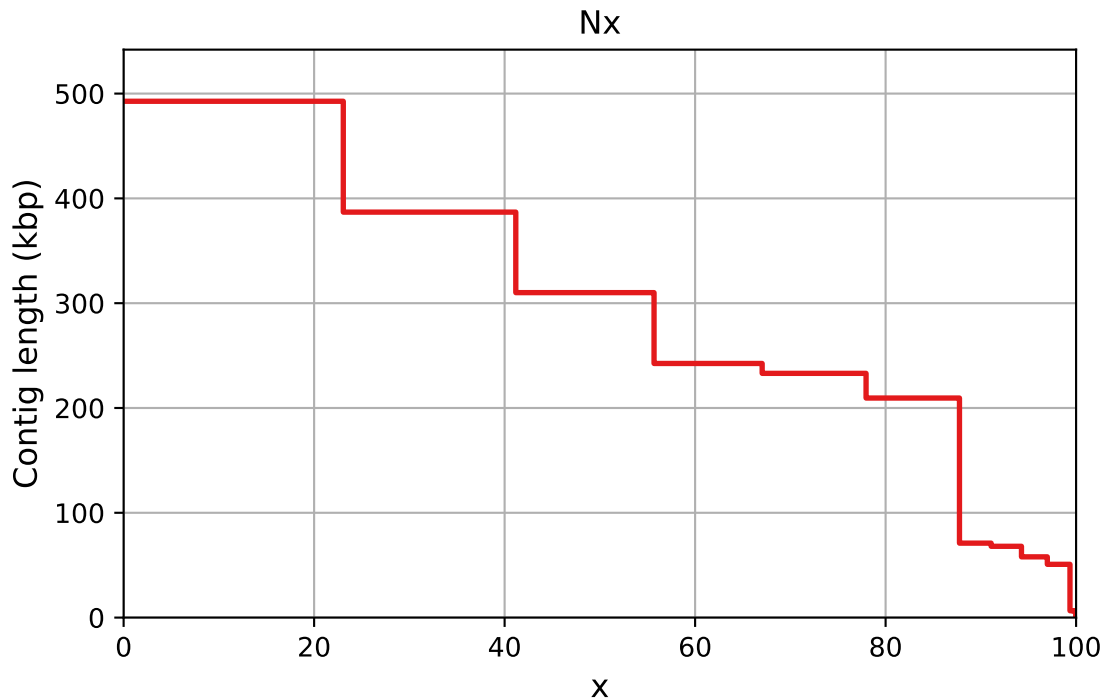


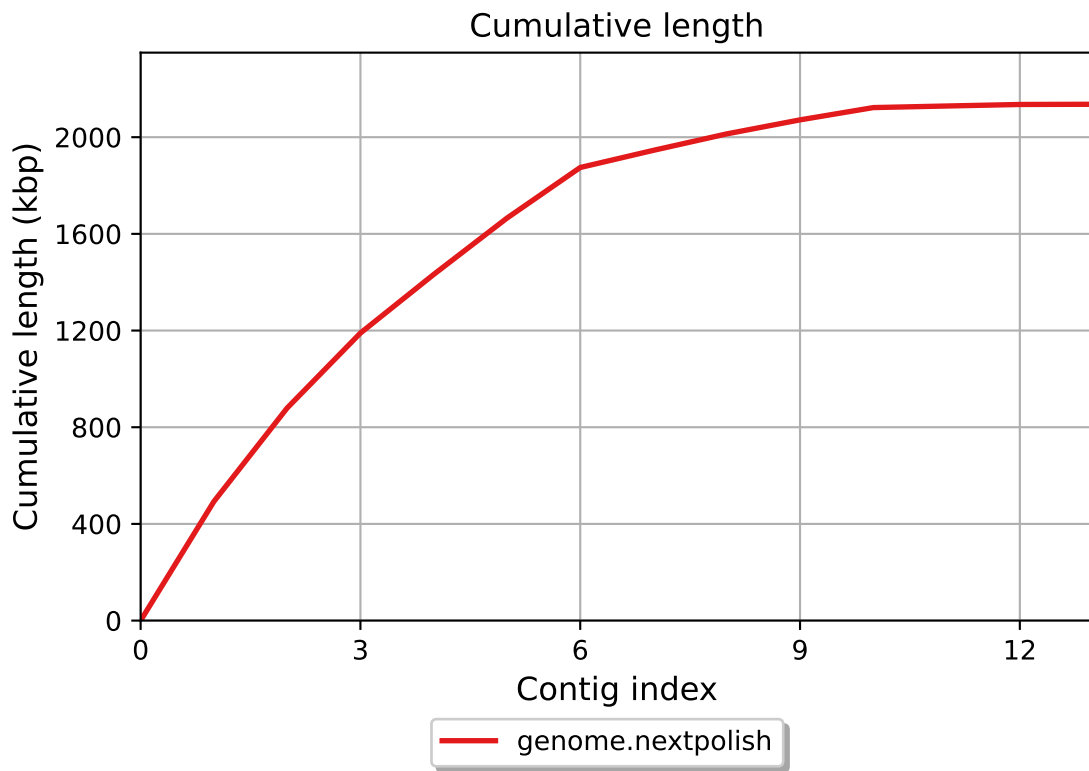
Report

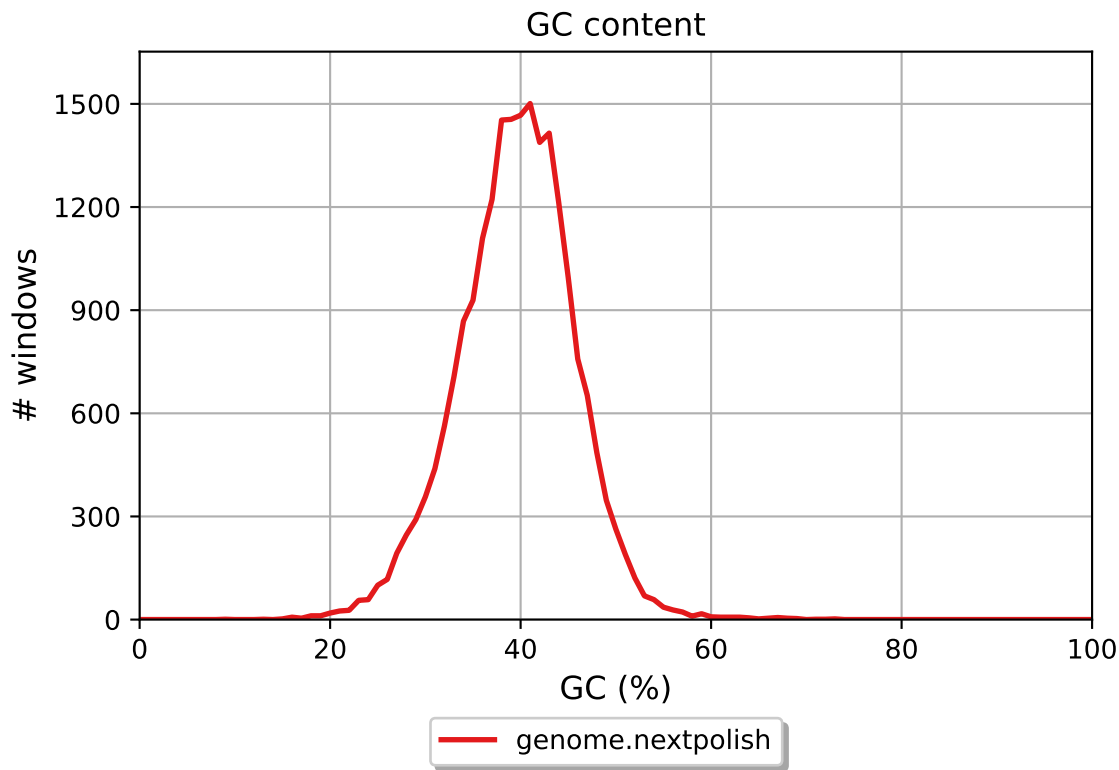
	genome.nextpolish
# contigs (>= 0 bp)	13
# contigs (>= 1000 bp)	12
# contigs (>= 5000 bp)	12
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	10
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	2136219
Total length (>= 1000 bp)	2135367
Total length (>= 5000 bp)	2135367
Total length (>= 10000 bp)	2122500
Total length (>= 25000 bp)	2122500
Total length (>= 50000 bp)	2122500
# contigs	13
Largest contig	492667
Total length	2136219
GC (%)	39.55
N50	310072
N90	71031
auN	309540.6
L50	3
L90	7
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 3 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

