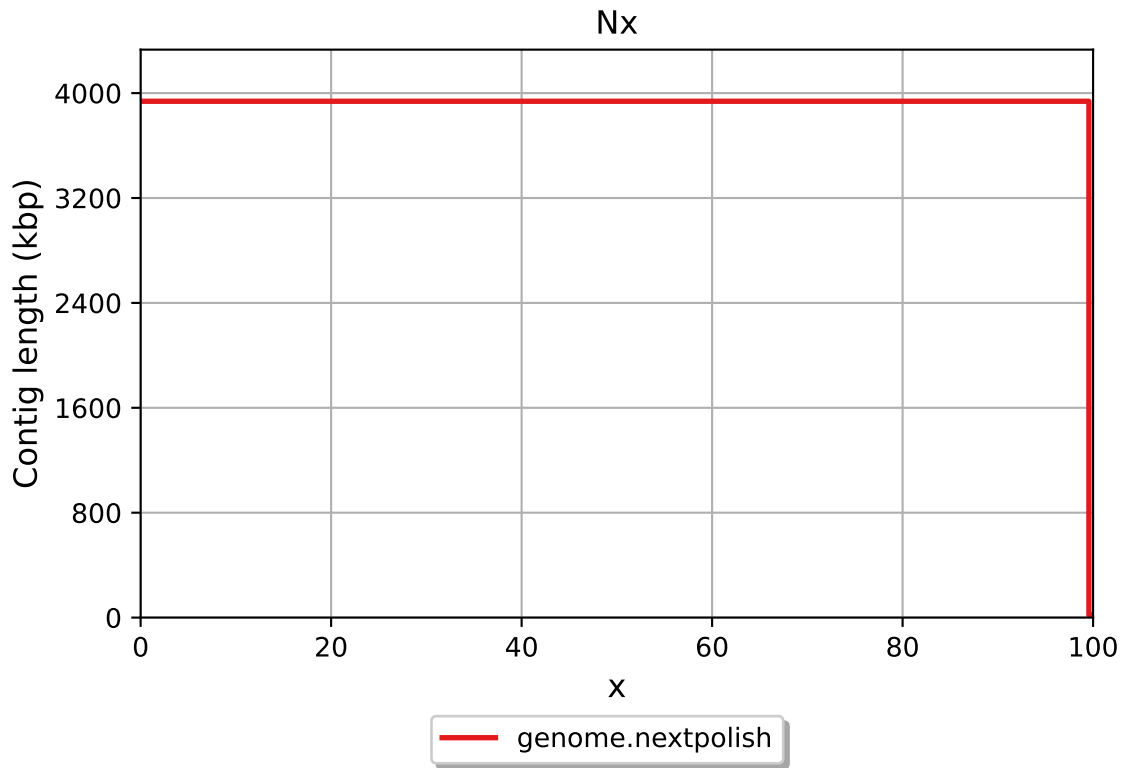
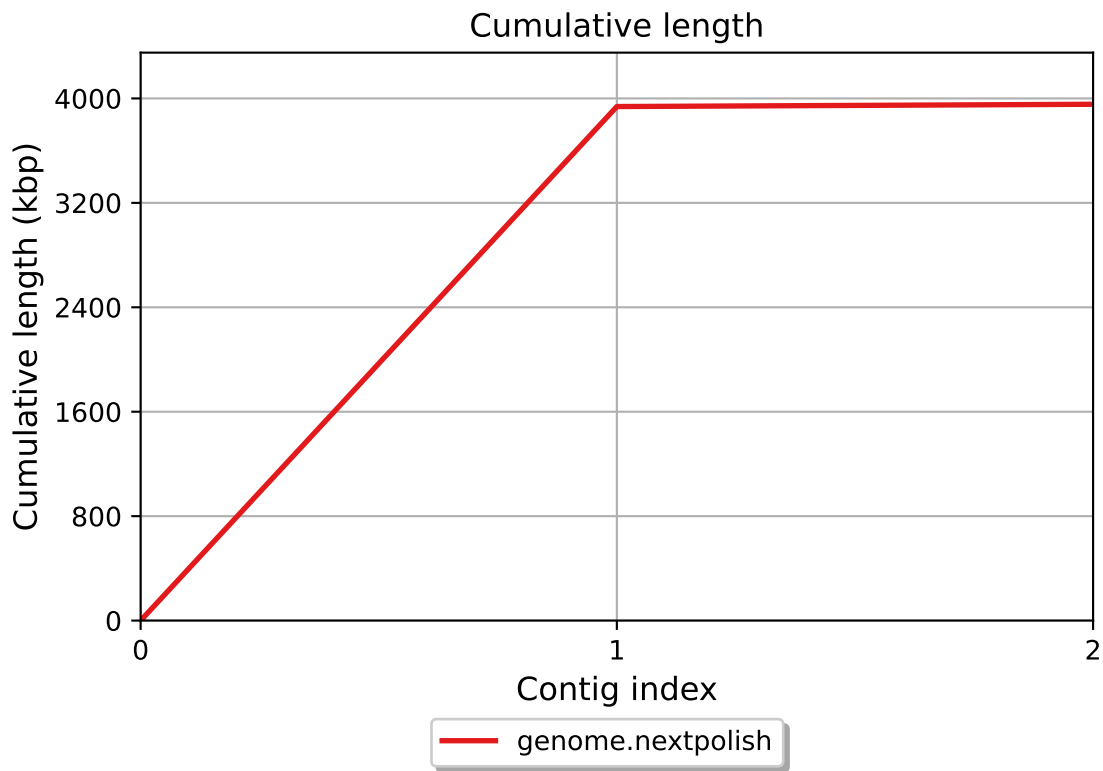


