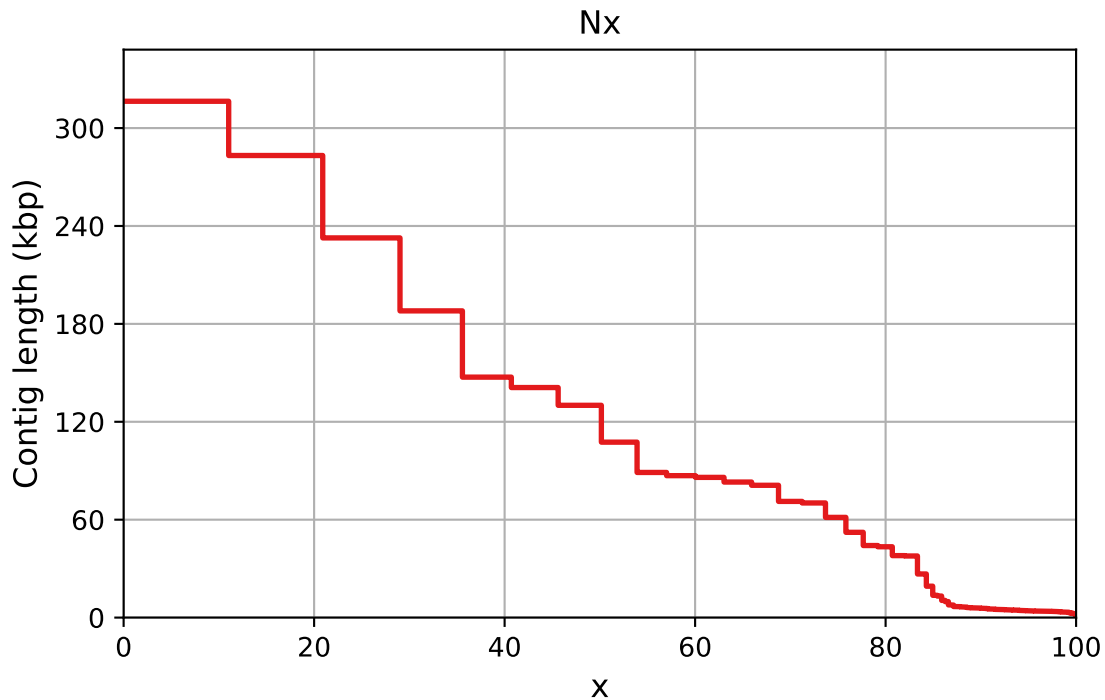


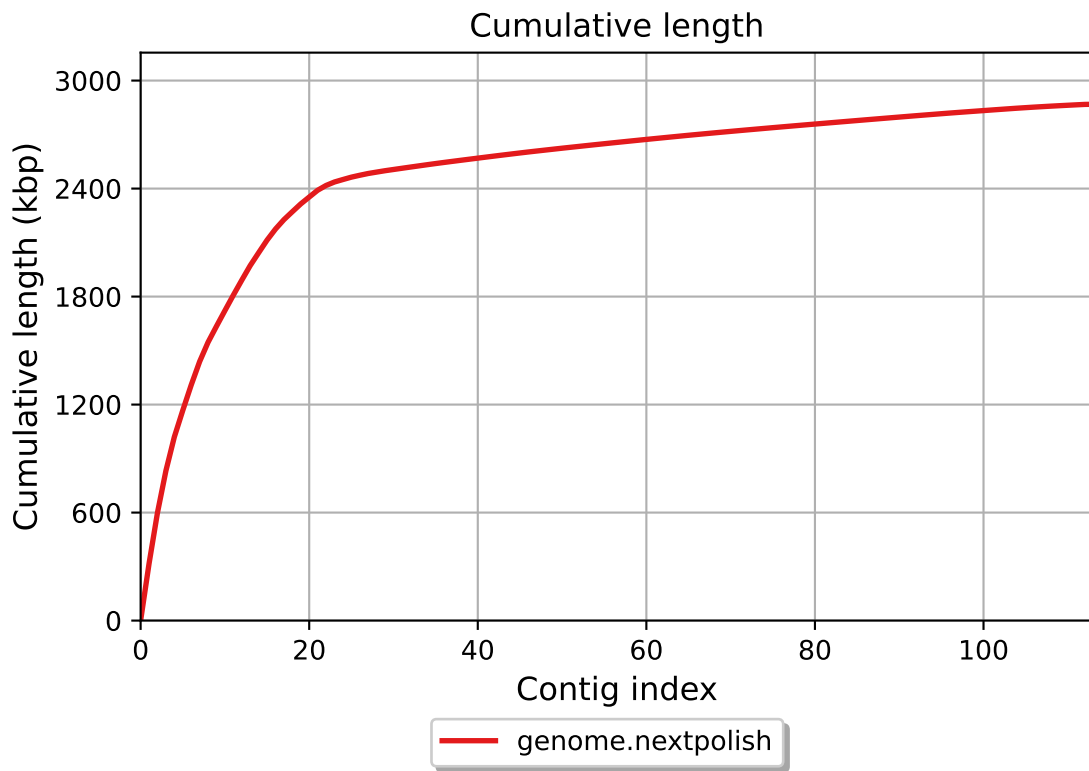
Report

	genome.nextpolish
# contigs (>= 0 bp)	113
# contigs (>= 1000 bp)	112
# contigs (>= 5000 bp)	52
# contigs (>= 10000 bp)	26
# contigs (>= 25000 bp)	22
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	2868358
Total length (>= 1000 bp)	2867566
Total length (>= 5000 bp)	2634322
