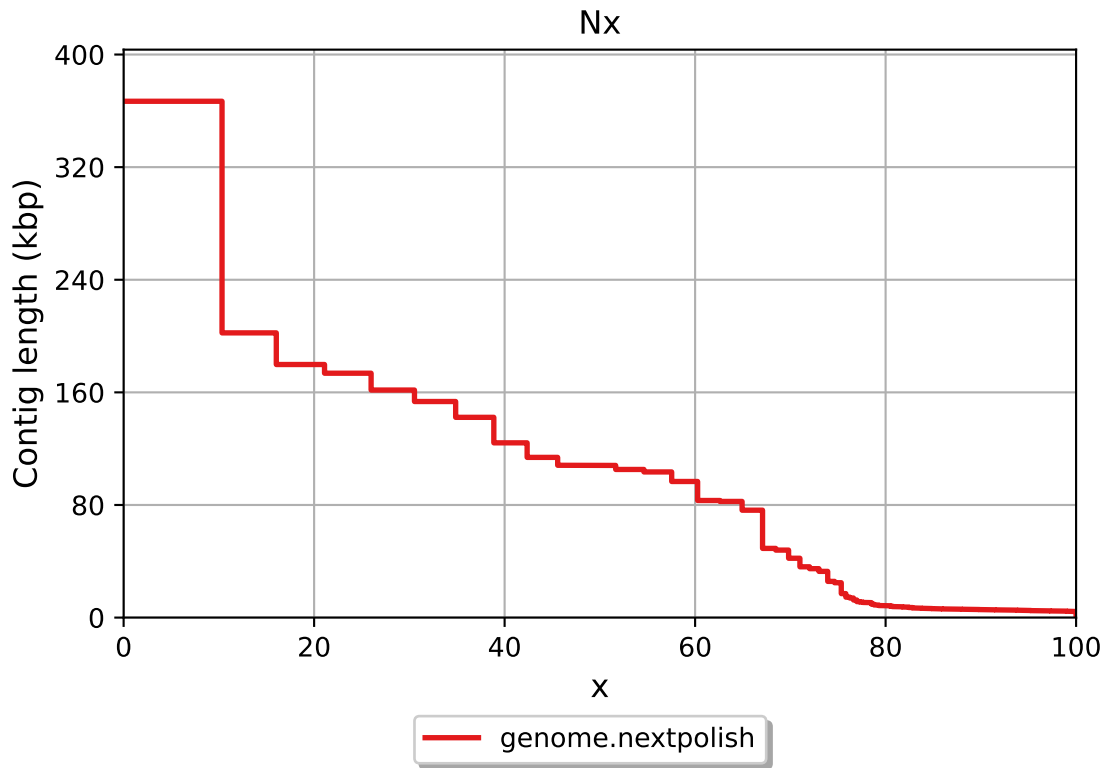
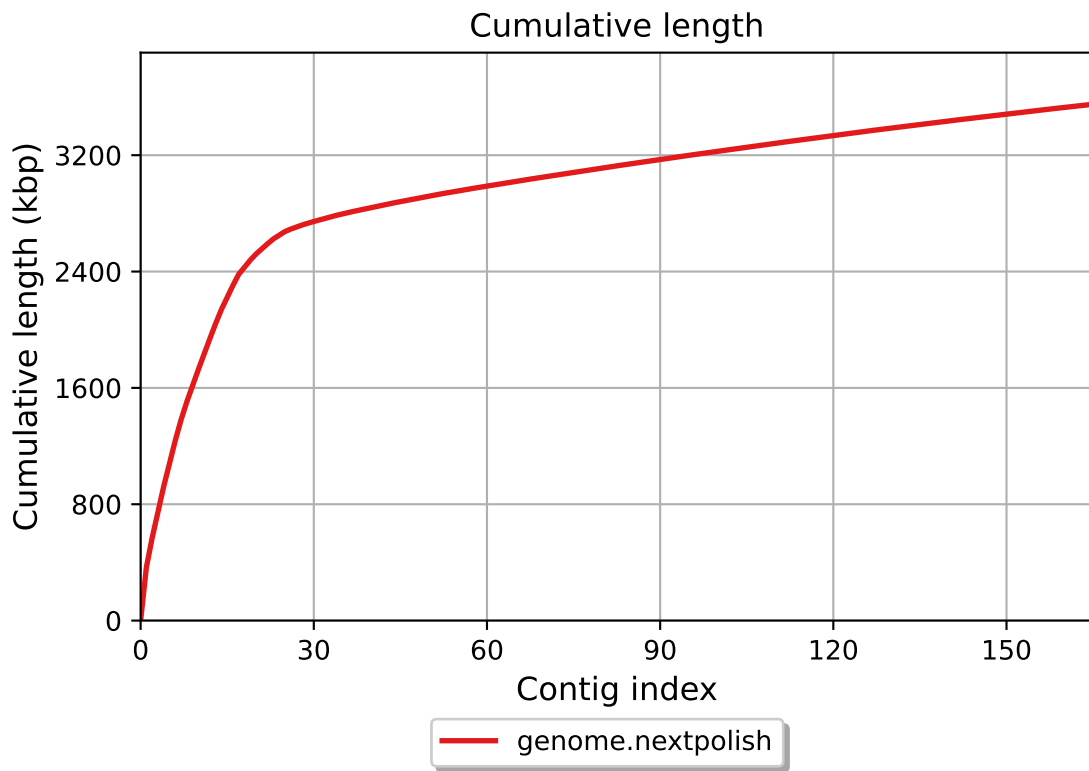


Report

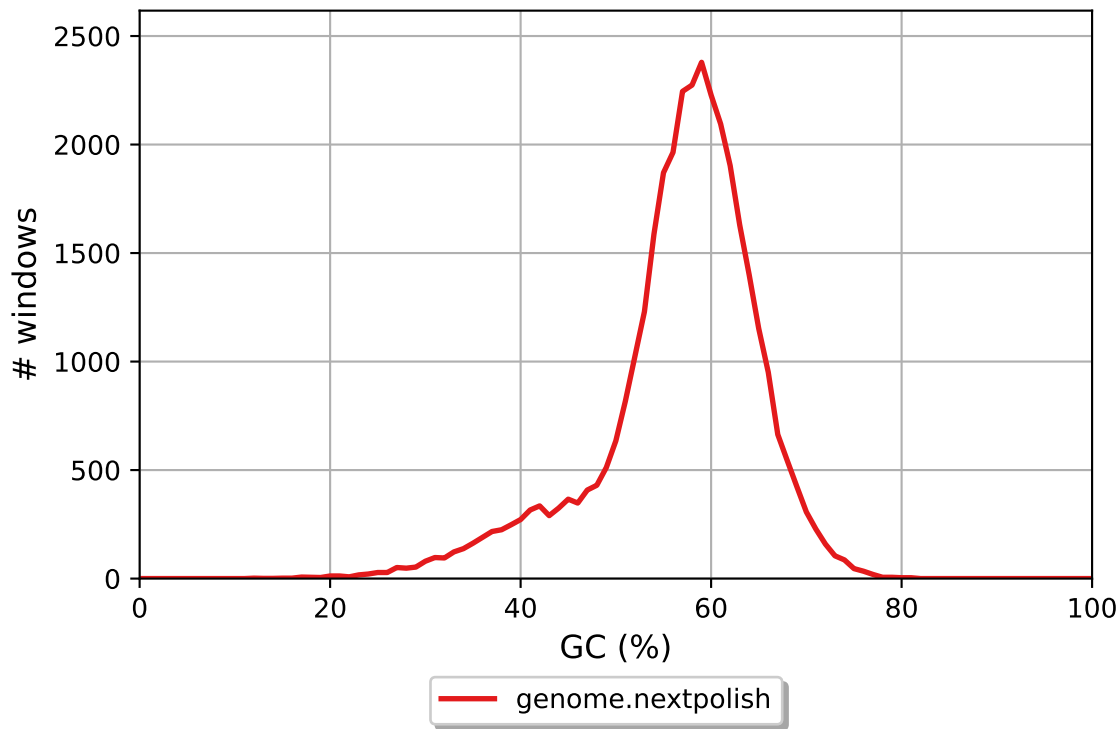
	genome.nextpolish
# contigs (>= 0 bp)	165
# contigs (>= 1000 bp)	165
# contigs (>= 5000 bp)	129
# contigs (>= 10000 bp)	34
# contigs (>= 25000 bp)	24
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	3550187
Total length (>= 1000 bp)	3550187
Total length (>= 5000 bp)	3381404
Total length (>= 10000 bp)	2787337
Total length (>= 25000 bp)	2650124
Total length (>= 50000 bp)	2381503
# contigs	165
Largest contig	366798
Total length	3550187
GC (%)	56.58
N50	108091
N90	5681
auN	120469.4
L50	11
L90	95
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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

