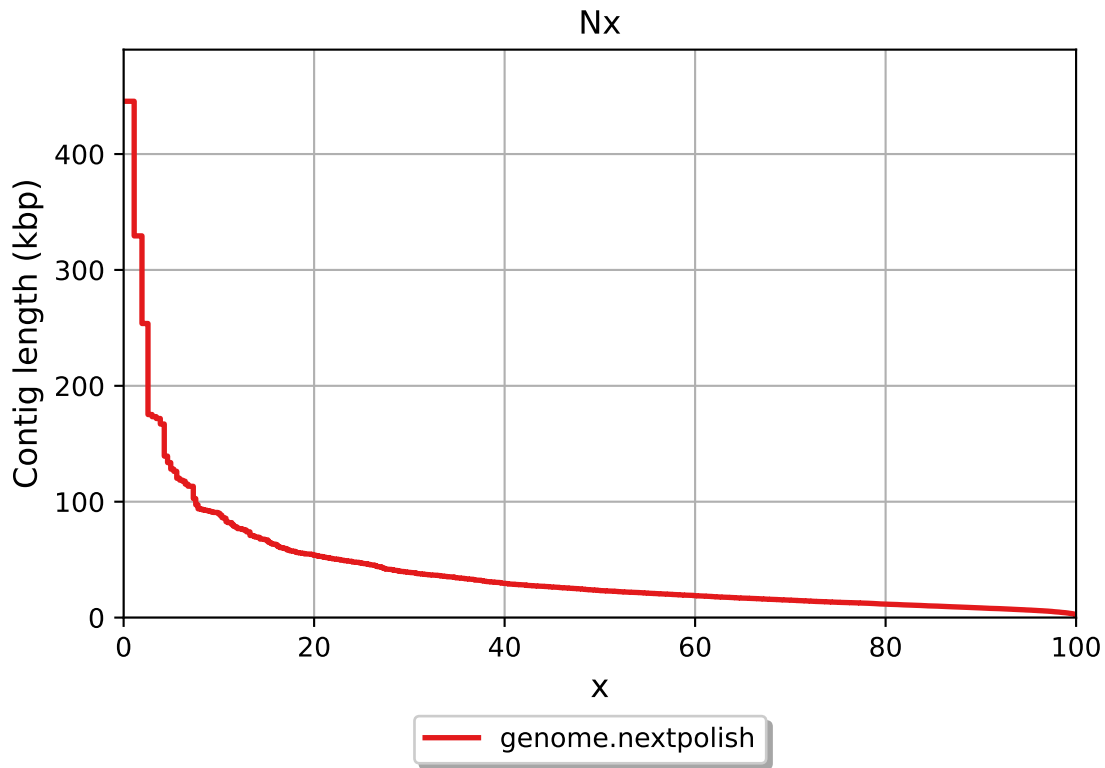


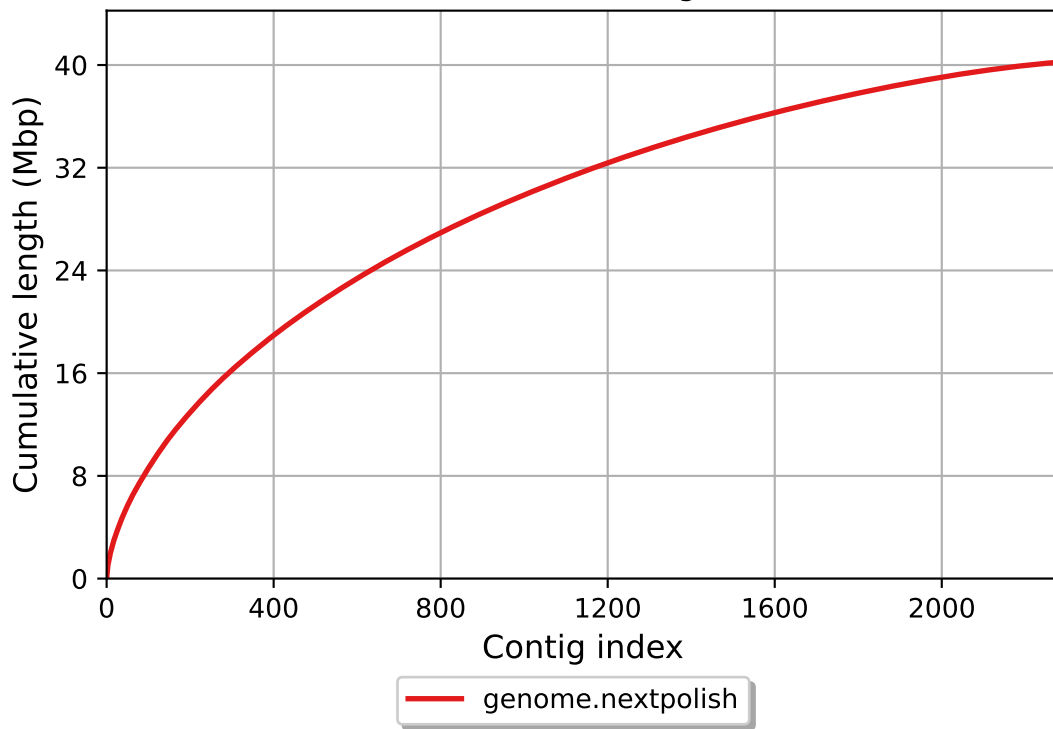
# Report

	genome.nextpolish
# contigs (>= 0 bp)	2281
# contigs (>= 1000 bp)	2279
# contigs (>= 5000 bp)	2054
# contigs (>= 10000 bp)	1369
# contigs (>= 25000 bp)	406
# contigs (>= 50000 bp)	111
Total length (>= 0 bp)	40214400
Total length (>= 1000 bp)	40212517
Total length (>= 5000 bp)	39336932
