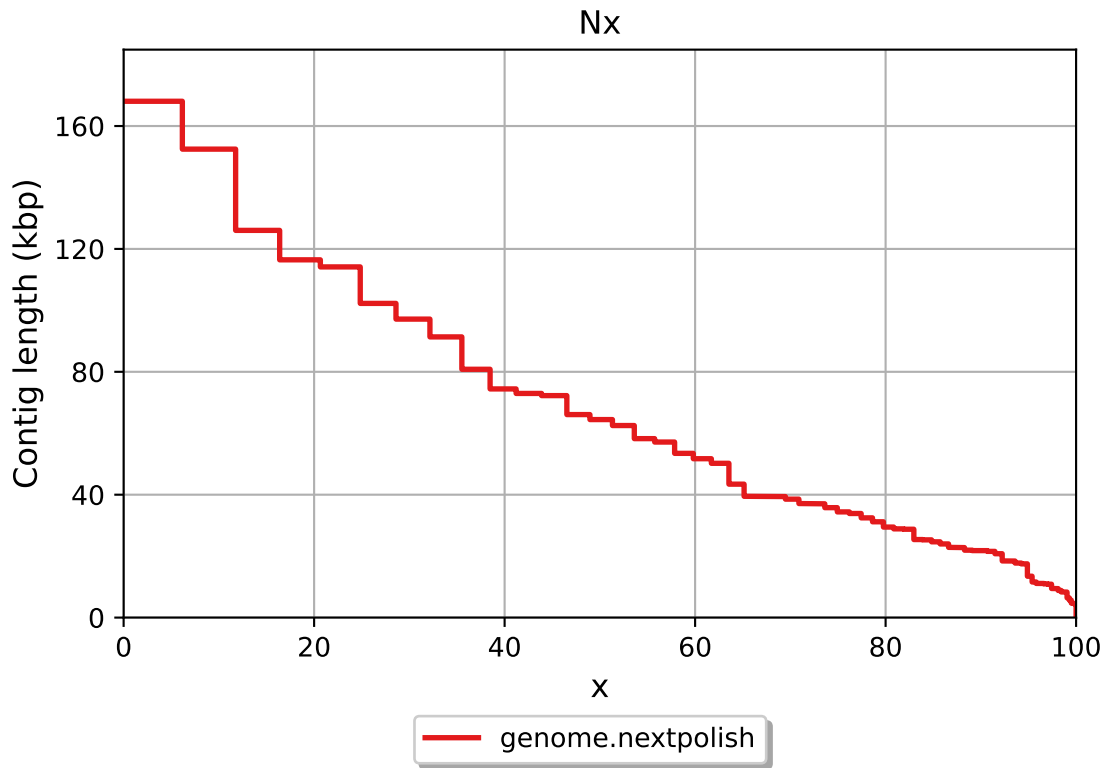
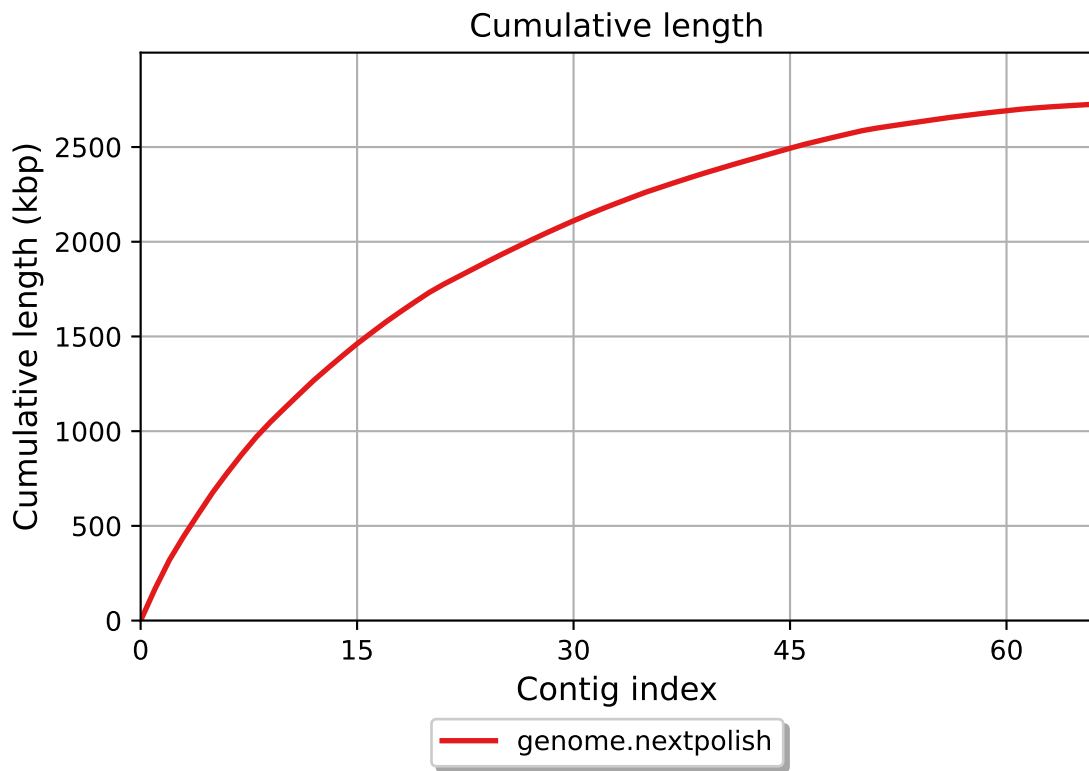


Report

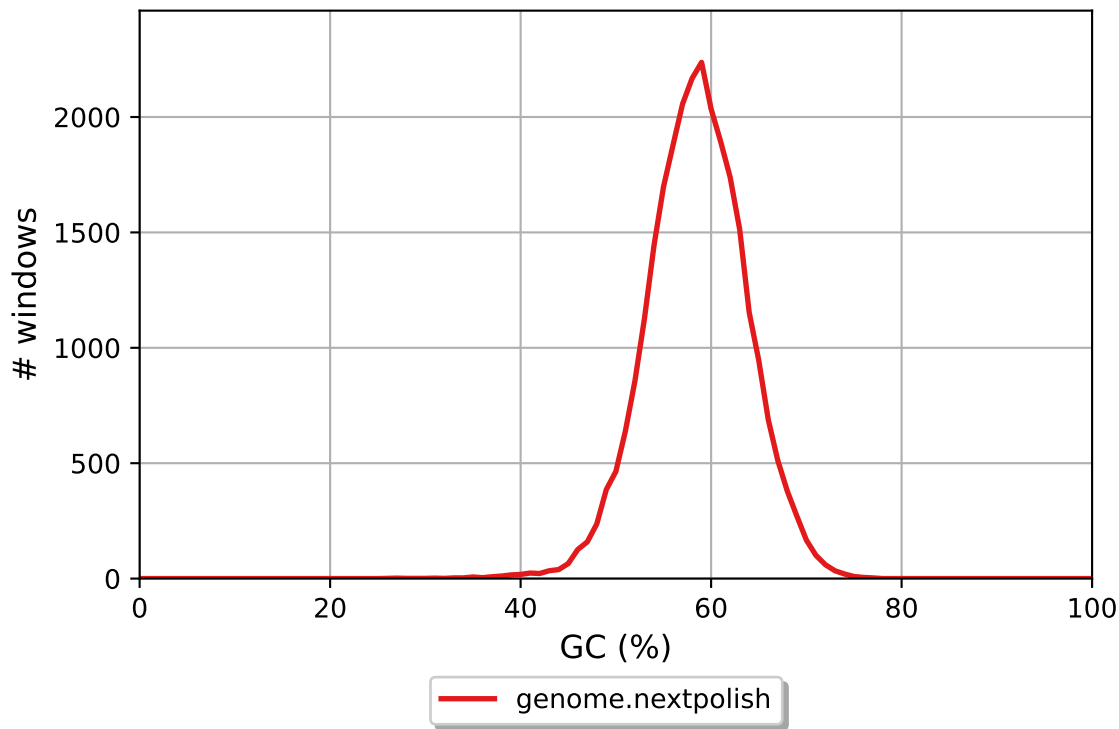
	genome.nextpolish
# contigs (>= 0 bp)	66
# contigs (>= 1000 bp)	66
# contigs (>= 5000 bp)	63
# contigs (>= 10000 bp)	56
# contigs (>= 25000 bp)	37
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	2725780
Total length (>= 1000 bp)	2725780
Total length (>= 5000 bp)	2712002
Total length (>= 10000 bp)	2655520
Total length (>= 25000 bp)	2312165
Total length (>= 50000 bp)	1732214
# contigs	66
Largest contig	168058
Total length	2725780
GC (%)	58.48
N50	64451
N90	21793
auN	73149.6
L50	14
L90	44
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 1 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