Total length (>= 10000 bp)	2473977
Total length (>= 25000 bp)	2417358
Total length (>= 50000 bp)	2227331
# contigs	113
Largest contig	316426
Total length	2868358
GC (%)	34.83
N50	130097
N90	5746
auN	140470.6
L50	7
L90	43
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 1 part

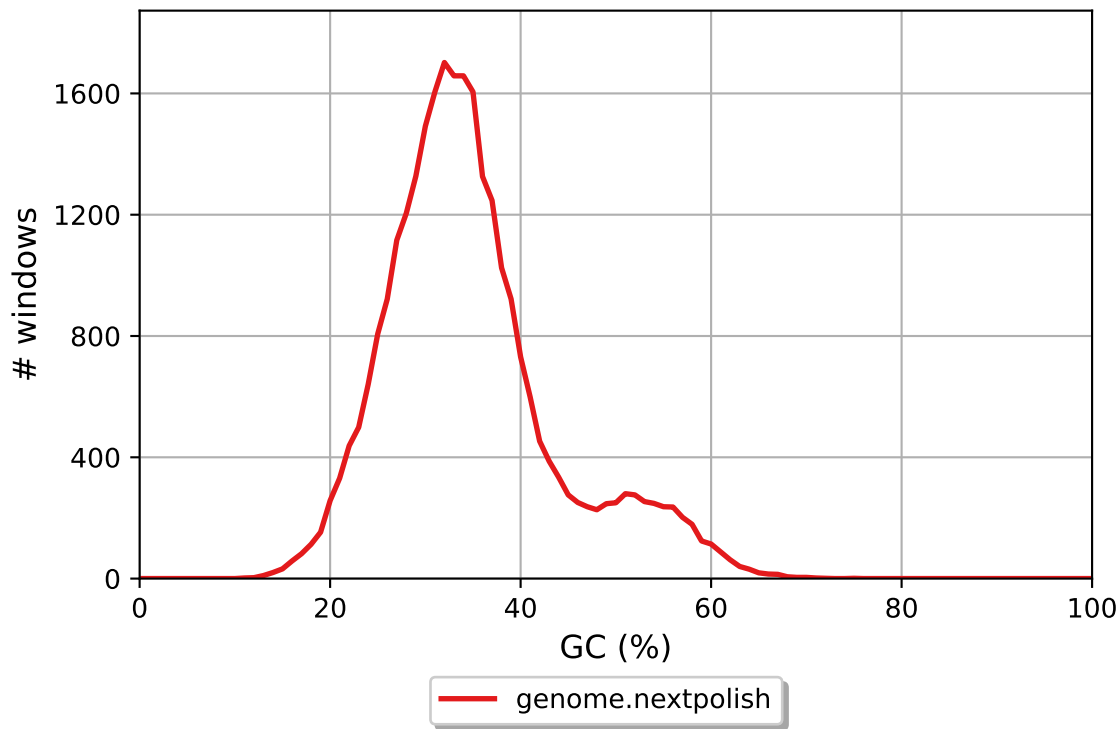
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

