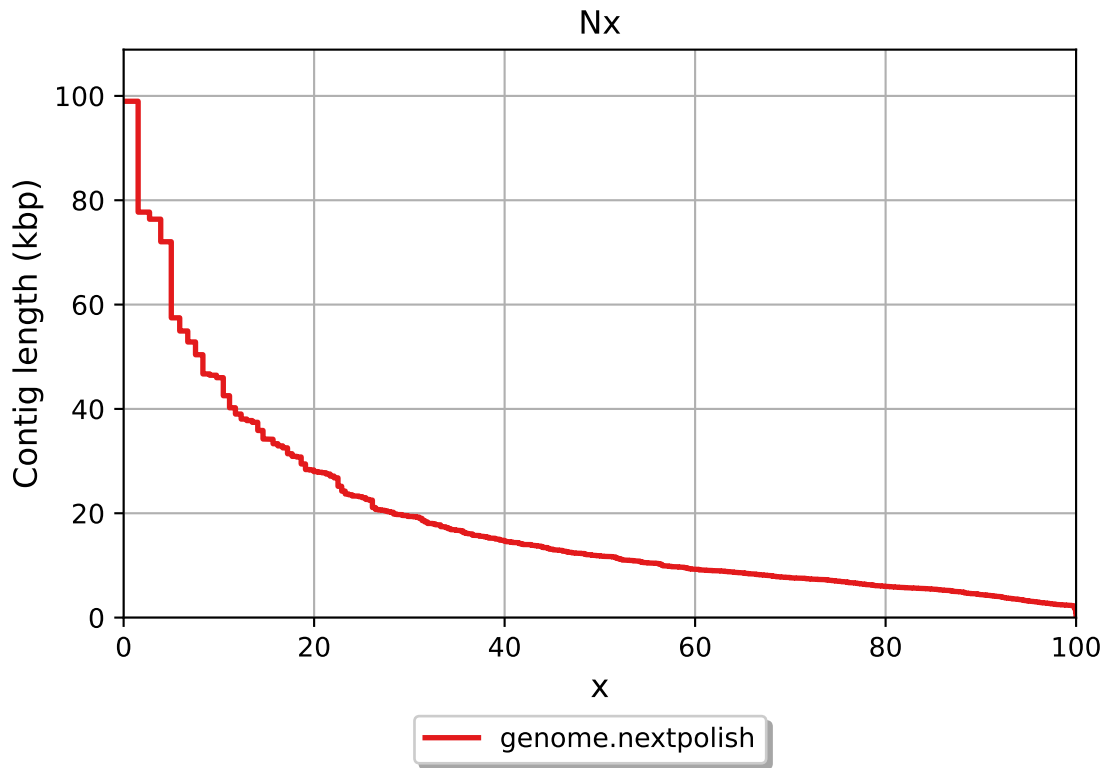


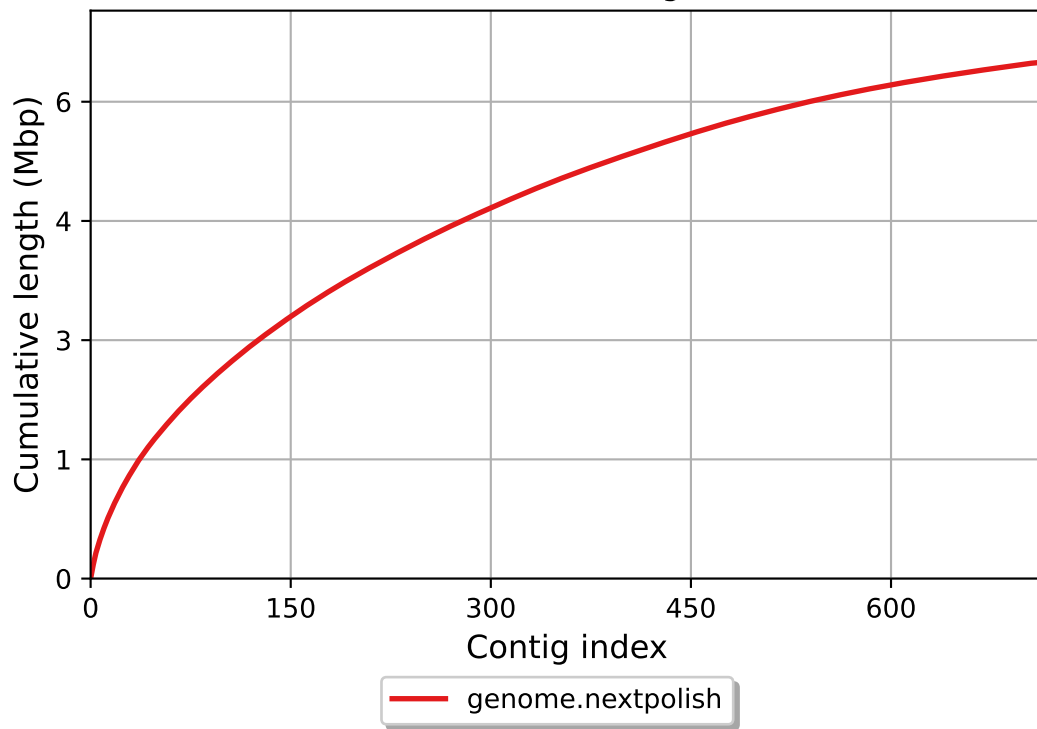
Report

	genome.nextpolish
# contigs (>= 0 bp)	714
# contigs (>= 1000 bp)	712
# contigs (>= 5000 bp)	465
# contigs (>= 10000 bp)	185
# contigs (>= 25000 bp)	36
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	6496530
Total length (>= 1000 bp)	6495008
Total length (>= 5000 bp)	5674713