Total length (>= 10000 bp)	34189765
Total length (>= 25000 bp)	19101542
Total length (>= 50000 bp)	9158247
# contigs	2281
Largest contig	445545
Total length	40214400
GC (%)	46.12
N50	23348
N90	8341
auN	42669.3
L50	448
L90	1588
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	126 + 7 part

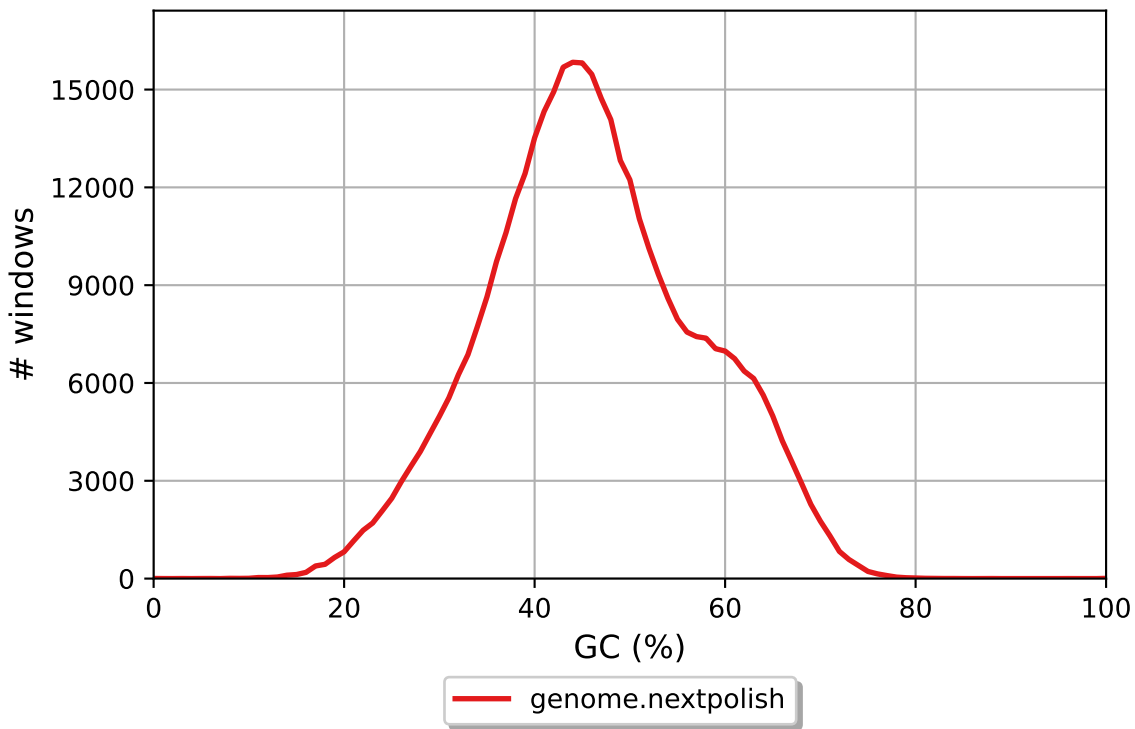
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

