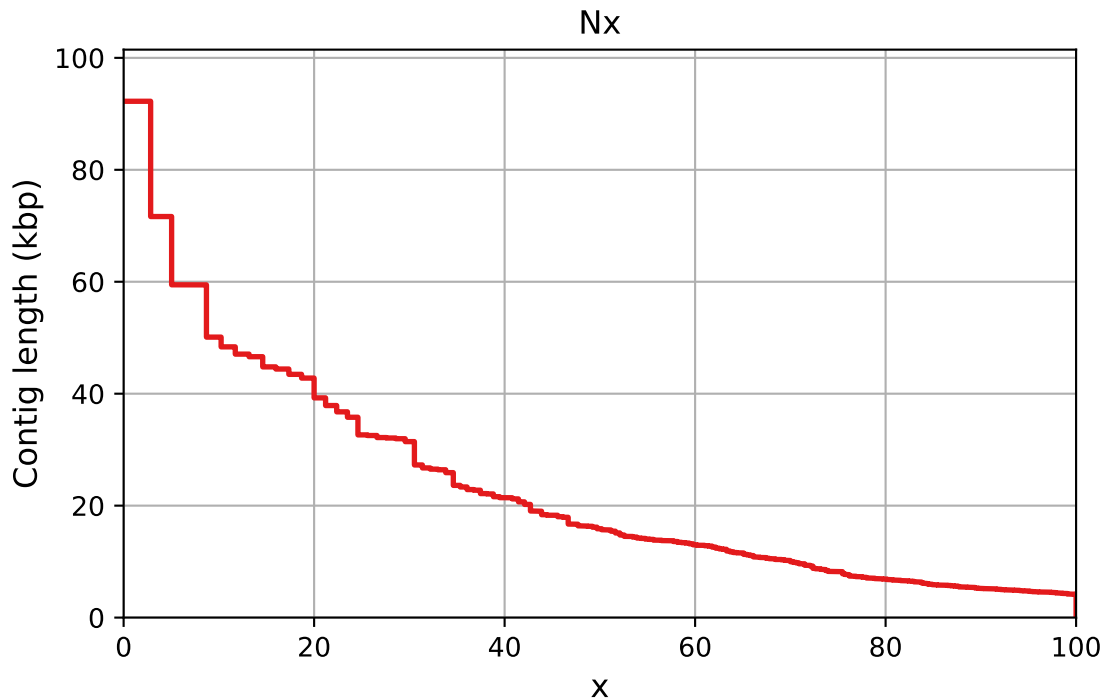


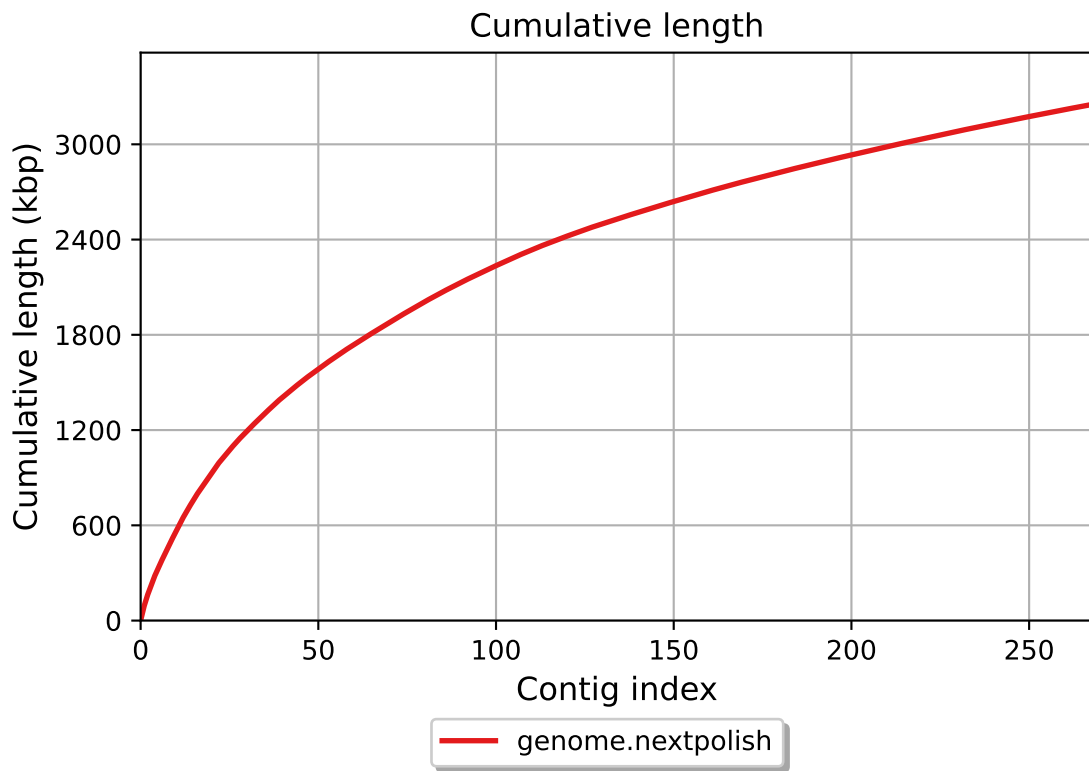
Report

	genome.nextpolish
# contigs (>= 0 bp)	268
# contigs (>= 1000 bp)	268
# contigs (>= 5000 bp)	213
# contigs (>= 10000 bp)	104
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	3252228
Total length (>= 1000 bp)	3252228
Total length (>= 5000 bp)	2999858
