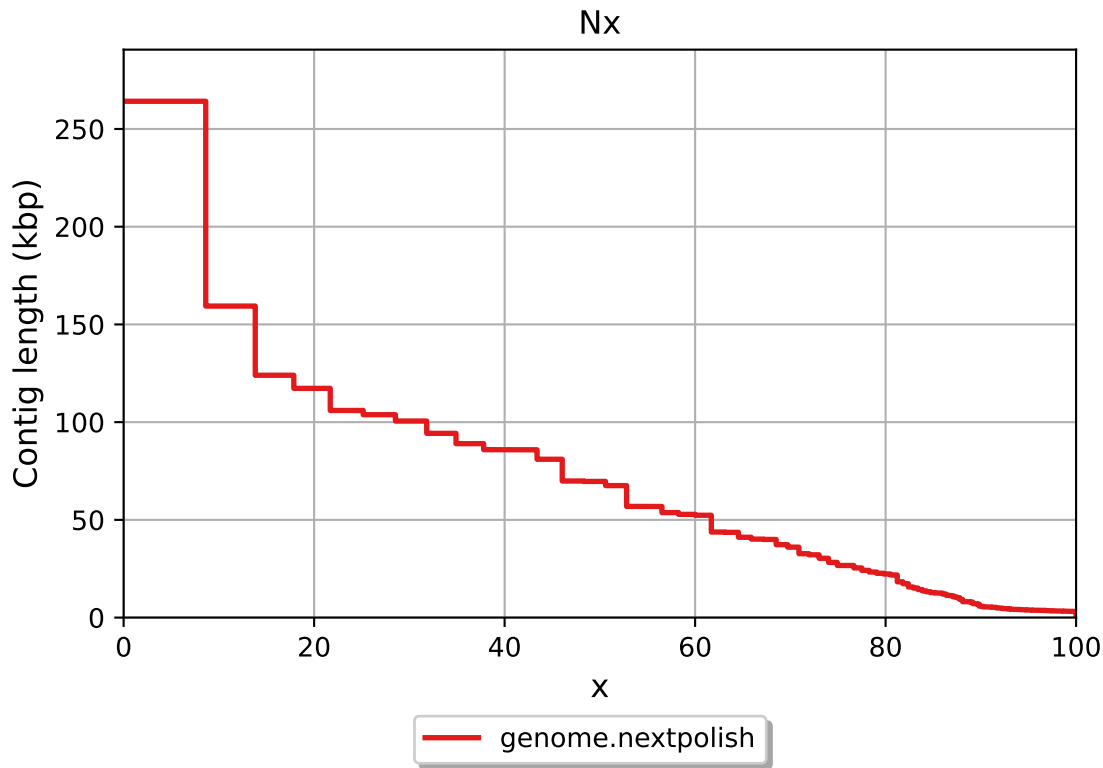
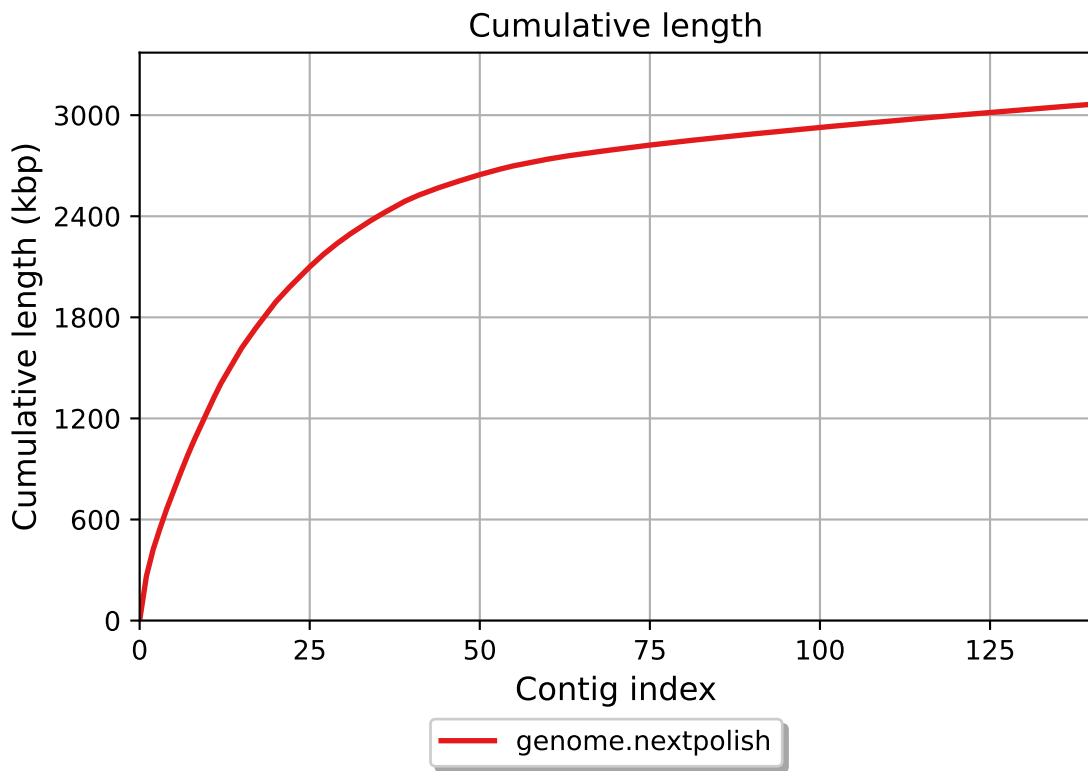


Report

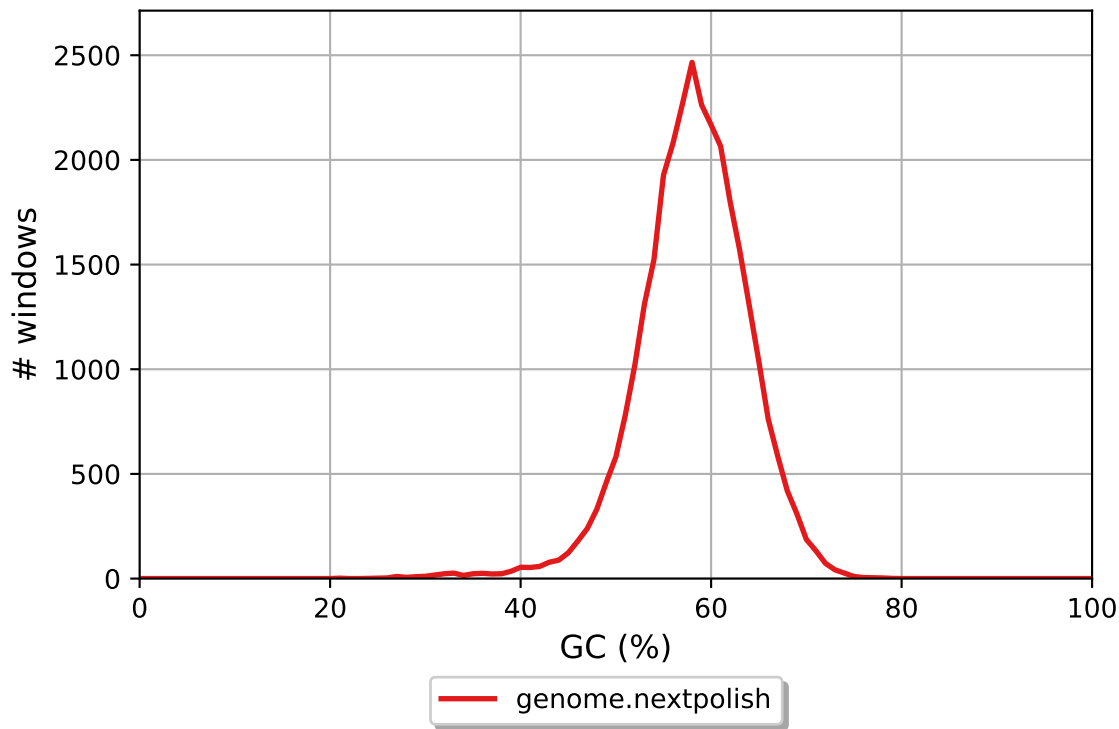
	genome.nextpolish
# contigs (>= 0 bp)	140
# contigs (>= 1000 bp)	140
# contigs (>= 5000 bp)	72
# contigs (>= 10000 bp)	54
# contigs (>= 25000 bp)	34
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	3064665
Total length (>= 1000 bp)	3064665
Total length (>= 5000 bp)	2807662
Total length (>= 10000 bp)	2690204
Total length (>= 25000 bp)	2375022
Total length (>= 50000 bp)	1890934
# contigs	140
Largest contig	264178
Total length	3064665
GC (%)	58.07
N50	69668
N90	5870
auN	80990.5
L50	14
L90	63
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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

