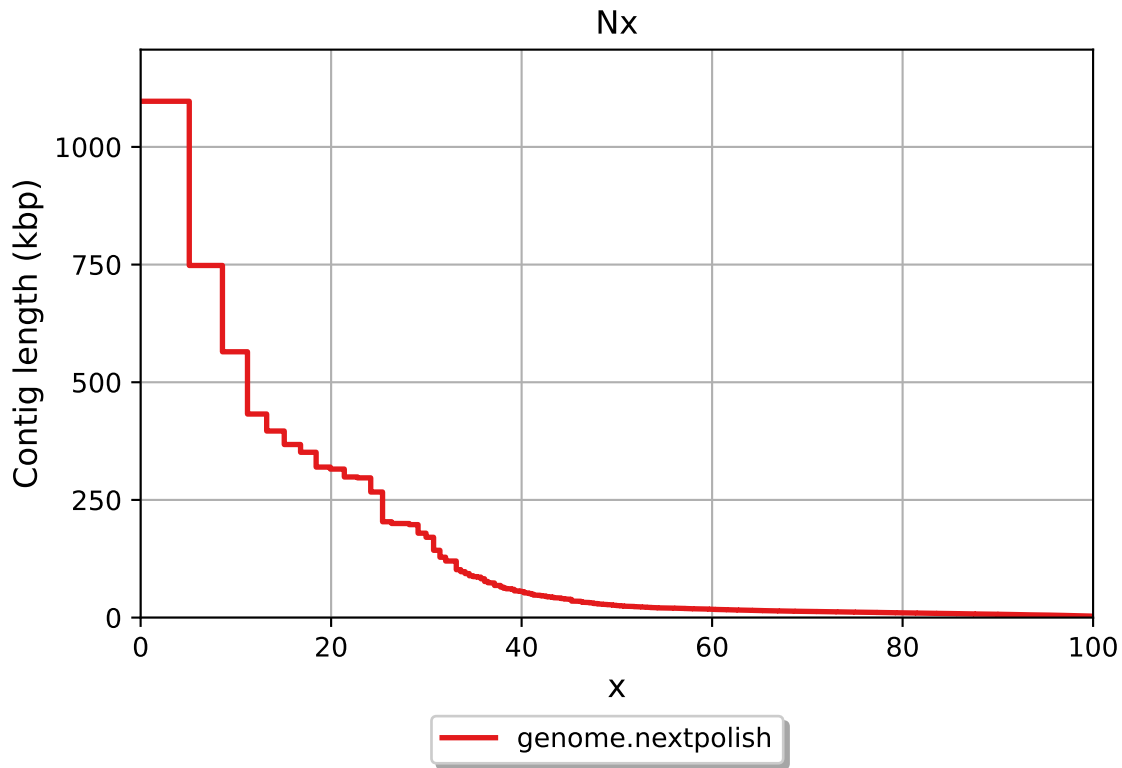


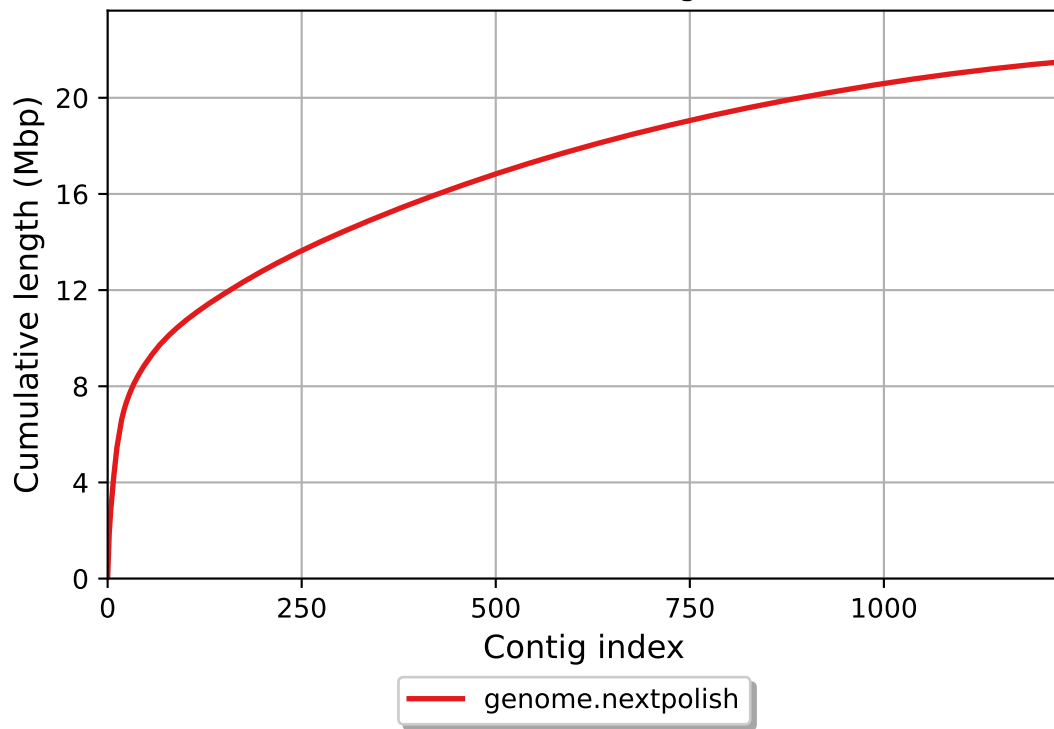
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1227
# contigs (>= 1000 bp)	1225
# contigs (>= 5000 bp)	1005
# contigs (>= 10000 bp)	541
# contigs (>= 25000 bp)	105
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	21477594
Total length (>= 1000 bp)	21476360
Total length (>= 5000 bp)	20620653
