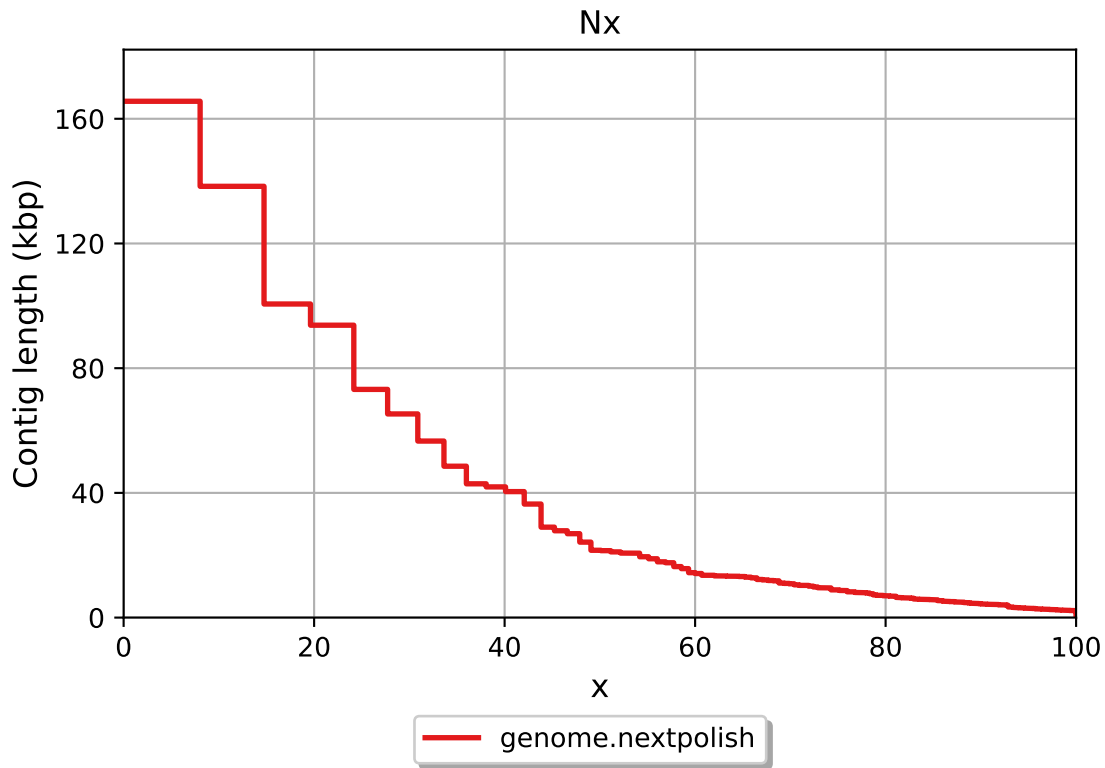
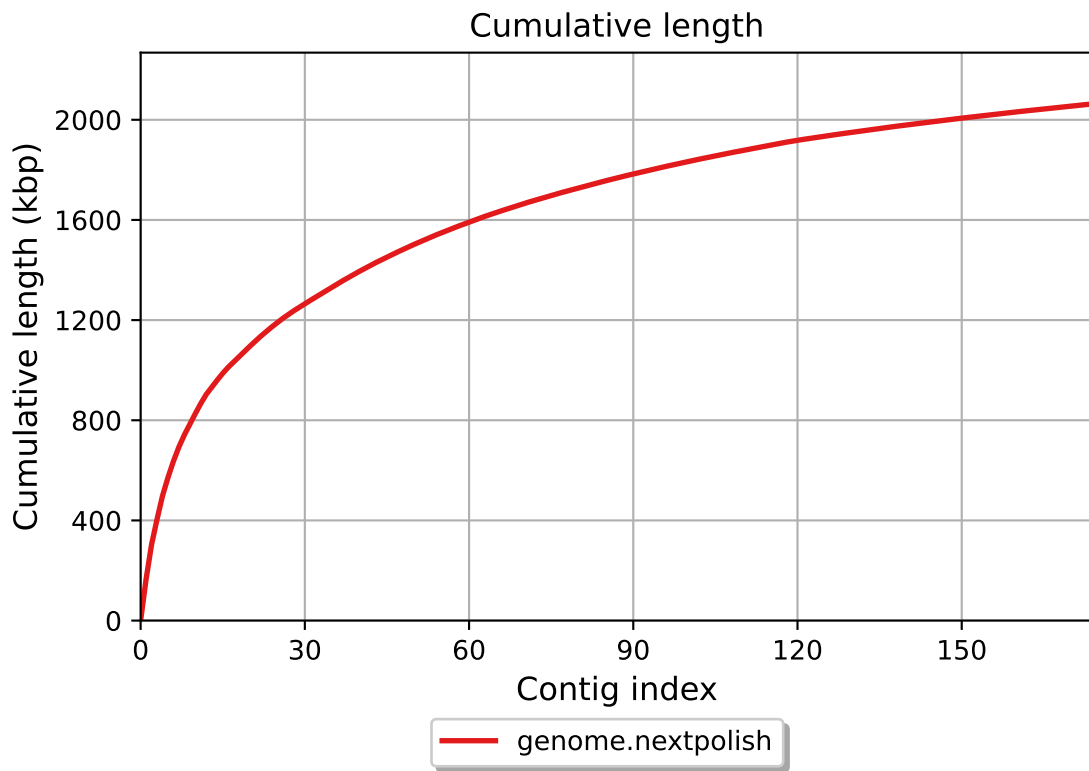


Report

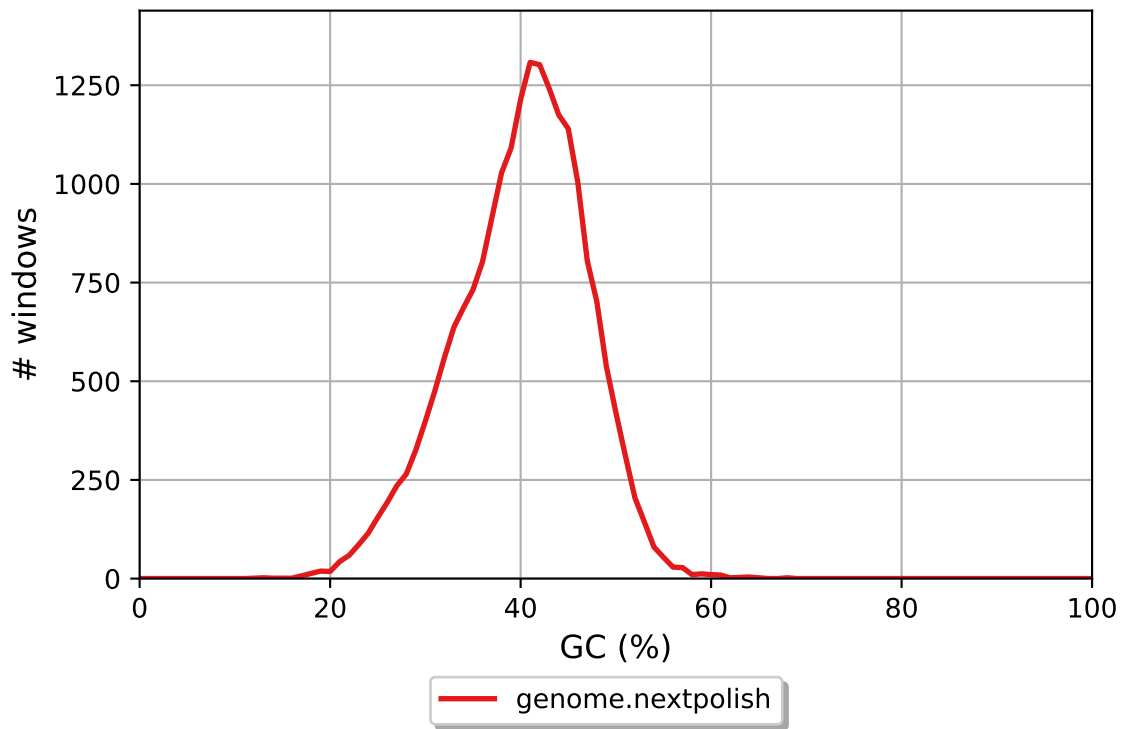
	genome.nextpolish
# contigs (>= 0 bp)	174
# contigs (>= 1000 bp)	173
# contigs (>= 5000 bp)	94
# contigs (>= 10000 bp)	49
# contigs (>= 25000 bp)	15
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	2061932
Total length (>= 1000 bp)	2061382
Total length (>= 5000 bp)	1804030
Total length (>= 10000 bp)	1492969
Total length (>= 25000 bp)	987434
Total length (>= 50000 bp)	693455
# contigs	174
Largest contig	165614
Total length	2061932
GC (%)	40.03
N50	21590
N90	4473
auN	48858.2
L50	17
L90	105
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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

