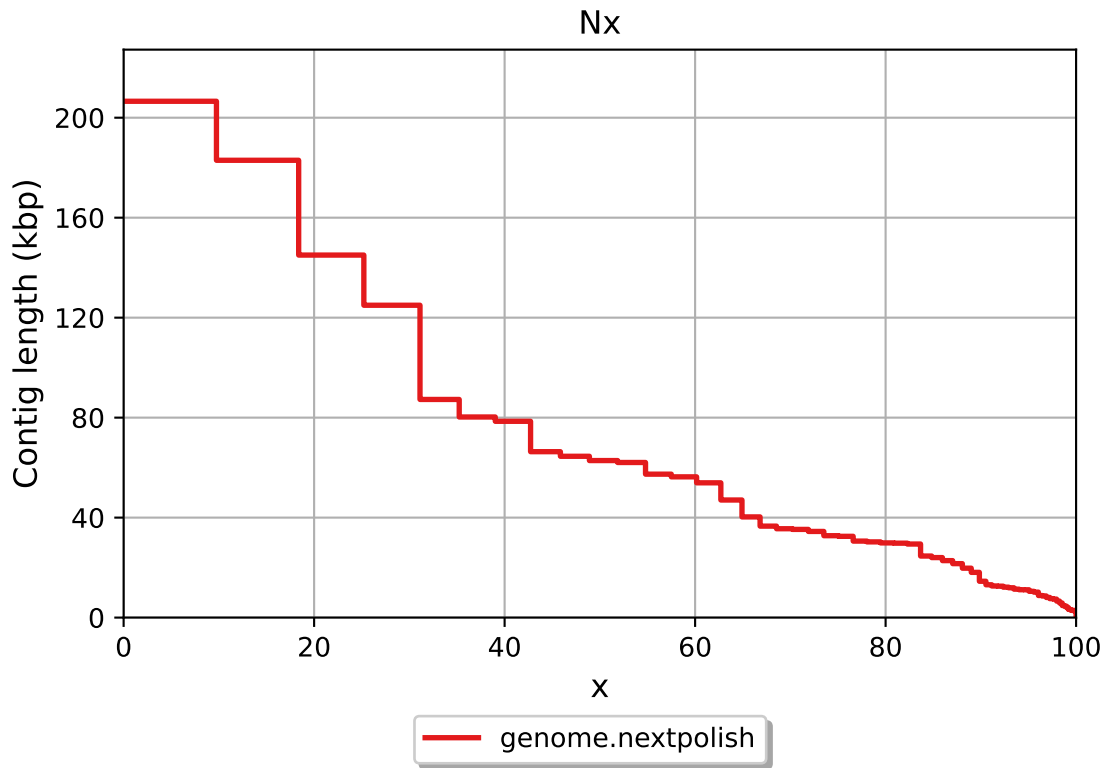
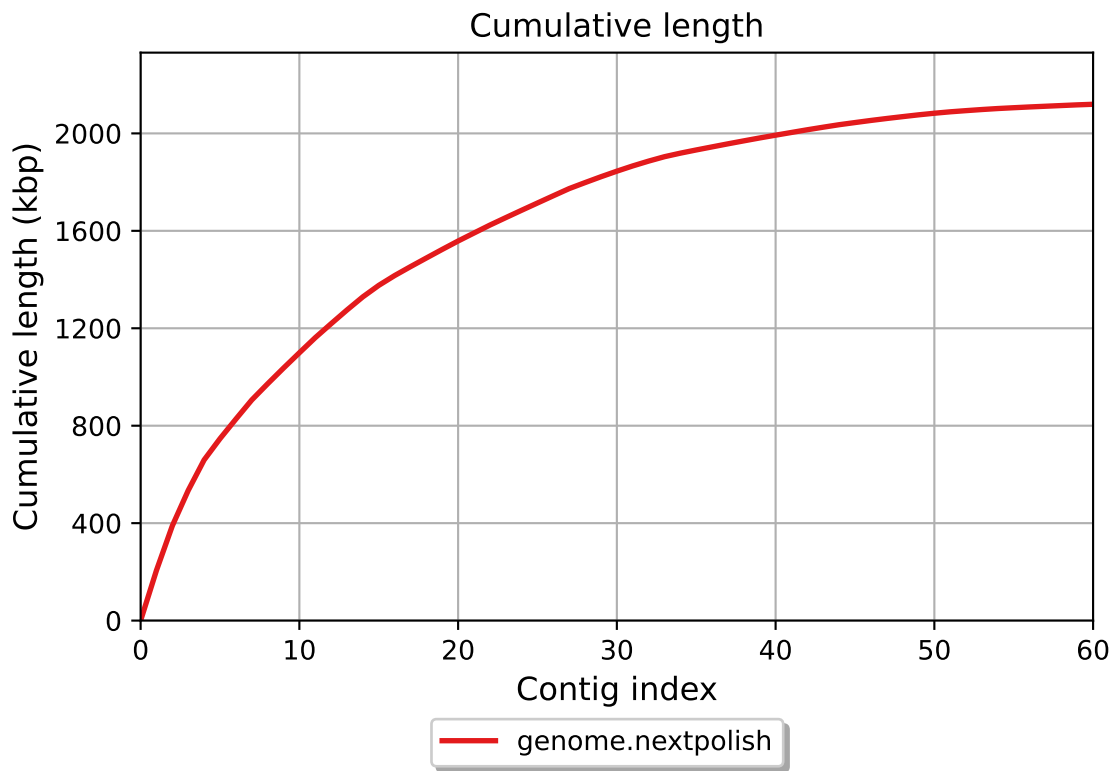


Report

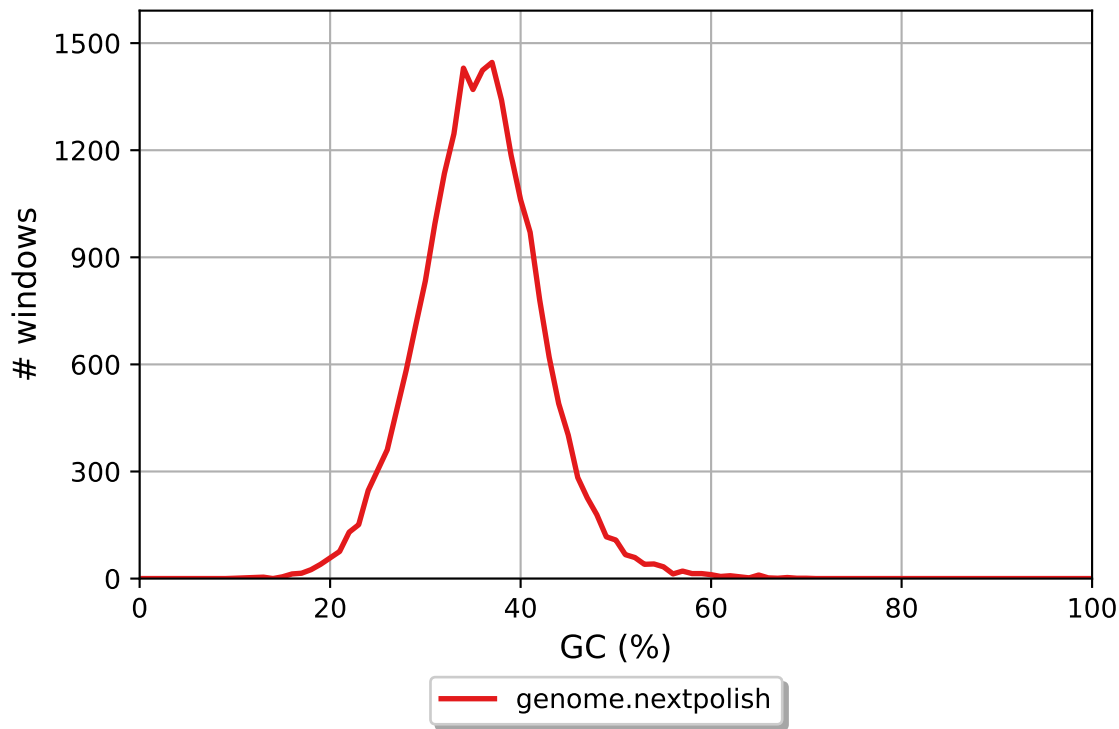
	genome.nextpolish
# contigs (>= 0 bp)	60
# contigs (>= 1000 bp)	60
# contigs (>= 5000 bp)	51
# contigs (>= 10000 bp)	44
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	14
Total length (>= 0 bp)	2119572
Total length (>= 1000 bp)	2119572
Total length (>= 5000 bp)	2088605
Total length (>= 10000 bp)	2035657
Total length (>= 25000 bp)	1773478
Total length (>= 50000 bp)	1329016
# contigs	60
Largest contig	206575
Total length	2119572
GC (%)	35.83
N50	62826
N90	14531
auN	84572.3
L50	10
L90	34
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 4 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

