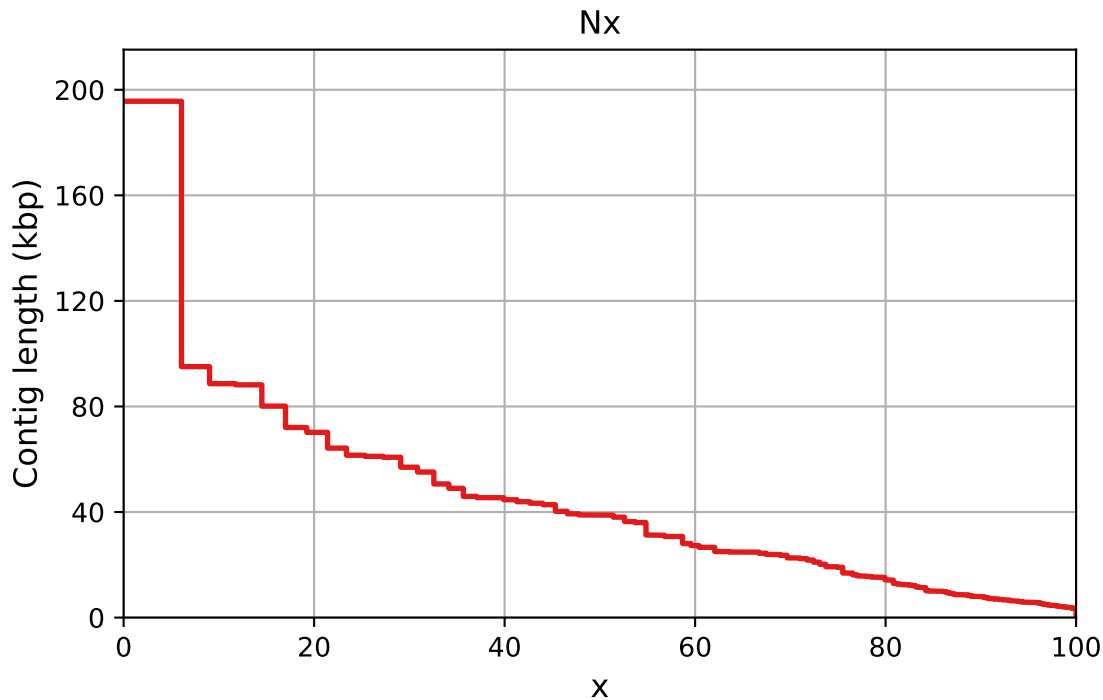


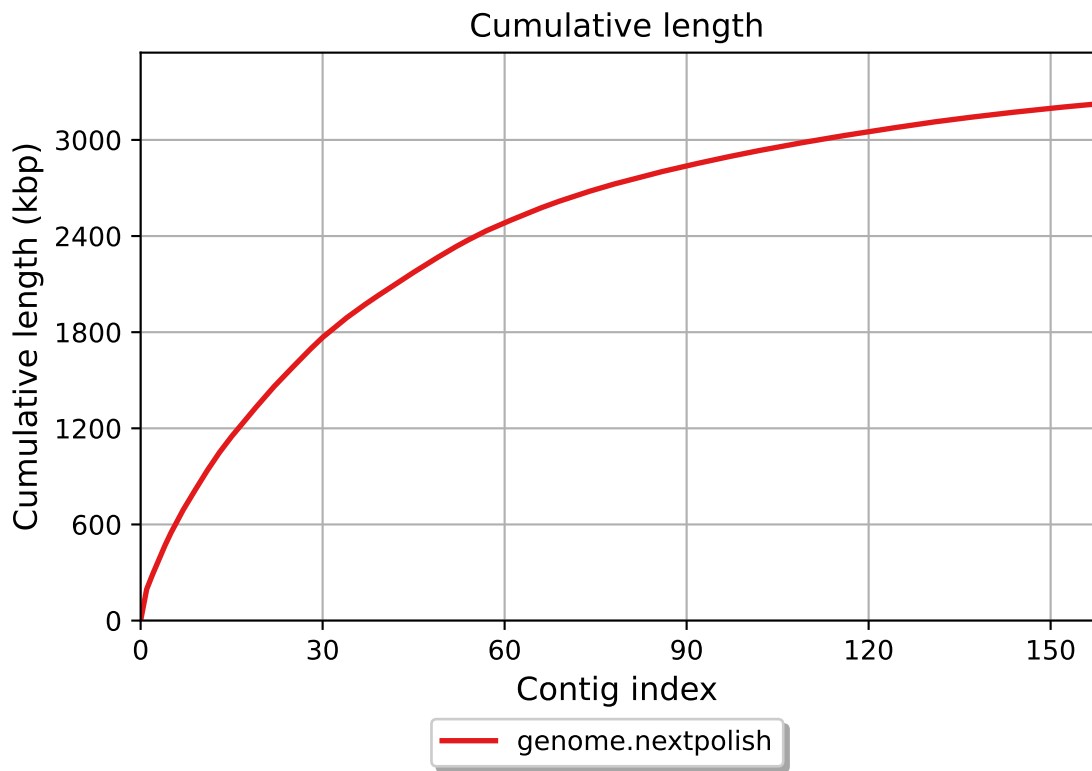
# Report

	genome.nextpolish
# contigs (>= 0 bp)	157
# contigs (>= 1000 bp)	157
# contigs (>= 5000 bp)	132
# contigs (>= 10000 bp)	82
# contigs (>= 25000 bp)	40
# contigs (>= 50000 bp)	14
Total length (>= 0 bp)	3222495
Total length (>= 1000 bp)	3222495
Total length (>= 5000 bp)	3118301
