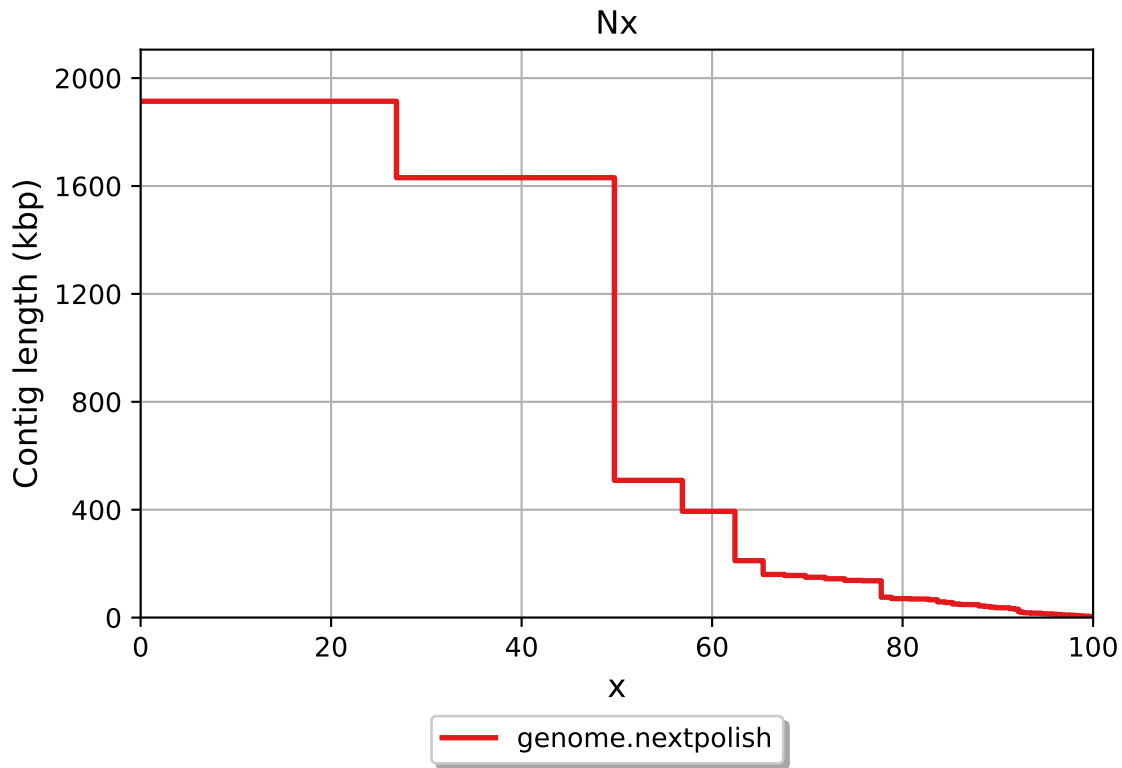


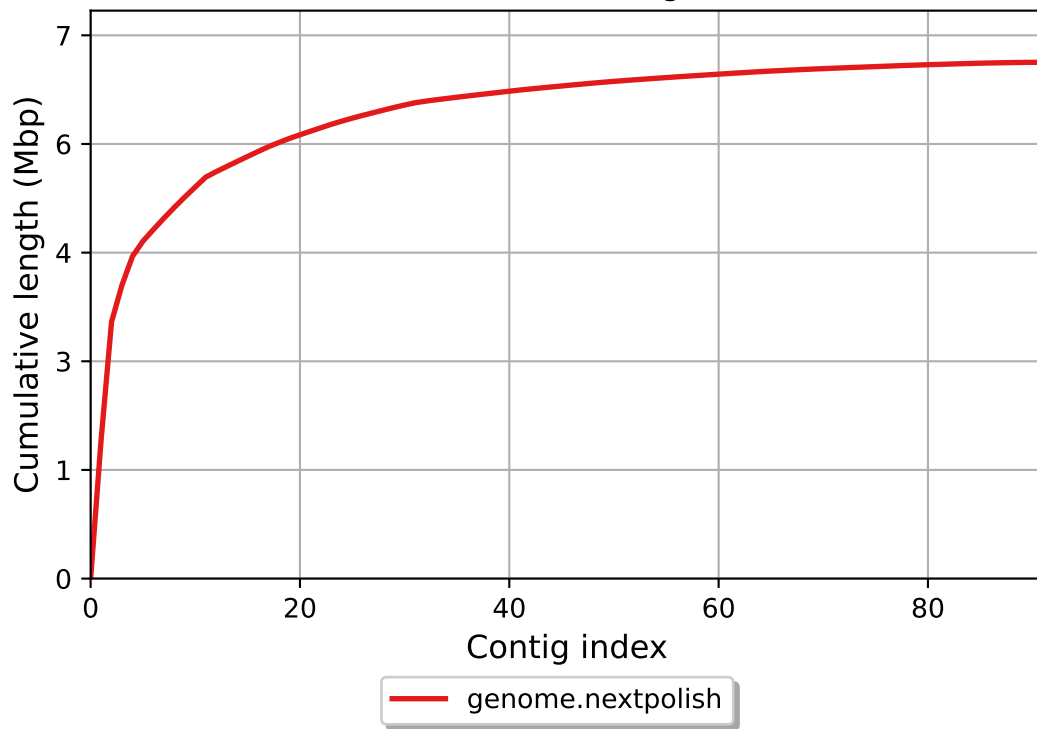
# Report

	genome.nextpolish
# contigs (>= 0 bp)	91
# contigs (>= 1000 bp)	91
# contigs (>= 5000 bp)	78
# contigs (>= 10000 bp)	55
# contigs (>= 25000 bp)	31
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	7127952
Total length (>= 1000 bp)	7127952
Total length (>= 5000 bp)	7084694
