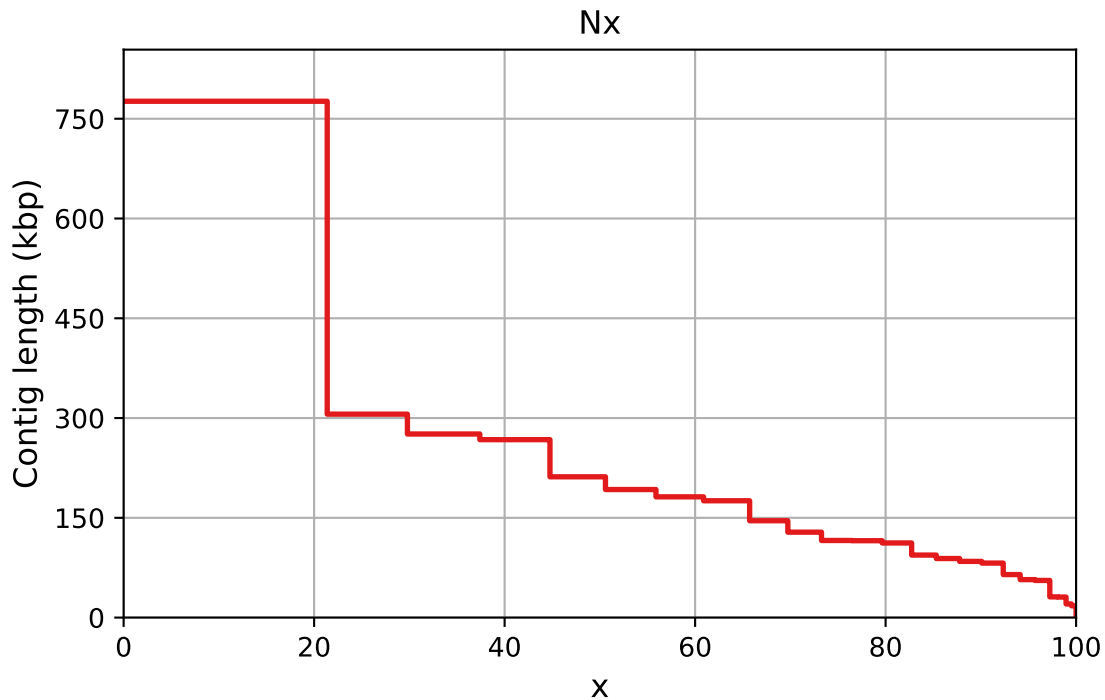


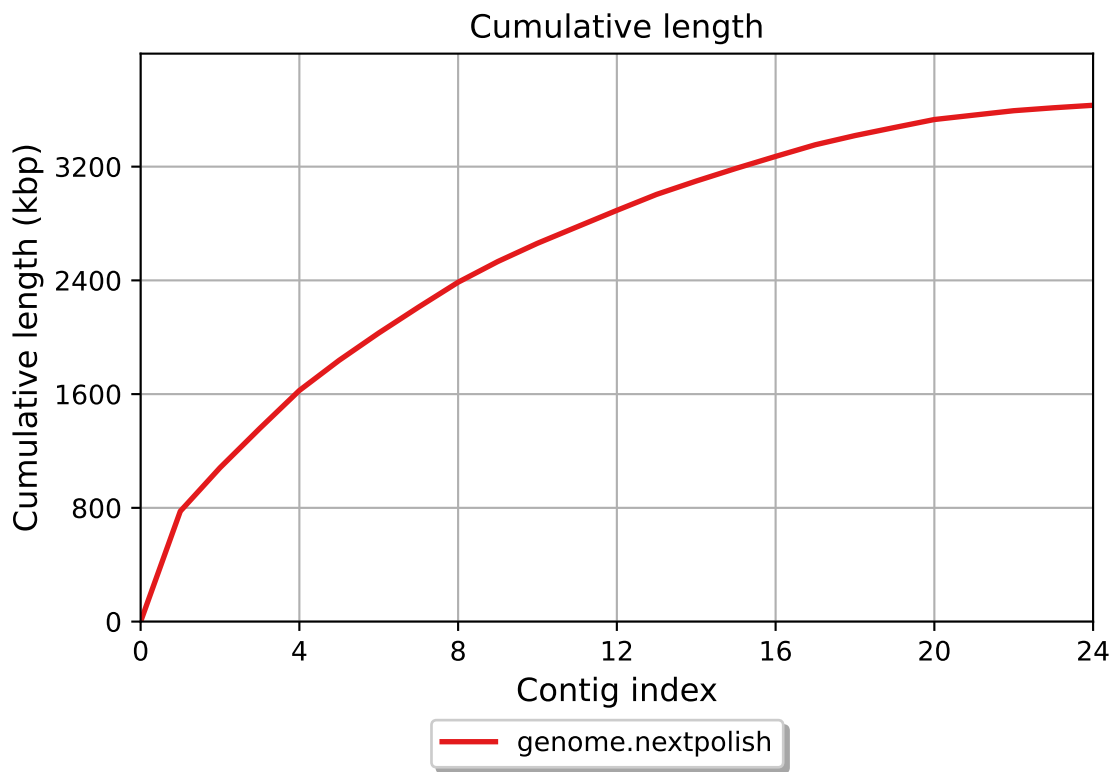
Report

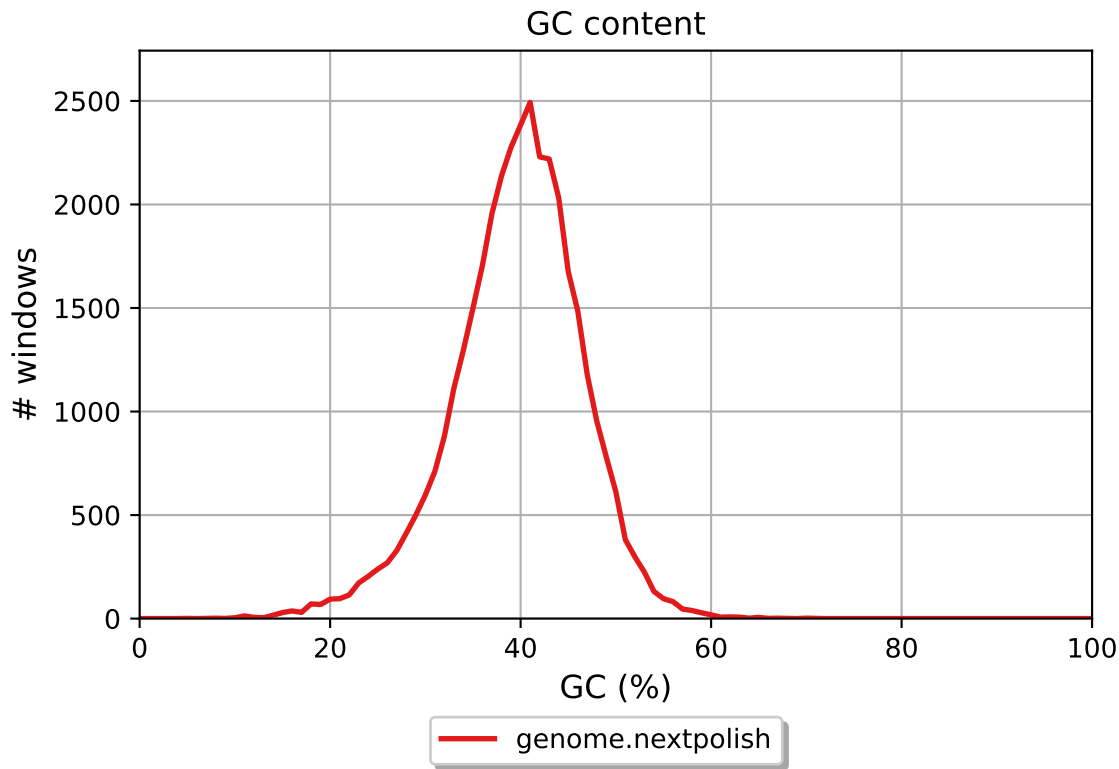
	genome.nextpolish
# contigs (>= 0 bp)	24
# contigs (>= 1000 bp)	24
# contigs (>= 5000 bp)	24
# contigs (>= 10000 bp)	24
# contigs (>= 25000 bp)	22
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	3631711
Total length (>= 1000 bp)	3631711
Total length (>= 5000 bp)	3631711
Total length (>= 10000 bp)	3631711
Total length (>= 25000 bp)	3593441
Total length (>= 50000 bp)	3531453
# contigs	24
Largest contig	776251
Total length	3631711
GC (%)	39.56
N50	211663
N90	84493
auN	305744.4
L50	5
L90	16
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 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).



— genome.nextpolish





genome.nextpolish GC content