Total length (>= 10000 bp)	2764239
Total length (>= 25000 bp)	2049863
Total length (>= 50000 bp)	1100254
# contigs	157
Largest contig	195644
Total length	3222495
GC (%)	57.91
N50	38850
N90	7955
auN	48410.7
L50	26
L90	98
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 0 part

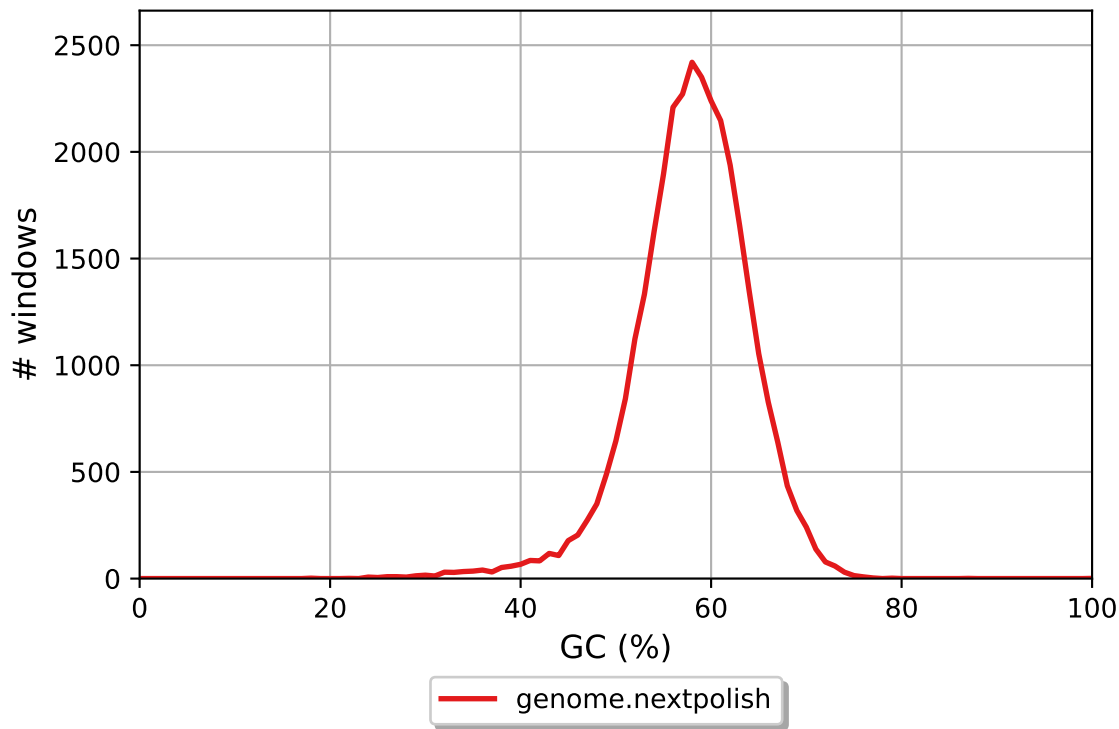
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