Total length (>= 10000 bp)	3677899
Total length (>= 25000 bp)	1487378
Total length (>= 50000 bp)	540728
# contigs	714
Largest contig	98972
Total length	6496530
GC (%)	42.31
N50	11862
N90	4401
auN	19011.7
L50	146
L90	502
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 3 part

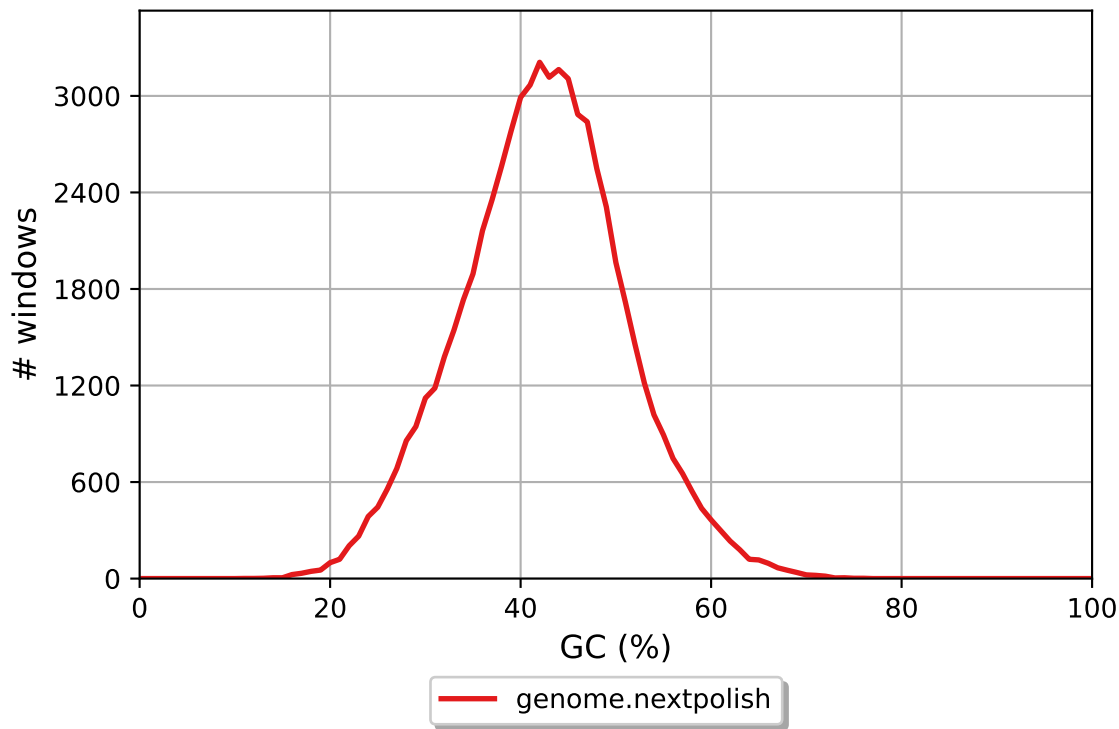
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

