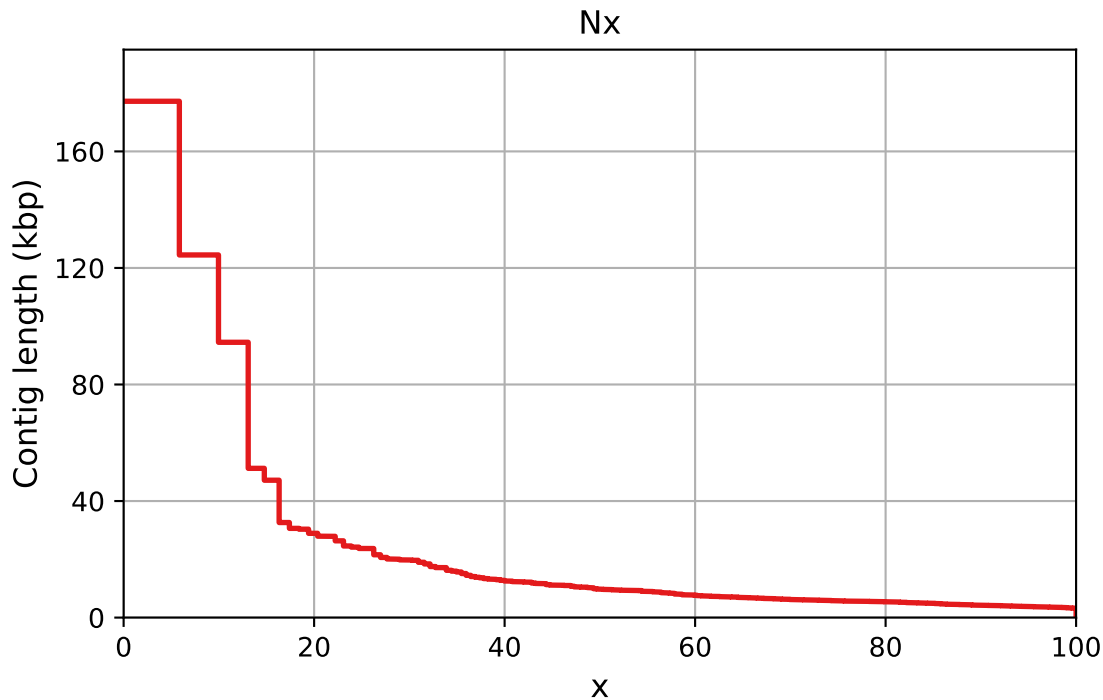


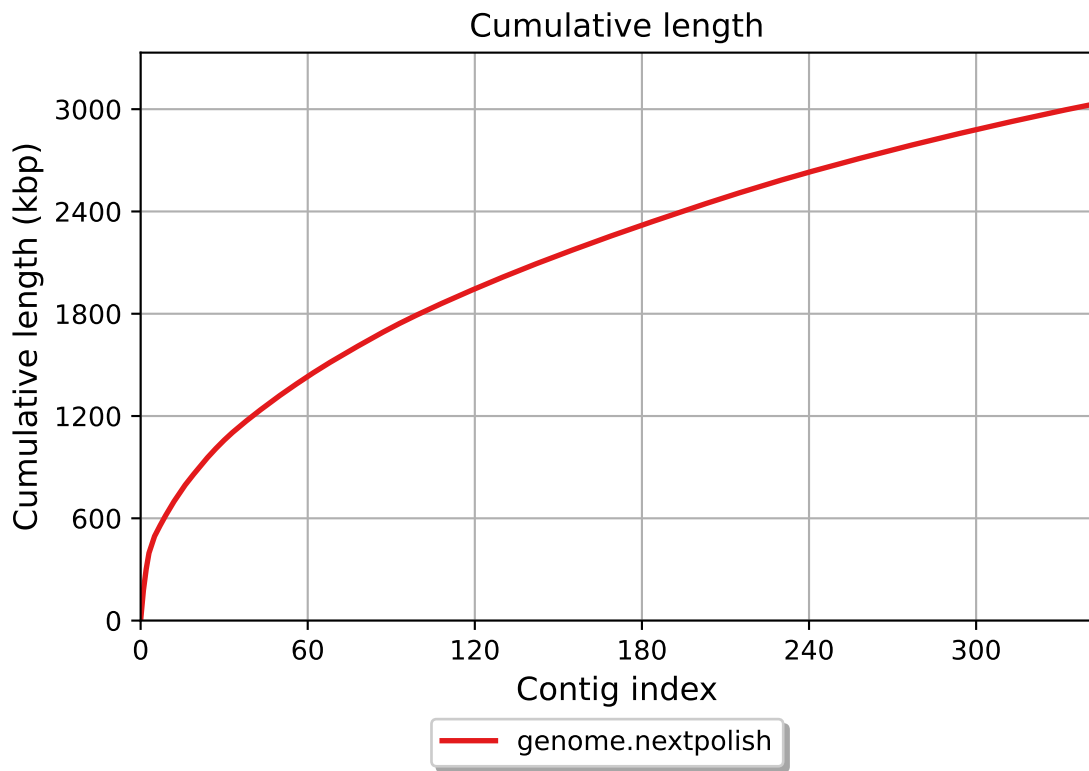
Report

	genome.nextpolish
# contigs (>= 0 bp)	342
# contigs (>= 1000 bp)	342
# contigs (>= 5000 bp)	220
# contigs (>= 10000 bp)	66
# contigs (>= 25000 bp)	12
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3028815
Total length (>= 1000 bp)	3028815
Total length (>= 5000 bp)	2534605
