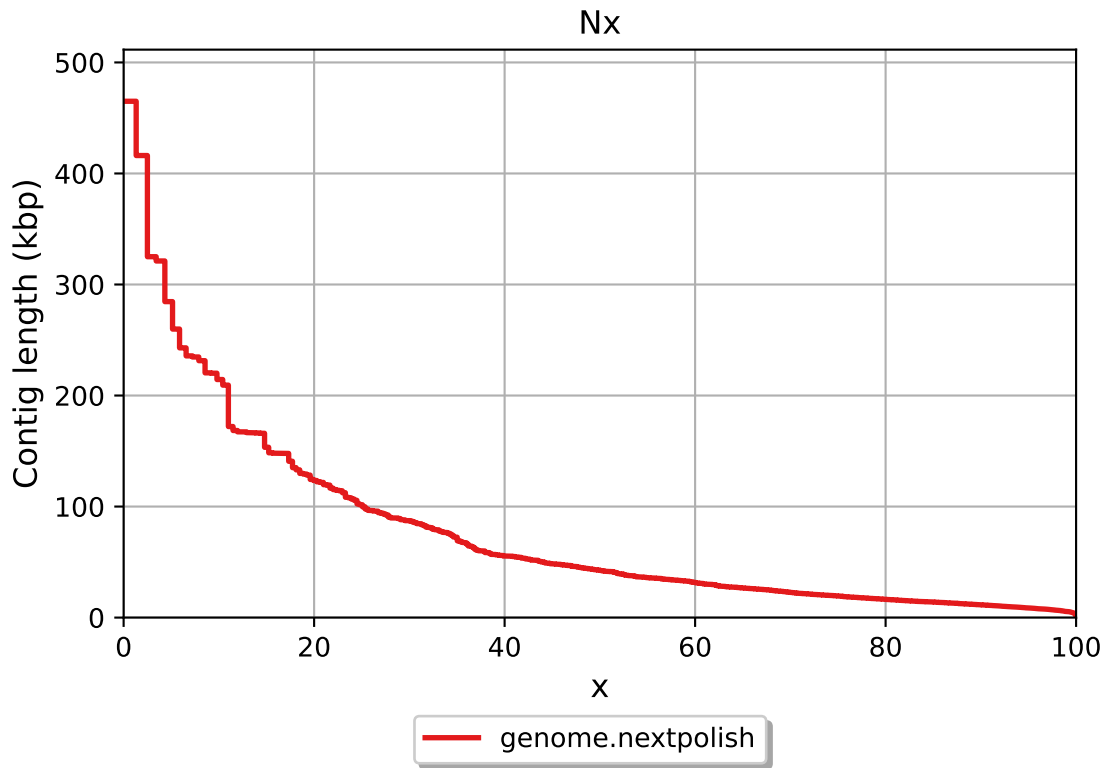


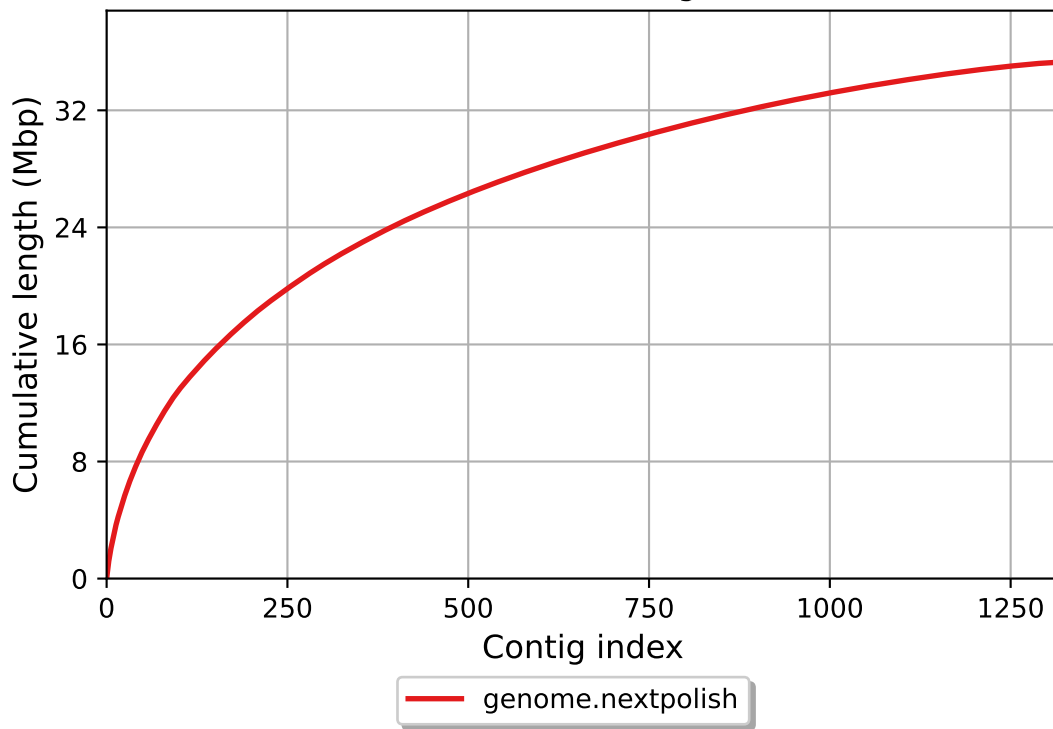
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1317
# contigs (>= 1000 bp)	1313
# contigs (>= 5000 bp)	1261
# contigs (>= 10000 bp)	949
# contigs (>= 25000 bp)	388
# contigs (>= 50000 bp)	148
Total length (>= 0 bp)	35283785
Total length (>= 1000 bp)	35280584
Total length (>= 5000 bp)	35076958
