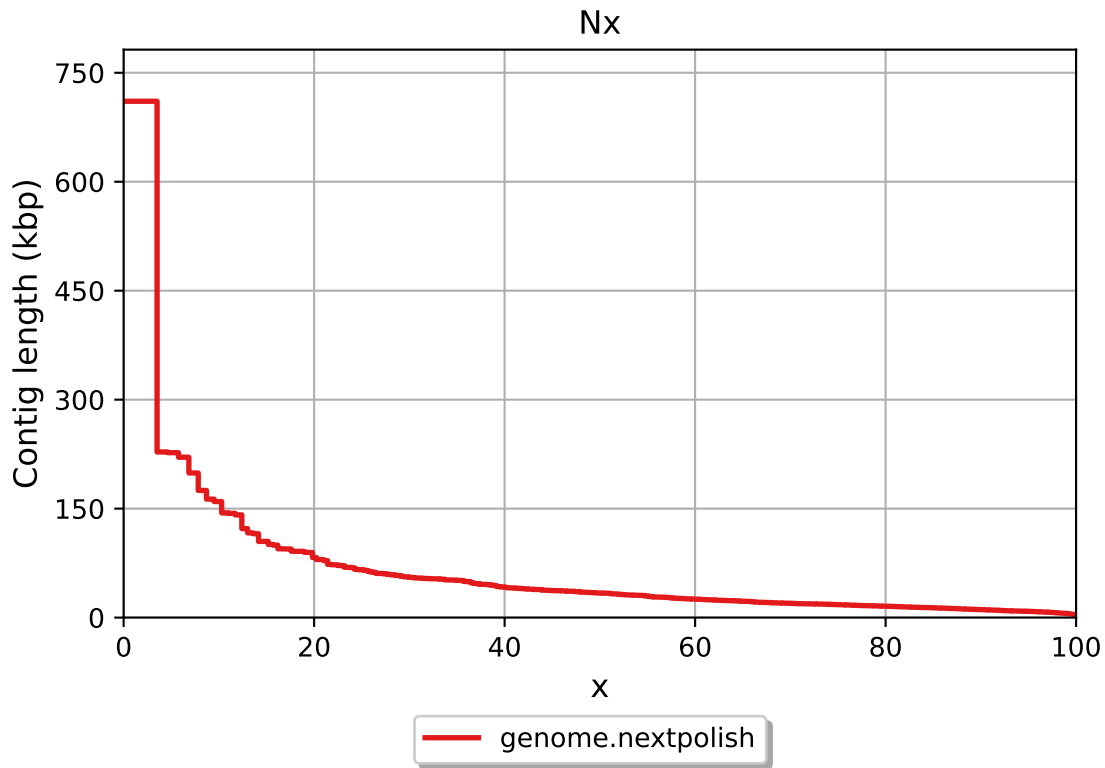


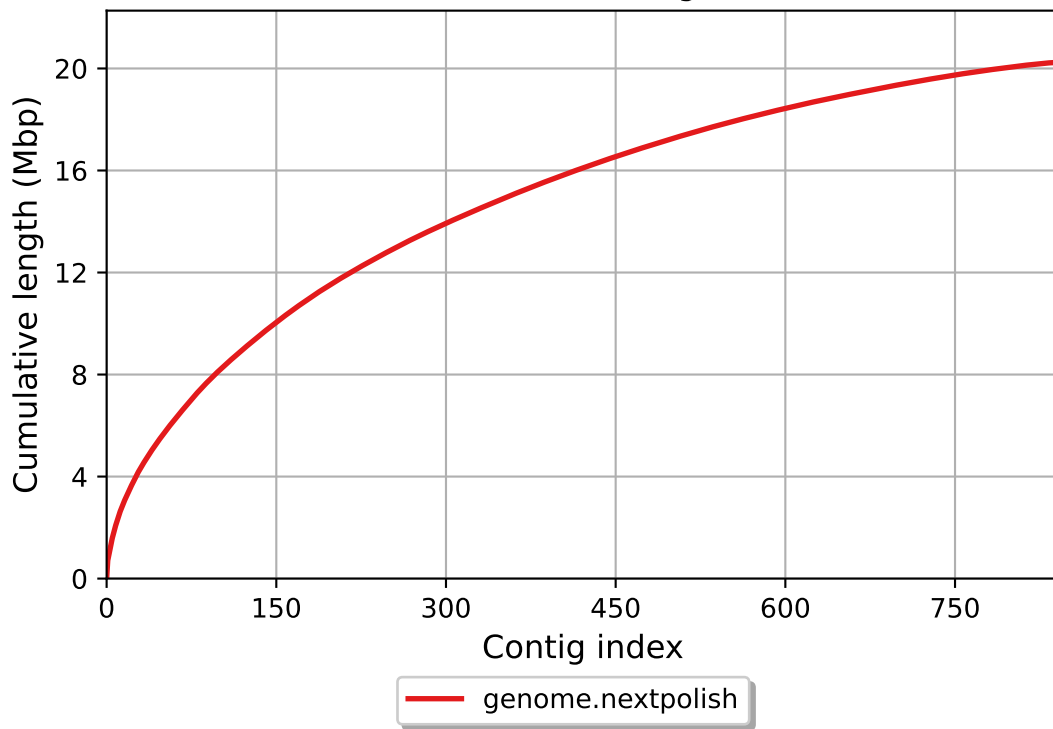
# Report

	genome.nextpolish
# contigs (>= 0 bp)	842
# contigs (>= 1000 bp)	841
# contigs (>= 5000 bp)	814
# contigs (>= 10000 bp)	617
# contigs (>= 25000 bp)	226
# contigs (>= 50000 bp)	79
Total length (>= 0 bp)	20243234
Total length (>= 1000 bp)	20242712
Total length (>= 5000 bp)	20134408