Total length (>= 10000 bp)	6916662
Total length (>= 25000 bp)	6568864
Total length (>= 50000 bp)	6126975
# contigs	91
Largest contig	1914198
Total length	7127952
GC (%)	39.12
N50	508823
N90	36815
auN	978747.6
L50	3
L90	27
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	44 + 3 part

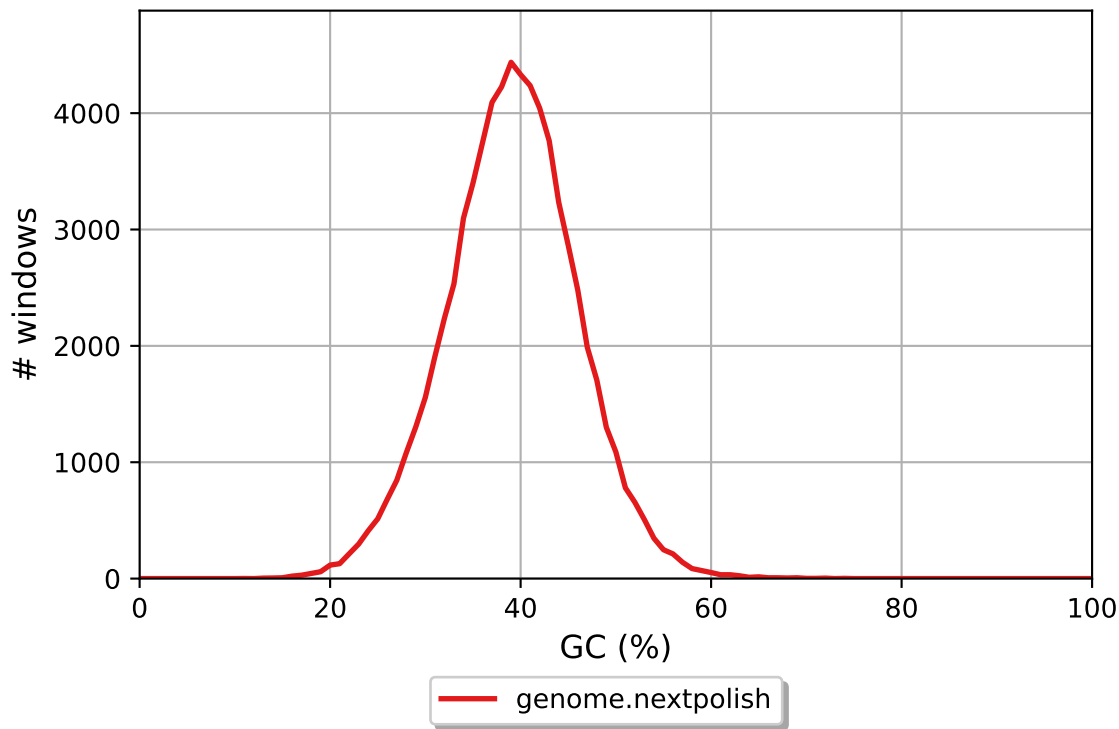
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

