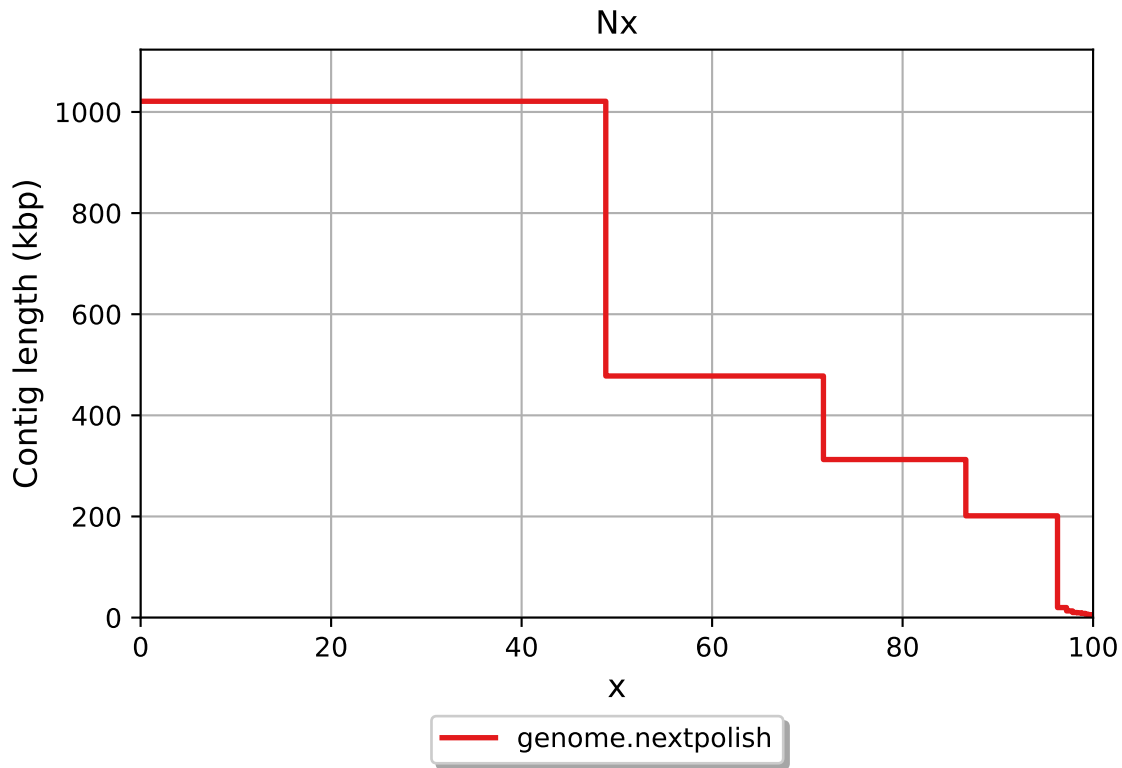
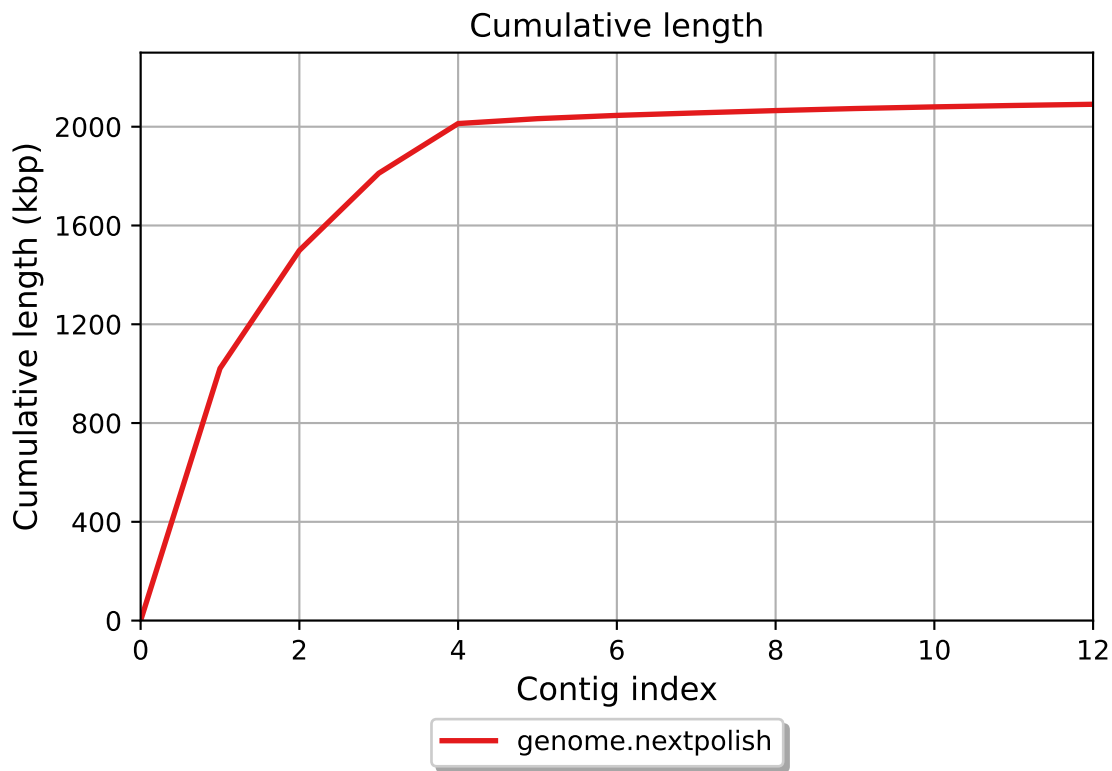


Report

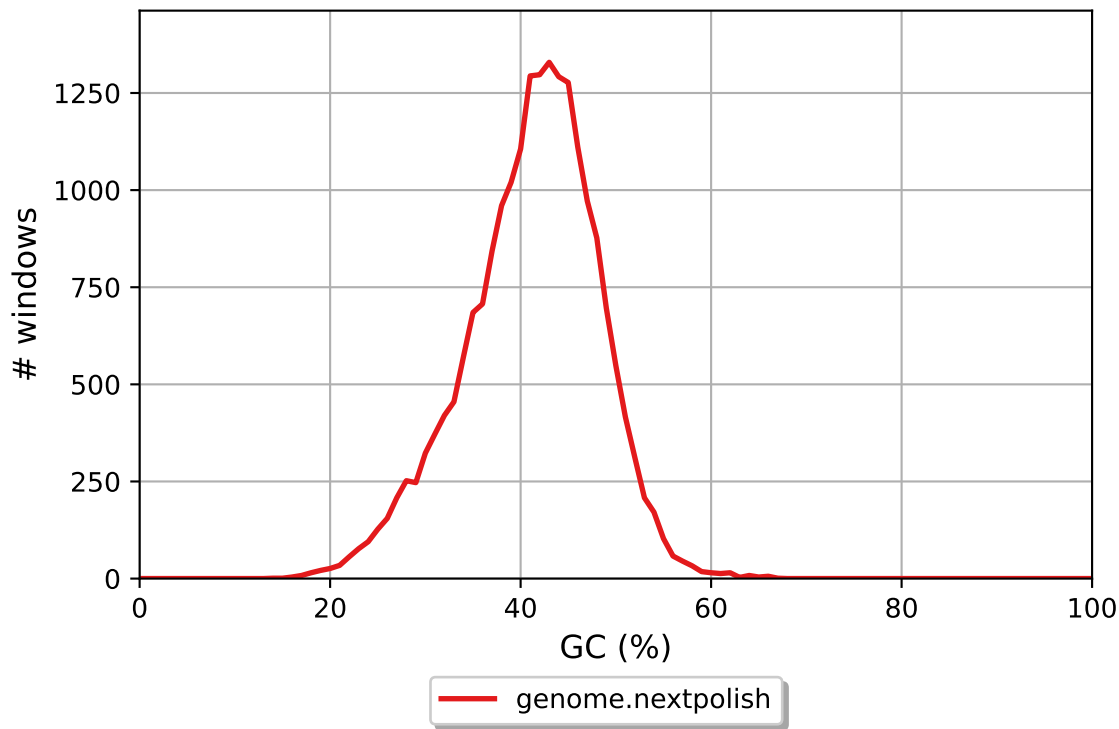
	genome.nextpolish
# contigs (>= 0 bp)	12
# contigs (>= 1000 bp)	12
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2090924
Total length (>= 1000 bp)	2090924
Total length (>= 5000 bp)	2086056
Total length (>= 10000 bp)	2055875
Total length (>= 25000 bp)	2012811
Total length (>= 50000 bp)	2012811
# contigs	12
Largest contig	1021152
Total length	2090924
GC (%)	41.14
N50	477811
N90	201266
auN	674437.8
L50	2
L90	4
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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

