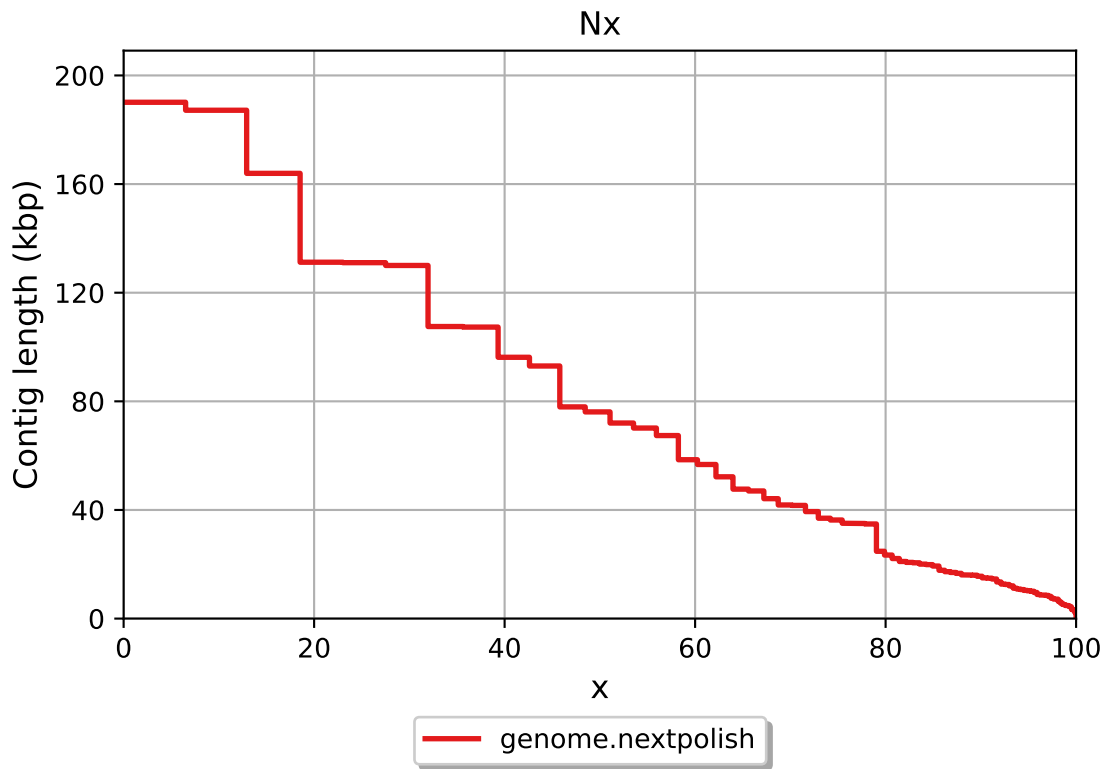
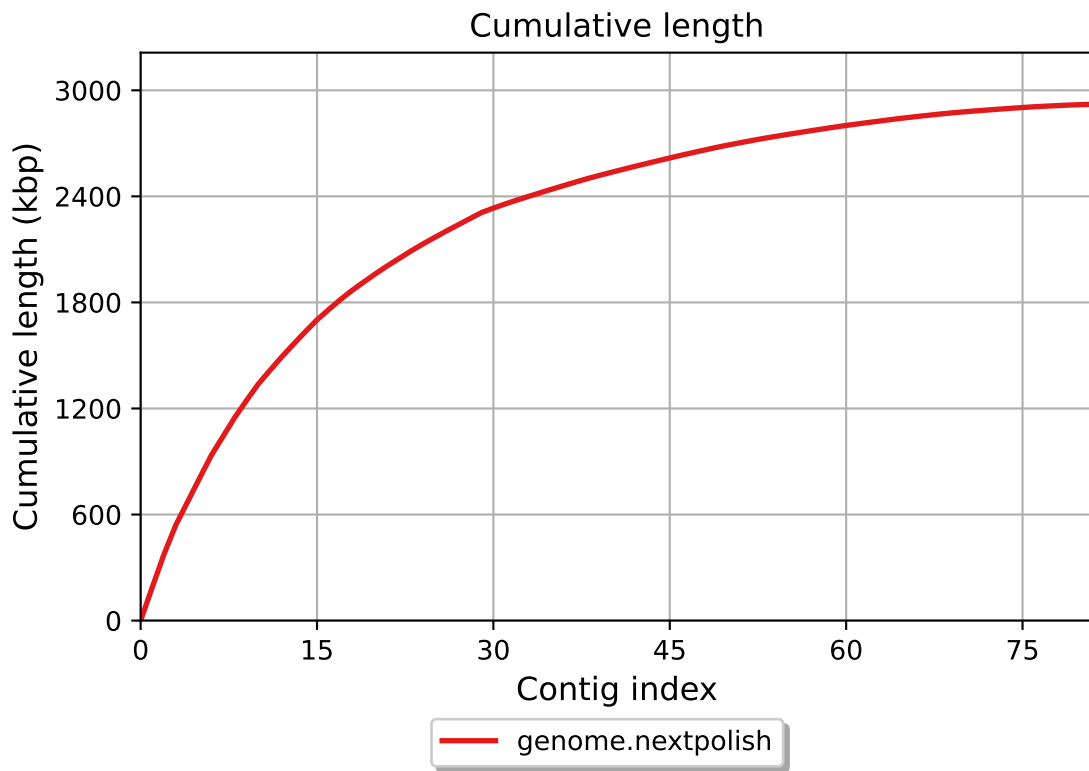


Report

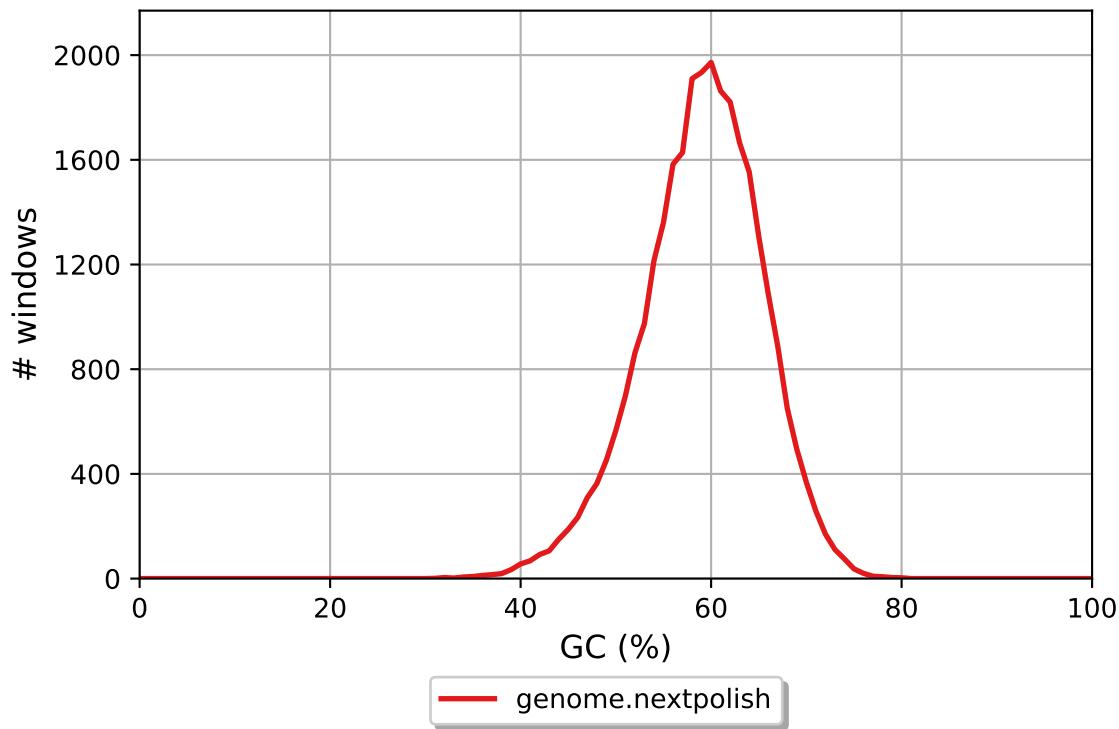
	genome.nextpolish
# contigs (>= 0 bp)	81
# contigs (>= 1000 bp)	81
# contigs (>= 5000 bp)	72
# contigs (>= 10000 bp)	59
# contigs (>= 25000 bp)	29
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	2921039
Total length (>= 1000 bp)	2921039
Total length (>= 5000 bp)	2888468
Total length (>= 10000 bp)	2791559
Total length (>= 25000 bp)	2308686
Total length (>= 50000 bp)	1868753
# contigs	81
Largest contig	190122
Total length	2921039
GC (%)	59.02
N50	76110
N90	15583
auN	86692.9
L50	12
L90	46
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 1 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

