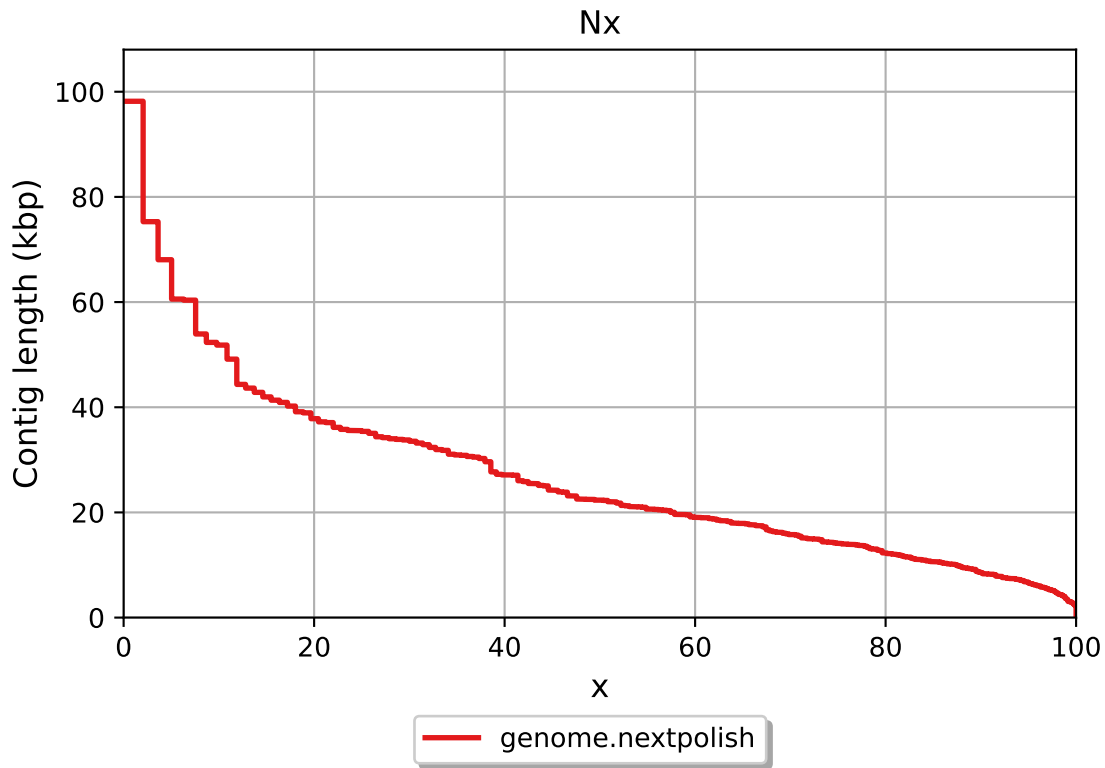
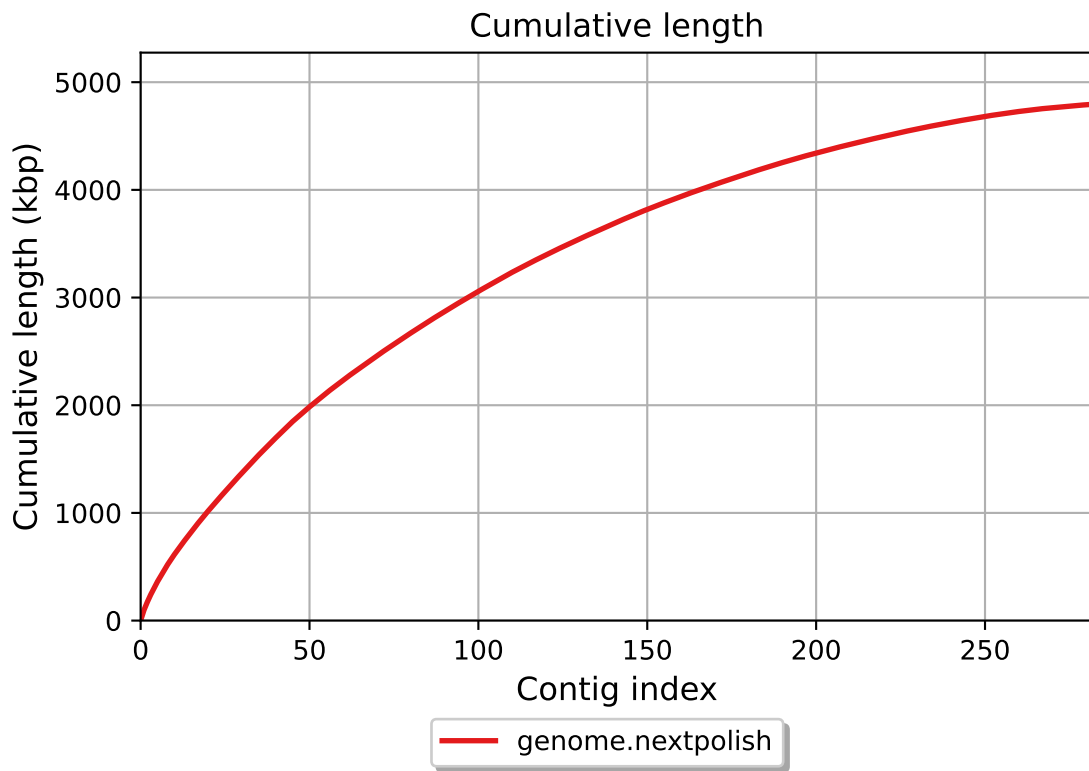


Report

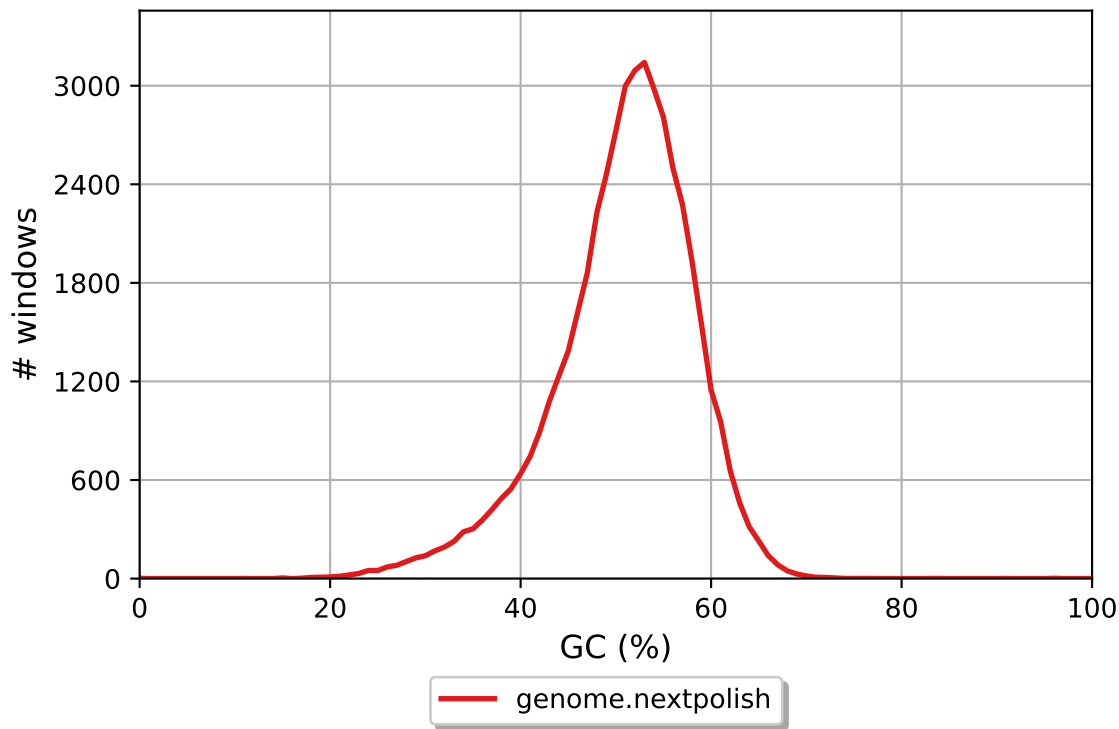
	genome.nextpolish
# contigs (>= 0 bp)	282
# contigs (>= 1000 bp)	282
# contigs (>= 5000 bp)	251
# contigs (>= 10000 bp)	184
# contigs (>= 25000 bp)	56
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	4794948
Total length (>= 1000 bp)	4794948
Total length (>= 5000 bp)	4687040
Total length (>= 10000 bp)	4195973
Total length (>= 25000 bp)	2137913
Total length (>= 50000 bp)	520552
# contigs	282
Largest contig	98186
Total length	4794948
GC (%)	50.73
N50	22332
N90	8612
auN	27328.1
L50	68
L90	197
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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