Total length (>= 10000 bp)	18609549
Total length (>= 25000 bp)	12265143
Total length (>= 50000 bp)	7228247
# contigs	842
Largest contig	710764
Total length	20243234
GC (%)	53.24
N50	33743
N90	11006
auN	73726.7
L50	153
L90	580
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	55 + 5 part

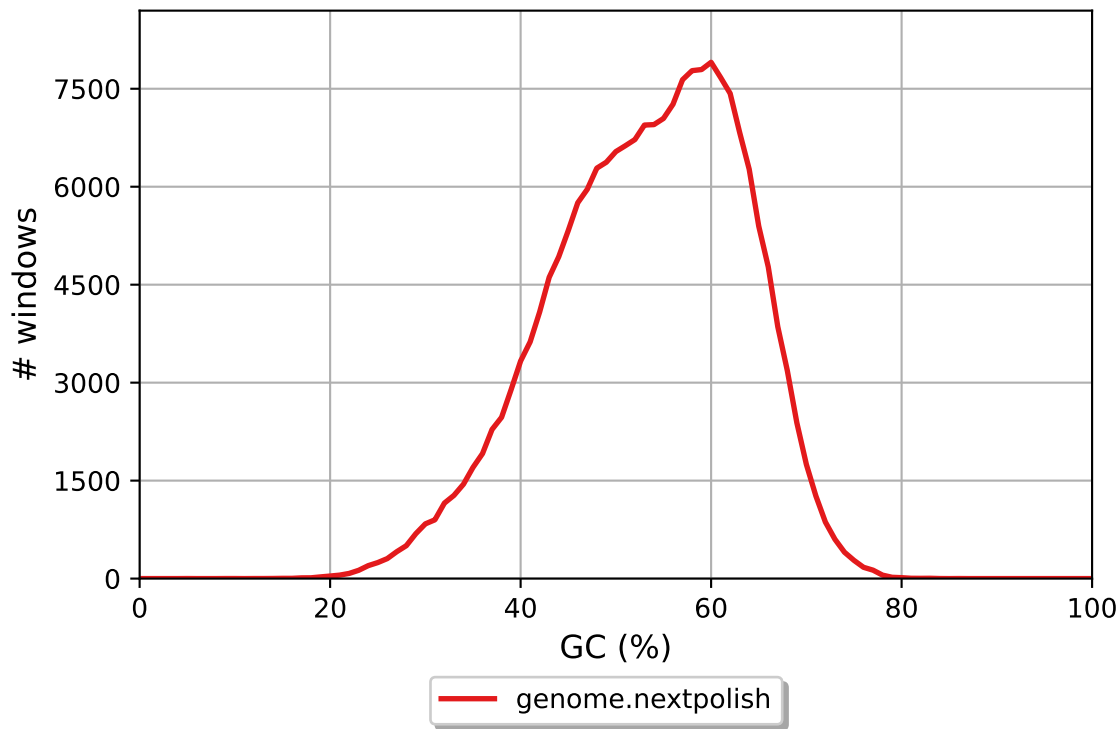
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

