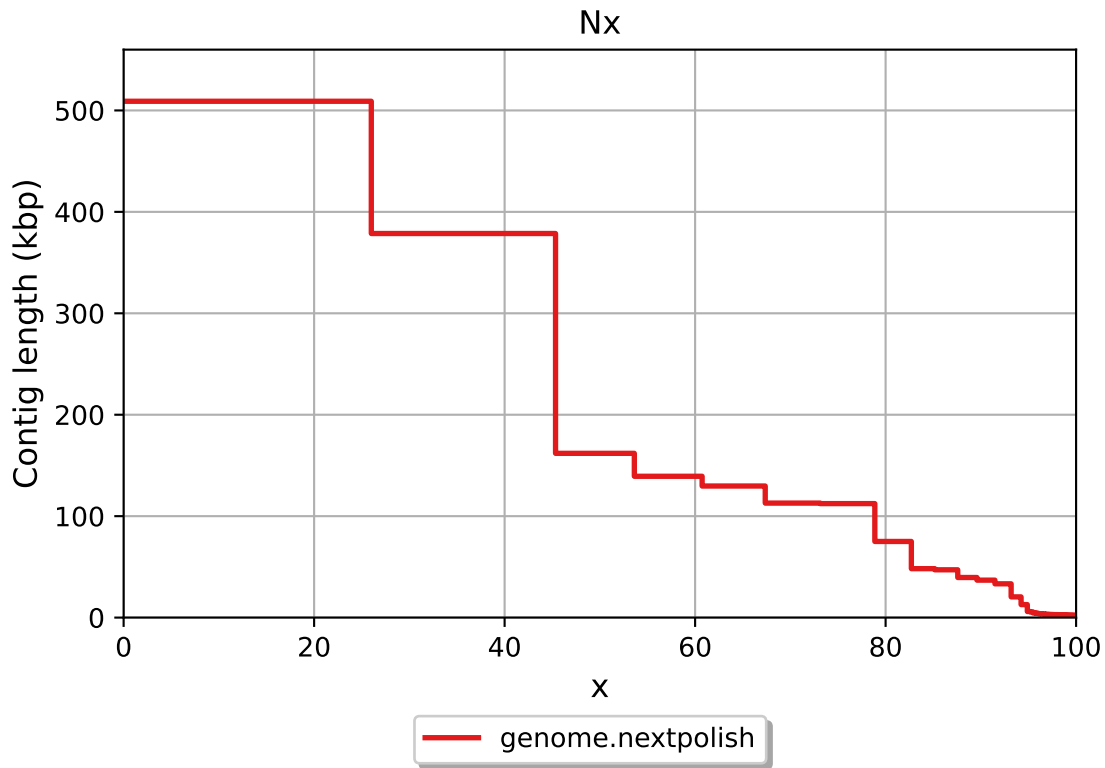
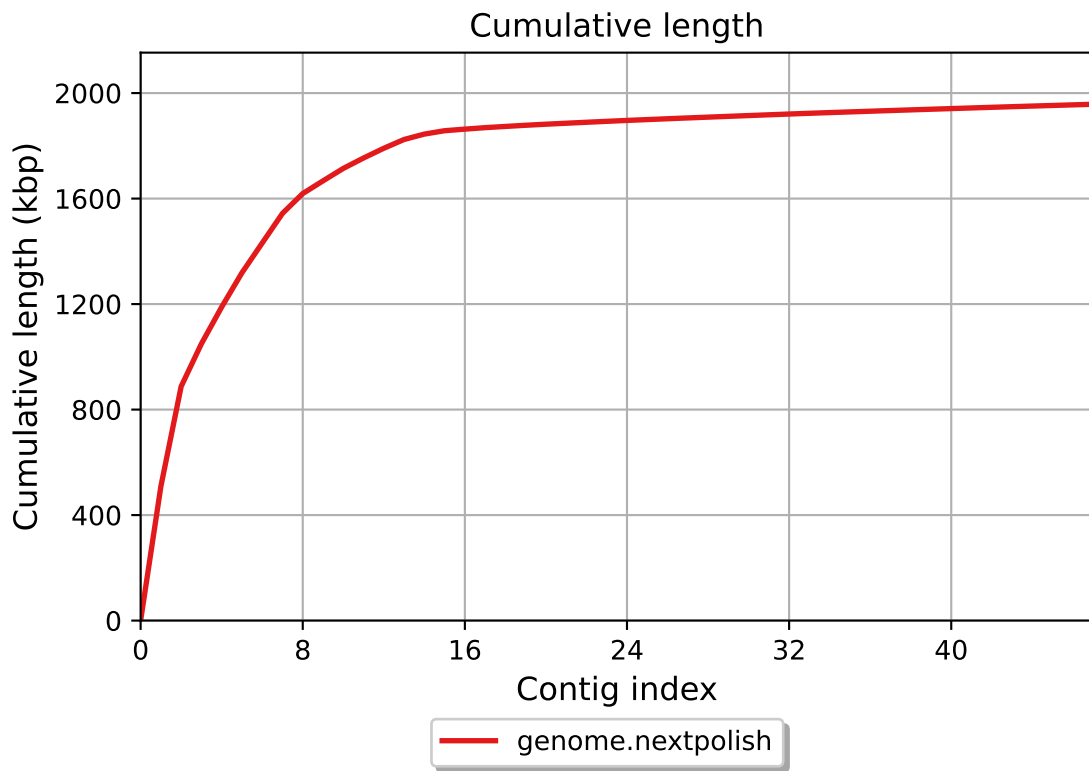


Report

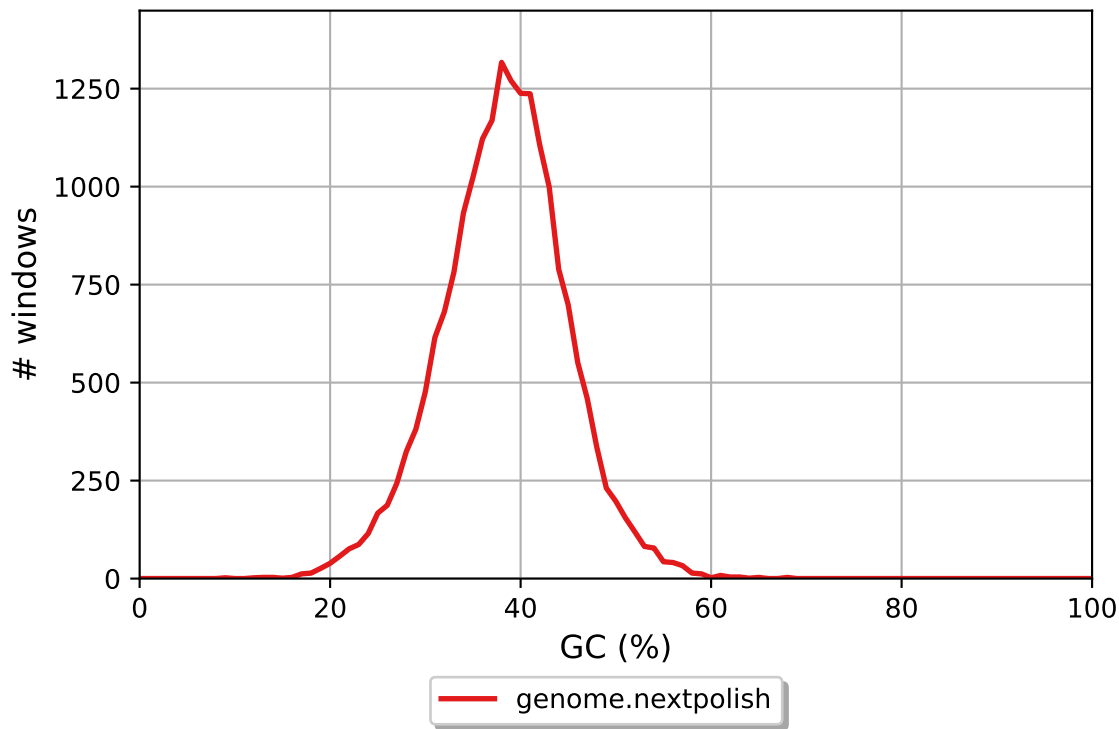
	genome.nextpolish
# contigs (>= 0 bp)	47
# contigs (>= 1000 bp)	47
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	1957564
Total length (>= 1000 bp)	1957564
Total length (>= 5000 bp)	1868924
Total length (>= 10000 bp)	1857303
Total length (>= 25000 bp)	1824057
Total length (>= 50000 bp)	1618967
# contigs	47
Largest contig	509039
Total length	1957564
GC (%)	38.28
N50	161936
N90	36911
auN	258211.7
L50	3
L90	12
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 2 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

