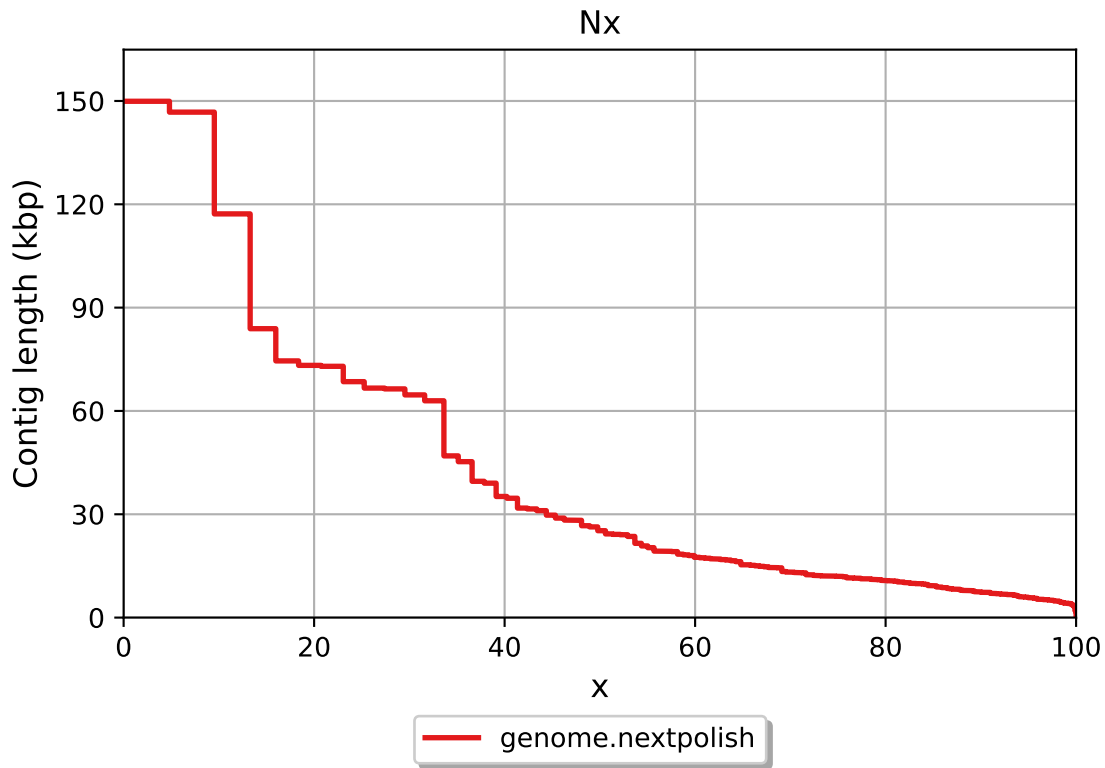
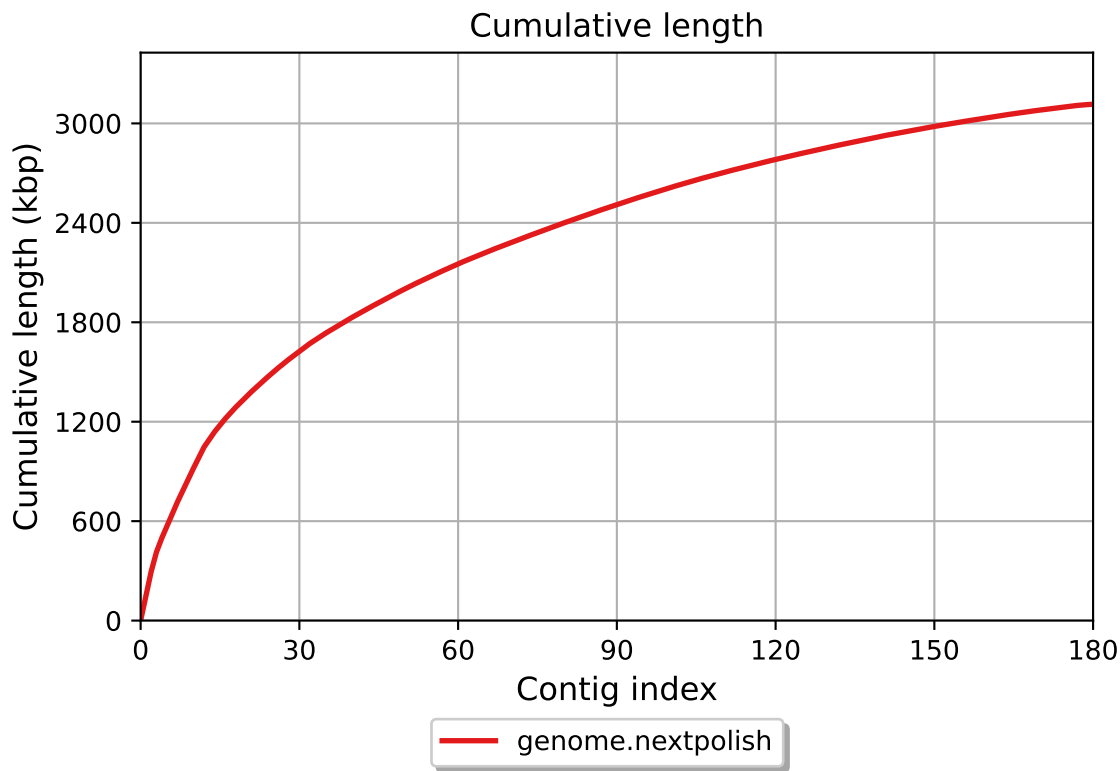


