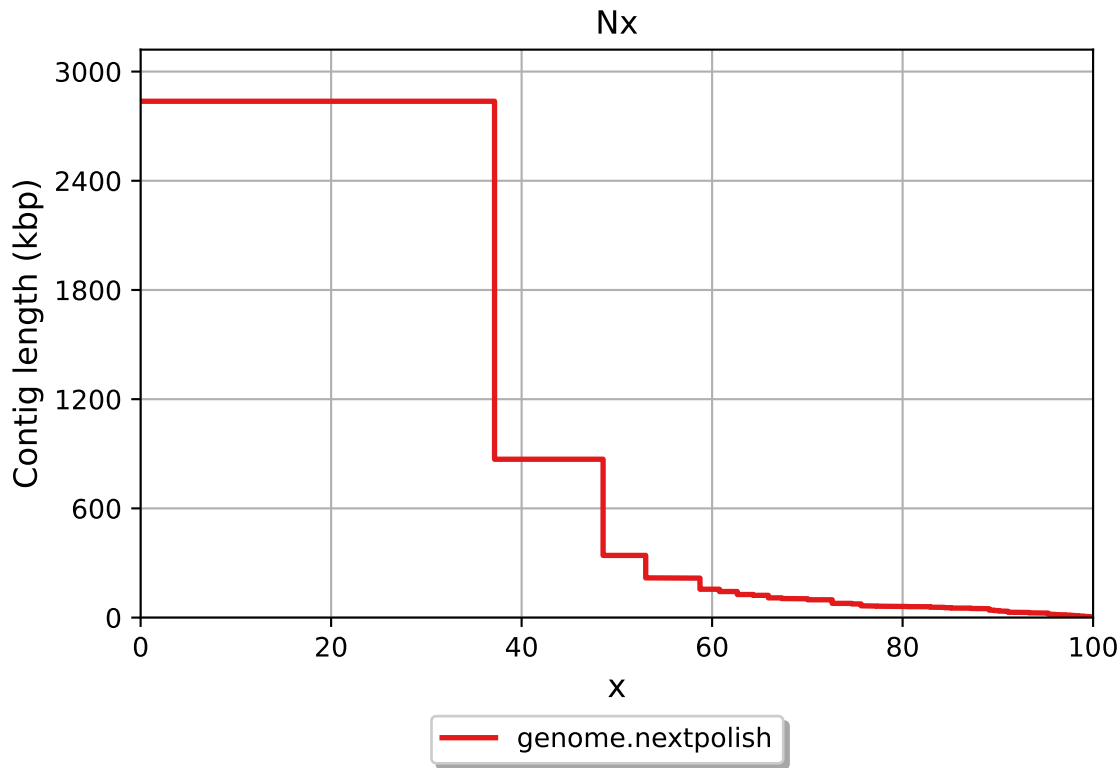


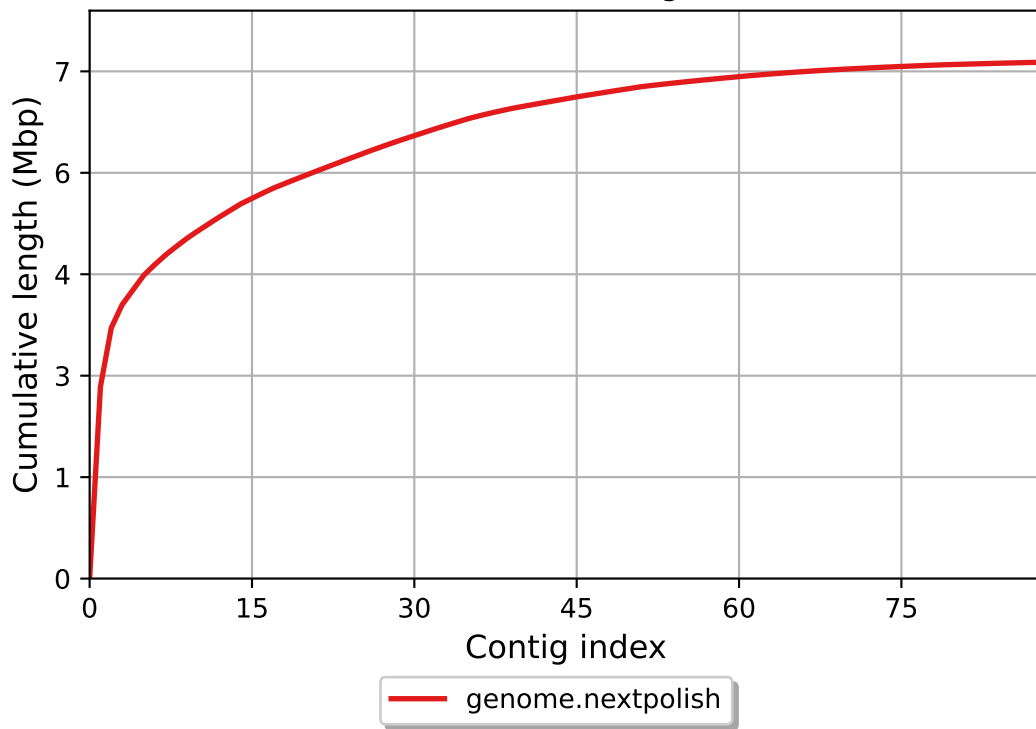
Report

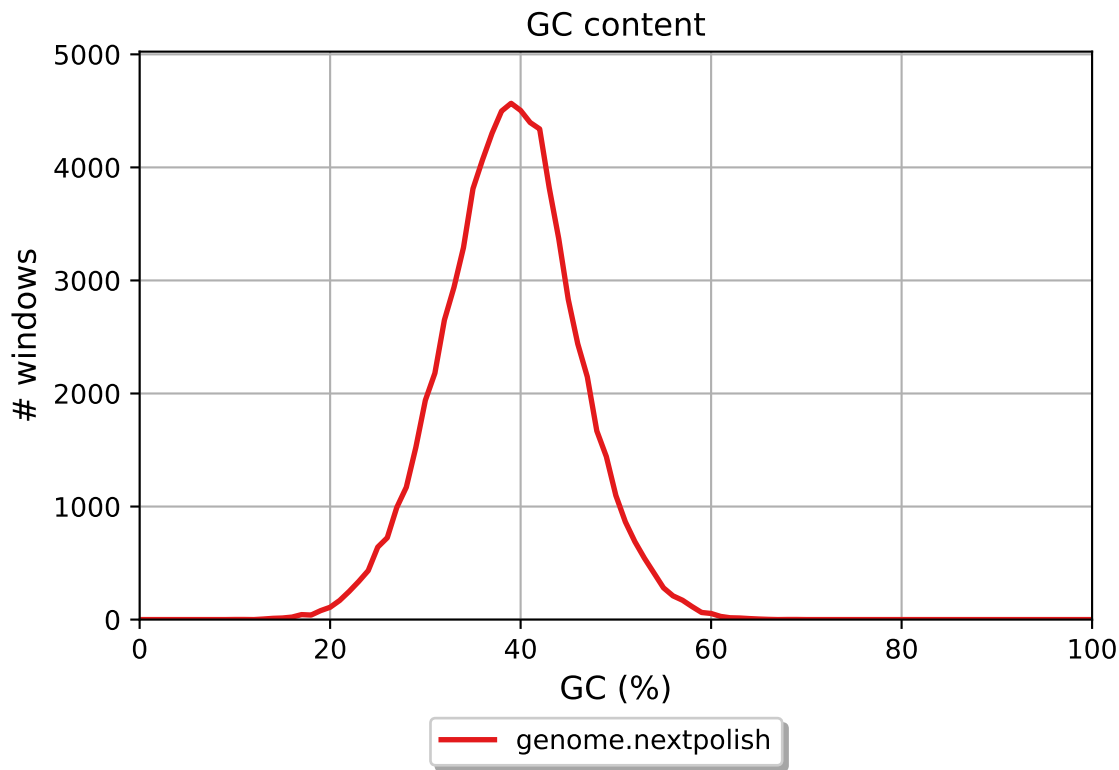
	genome.nextpolish
# contigs (>= 0 bp)	88
# contigs (>= 1000 bp)	88
# contigs (>= 5000 bp)	79
# contigs (>= 10000 bp)	67
# contigs (>= 25000 bp)	48
# contigs (>= 50000 bp)	33
Total length (>= 0 bp)	7634449
Total length (>= 1000 bp)	7634449
Total length (>= 5000 bp)	7597477
Total length (>= 10000 bp)	7507261
Total length (>= 25000 bp)	7199296
Total length (>= 50000 bp)	6703693
# contigs	88
Largest contig	2836856
Total length	7634449
GC (%)	38.86
N50	341572
N90	38392
auN	1210282.0
L50	3
L90	37
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	51 + 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).



Cumulative length





genome.nextpolish GC content