Report

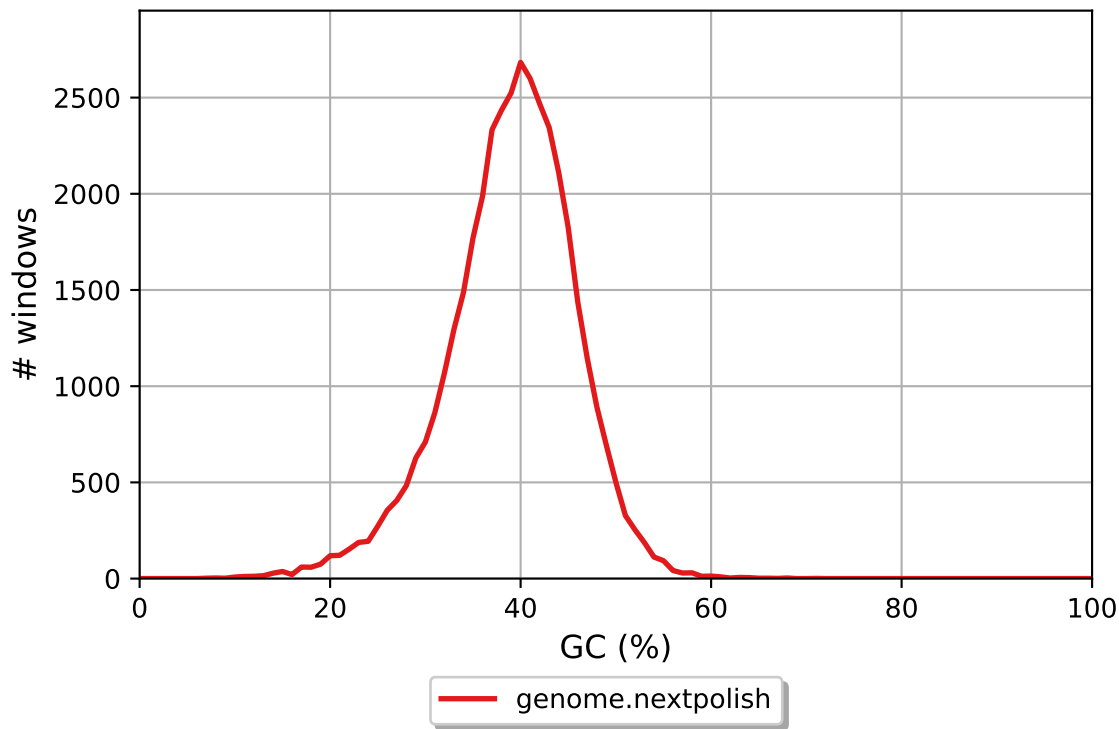
	genome.nextpolish
# contigs (>= 0 bp)	2
# contigs (>= 1000 bp)	2
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	3955283
Total length (>= 1000 bp)	3955283
Total length (>= 5000 bp)	3955283
Total length (>= 10000 bp)	3955283
Total length (>= 25000 bp)	3937839
Total length (>= 50000 bp)	3937839
# contigs	2
Largest contig	3937839
Total length	3955283
GC (%)	39.01
N50	3937839
N90	3937839
auN	3920548.9
L50	1
L90	1
# 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).





GC content



genome.nextpolish GC content