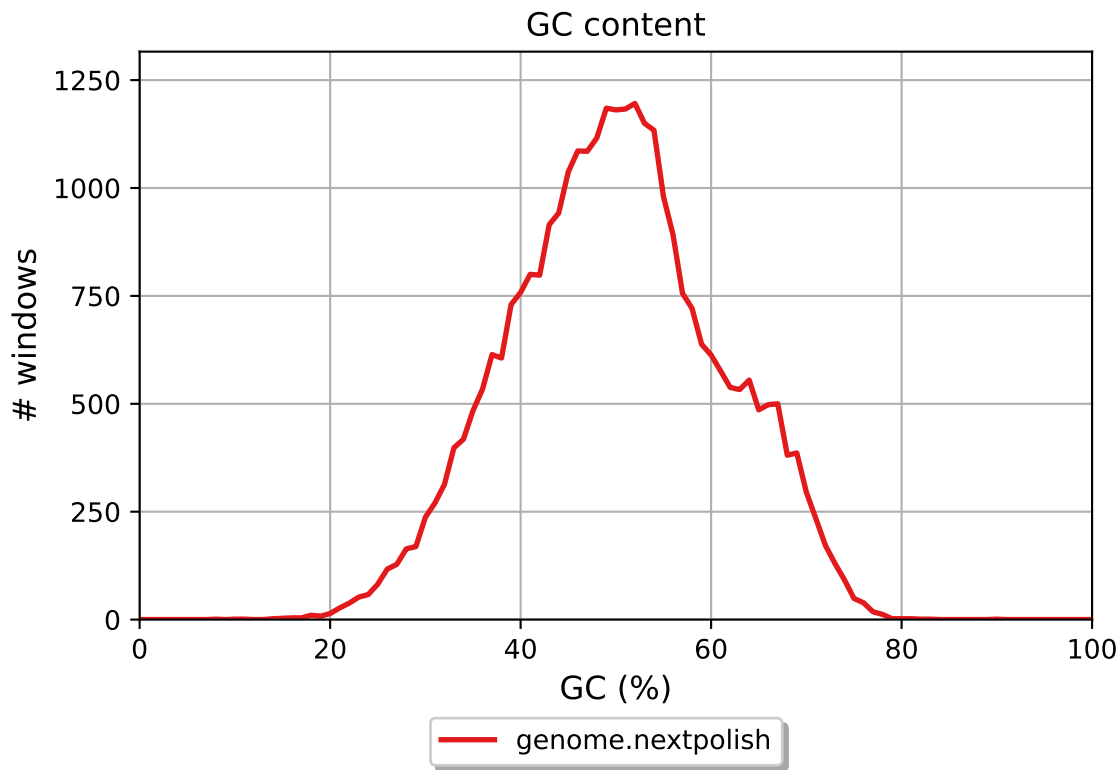
Report

	genome.nextpolish
# contigs (>= 0 bp)	180
# contigs (>= 1000 bp)	180
# contigs (>= 5000 bp)	161
# contigs (>= 10000 bp)	96
# contigs (>= 25000 bp)	28
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	3115456
Total length (>= 1000 bp)	3115456
Total length (>= 5000 bp)	3039288
Total length (>= 10000 bp)	2572216
Total length (>= 25000 bp)	1576069
Total length (>= 50000 bp)	1047624
# contigs	180
Largest contig	149913
Total length	3115456
GC (%)	49.86
N50	25229
N90	7455
auN	45008.2
L50	28
L90	123
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).







genome.nextpolish GC content