Total length (>= 10000 bp)	17255833
Total length (>= 25000 bp)	10846497
Total length (>= 50000 bp)	8805438
# contigs	1227
Largest contig	1096976
Total length	21477594
GC (%)	46.06
N50	25707
N90	7082
auN	174582.2
L50	101
L90	788
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	49 + 7 part

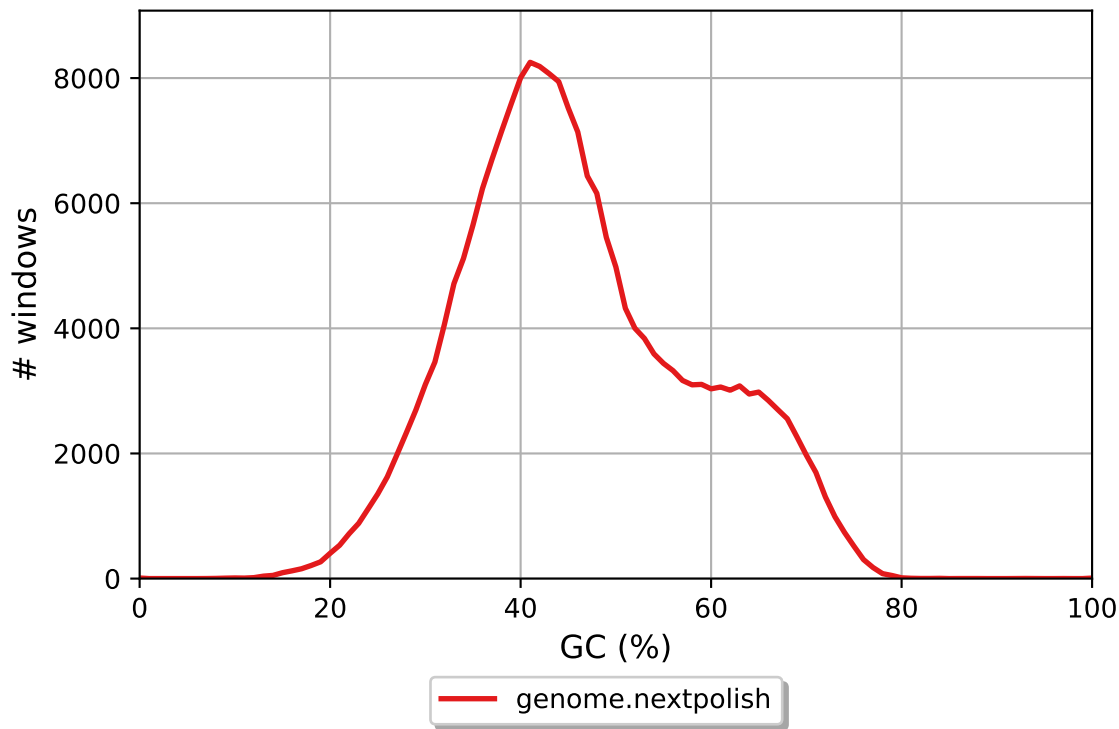
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

