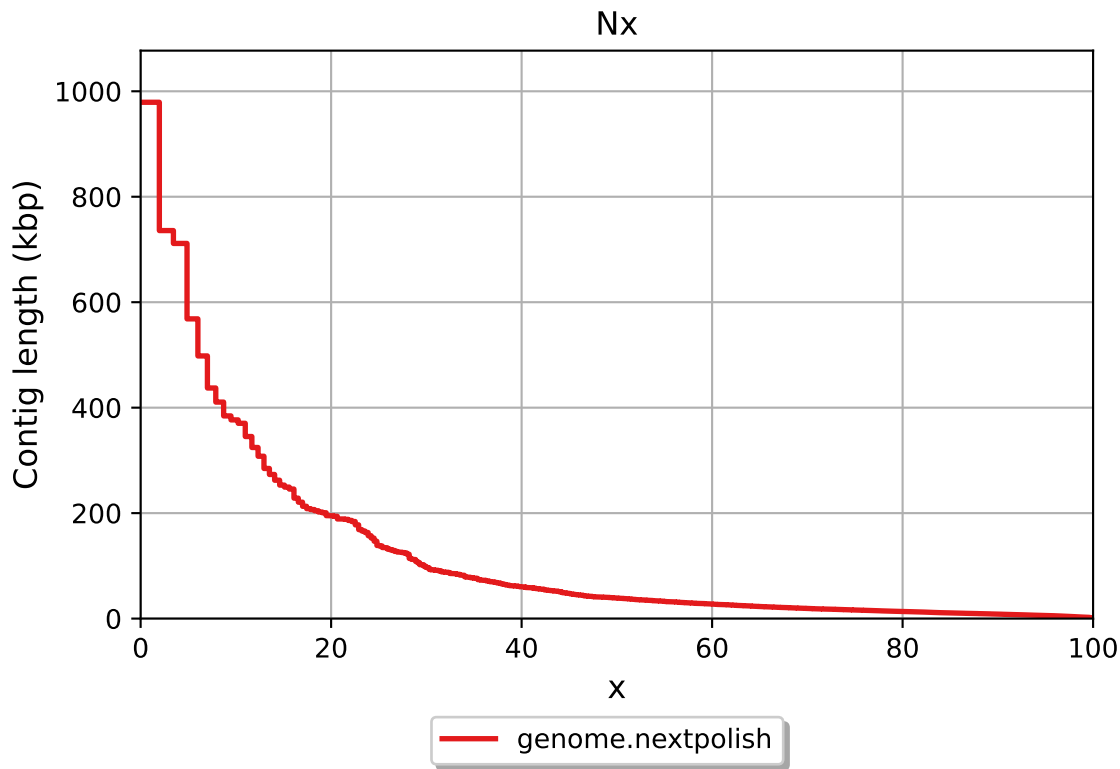


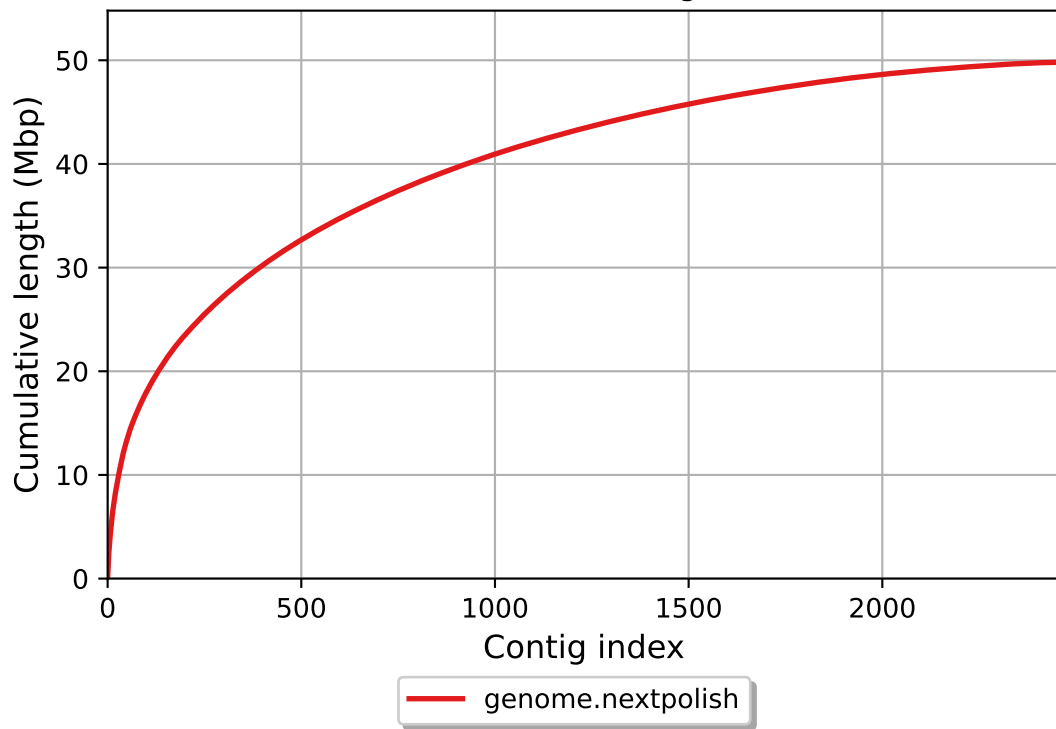
# Report

	genome.nextpolish
# contigs (>= 0 bp)	2460
# contigs (>= 1000 bp)	2424
# contigs (>= 5000 bp)	1859
# contigs (>= 10000 bp)	1200
# contigs (>= 25000 bp)	436
# contigs (>= 50000 bp)	167
Total length (>= 0 bp)	49808475
Total length (>= 1000 bp)	49783096
Total length (>= 5000 bp)	48006594
