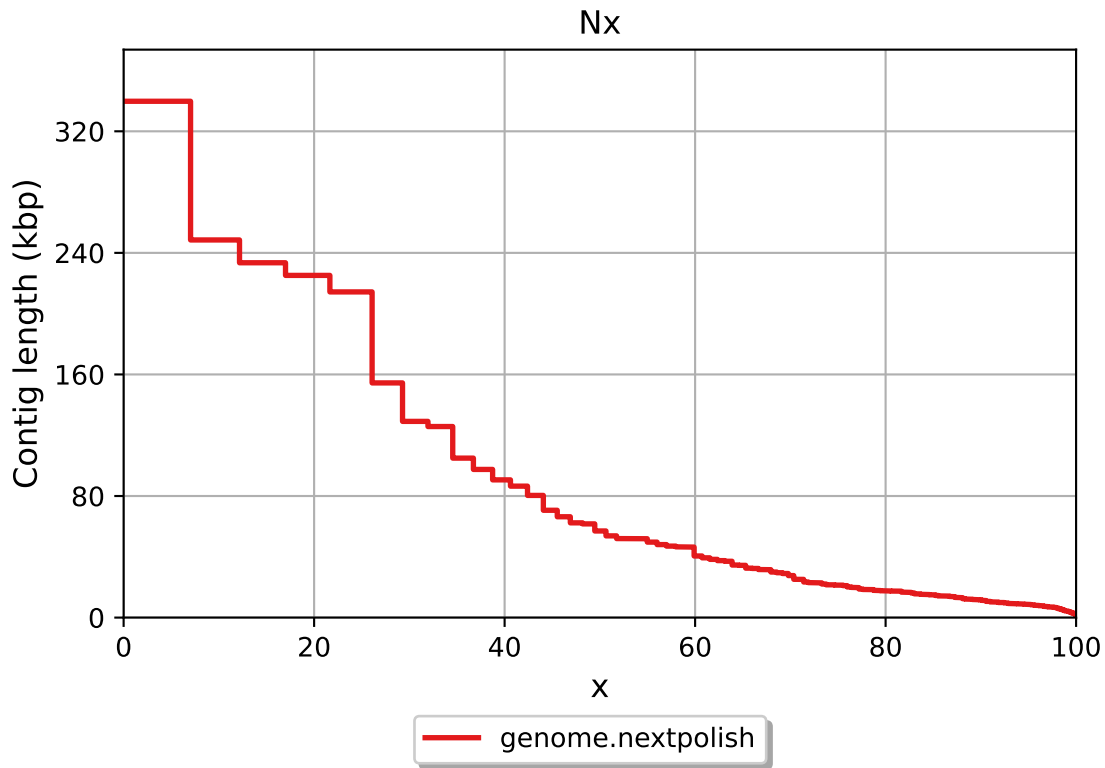
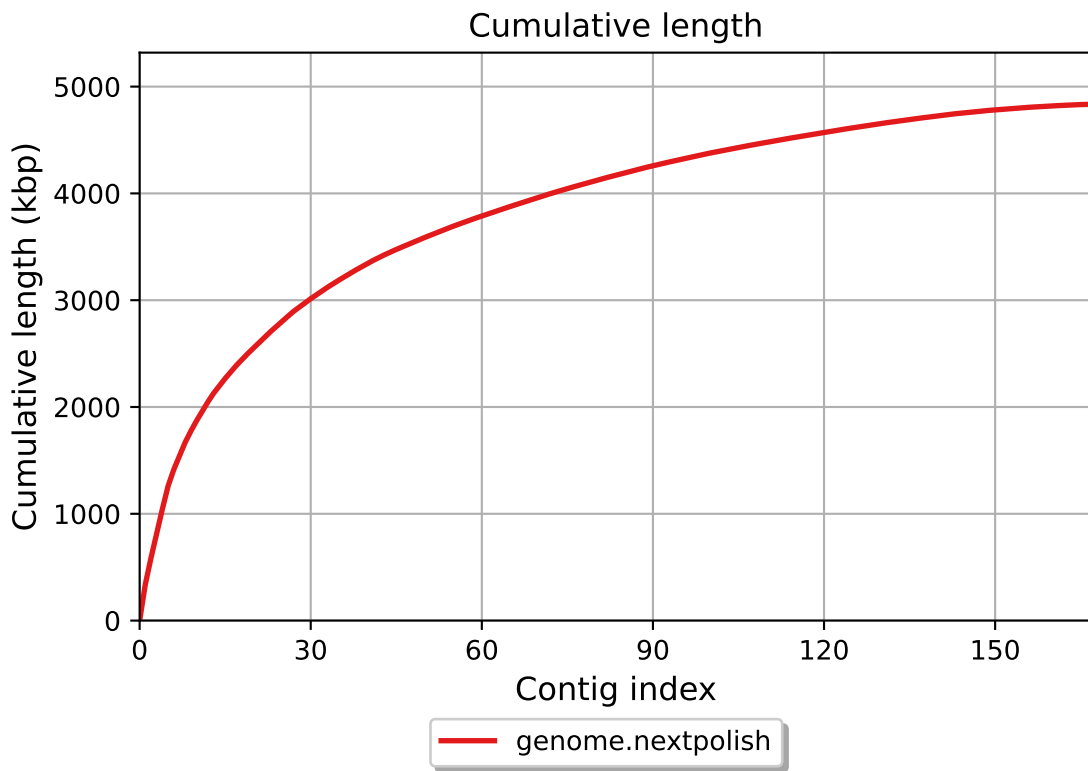


# Report

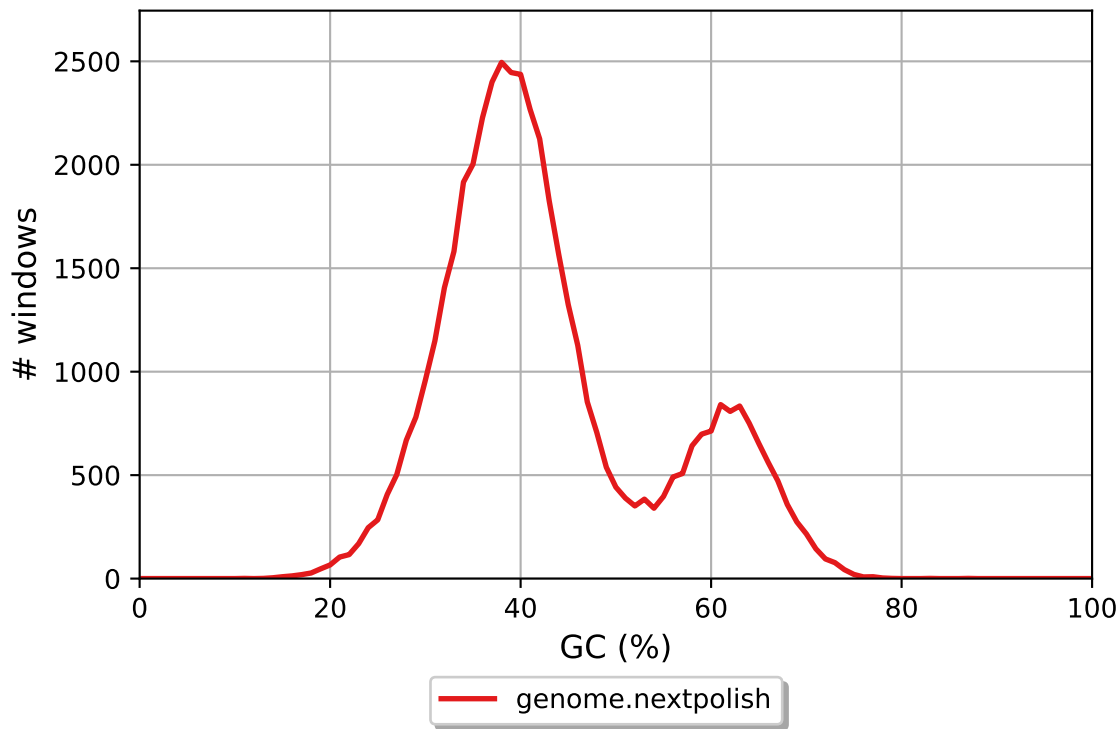
	genome.nextpolish
# contigs (>= 0 bp)	167
# contigs (>= 1000 bp)	167
# contigs (>= 5000 bp)	148
# contigs (>= 10000 bp)	106
# contigs (>= 25000 bp)	44
# contigs (>= 50000 bp)	22
Total length (>= 0 bp)	4834590
Total length (>= 1000 bp)	4834590
Total length (>= 5000 bp)	4772867
Total length (>= 10000 bp)	4439992
Total length (>= 25000 bp)	3451812
Total length (>= 50000 bp)	2658003
# contigs	167
Largest contig	339761
Total length	4834590
GC (%)	43.12
N50	56994
N90	11696
auN	105093.1
L50	18
L90	98
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 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