Total length (>= 10000 bp)	32697617
Total length (>= 25000 bp)	23874788
Total length (>= 50000 bp)	15558482
# contigs	1317
Largest contig	465026
Total length	35283785
GC (%)	48.00
N50	42537
N90	11473
auN	79262.1
L50	194
L90	862
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	92 + 8 part

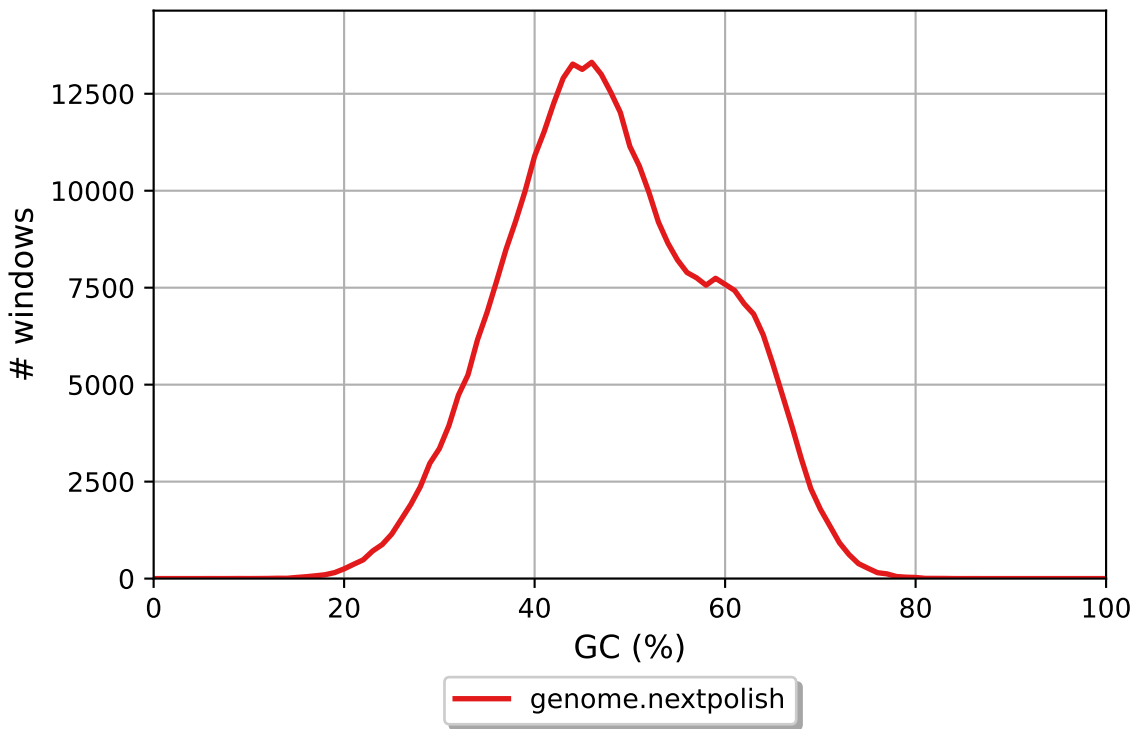
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