Total length (>= 10000 bp)	43157394
Total length (>= 25000 bp)	31162688
Total length (>= 50000 bp)	22009712
# contigs	2459
Largest contig	979155
Total length	49808002
GC (%)	44.63
N50	38940
N90	8487
auN	126585.0
L50	235
L90	1382
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	132 + 7 part

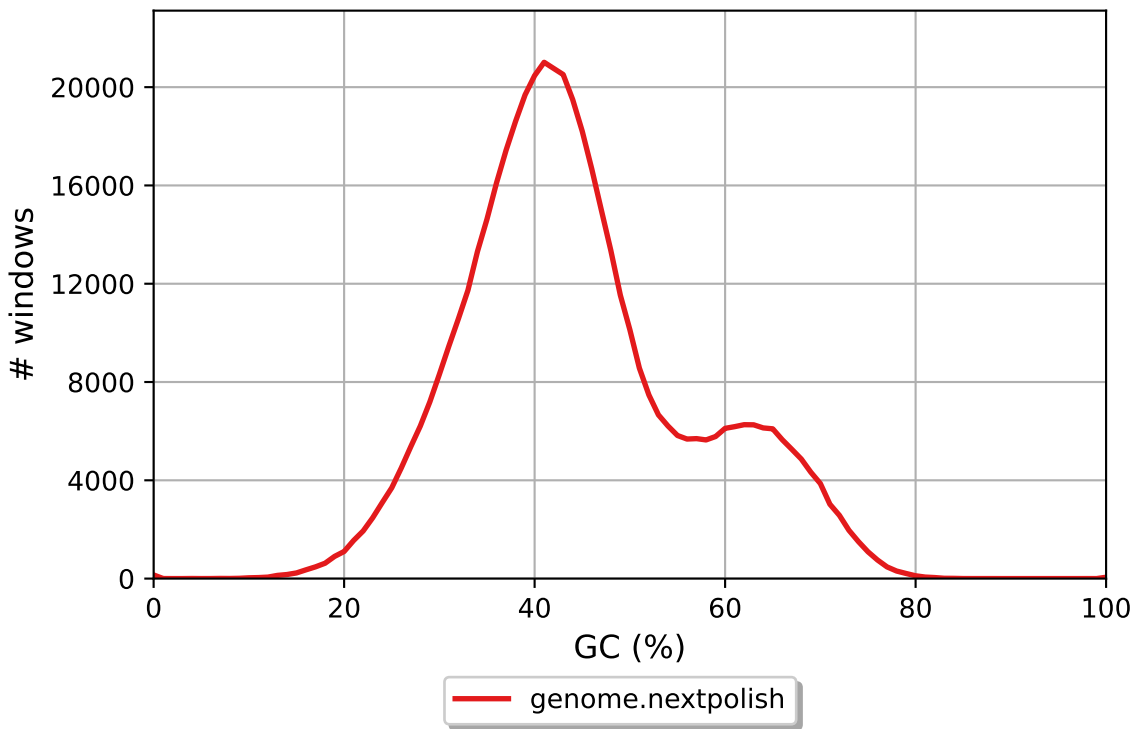
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