Total length (>= 10000 bp)	1494940
Total length (>= 25000 bp)	699191
Total length (>= 50000 bp)	447377
# contigs	342
Largest contig	177207
Total length	3028815
GC (%)	63.15
N50	9794
N90	4236
auN	29275.6
L50	68
L90	262
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

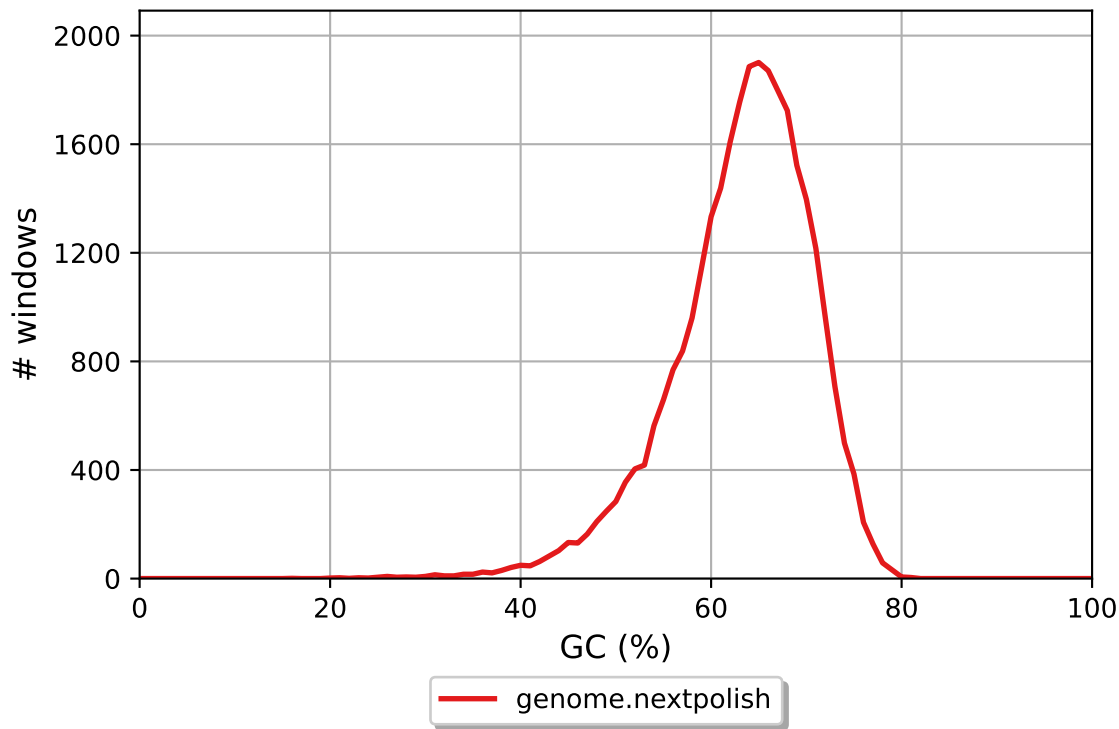
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