Total length (>= 10000 bp)	2276742
Total length (>= 25000 bp)	1125719
Total length (>= 50000 bp)	332897
# contigs	268
Largest contig	92244
Total length	3252228
GC (%)	56.24
N50	15873
N90	5227
auN	23810.8
L50	53
L90	199
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 0 part

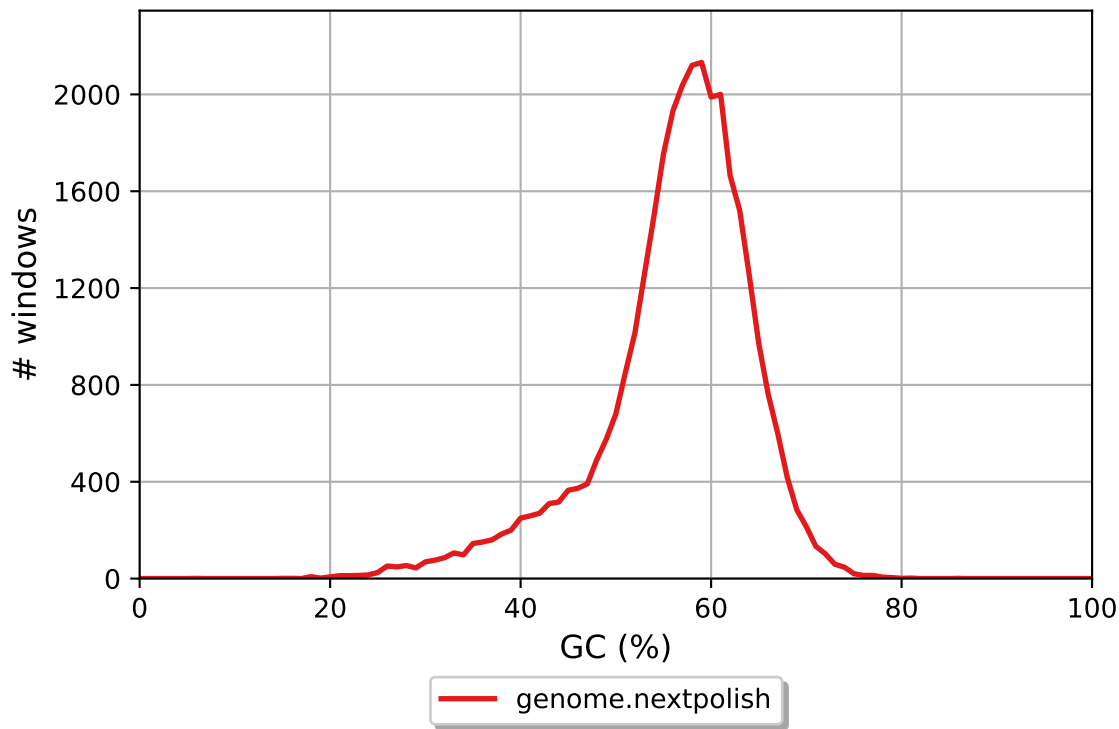
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

