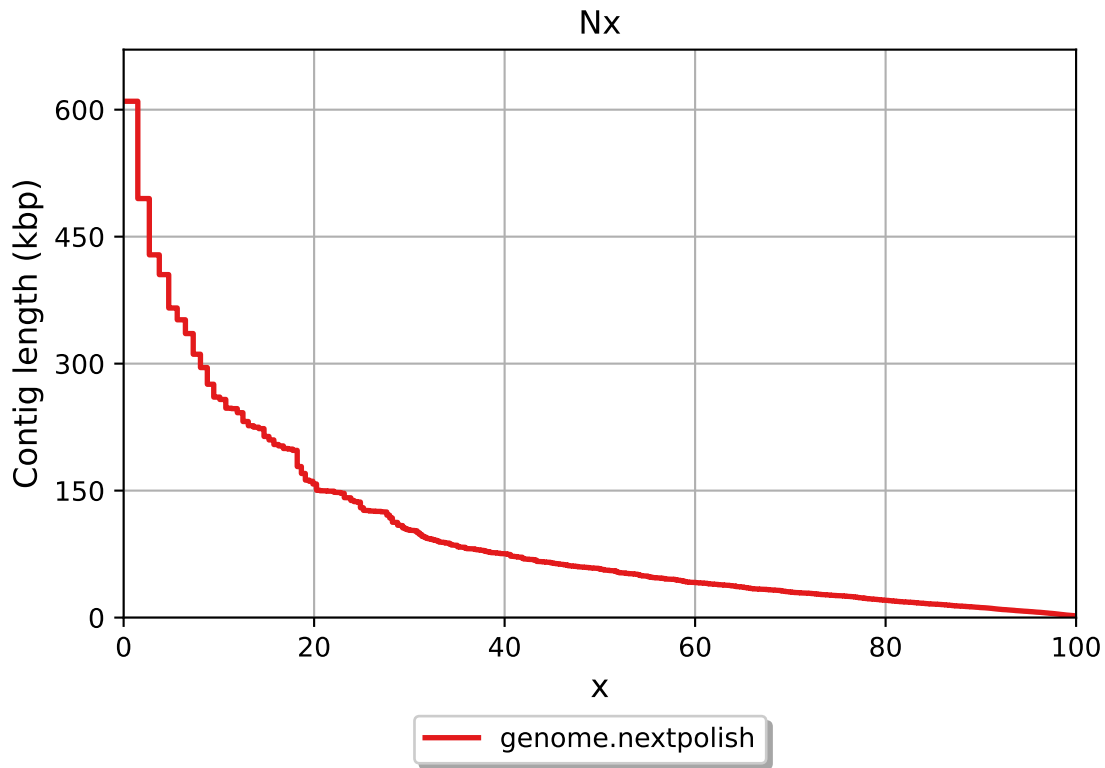


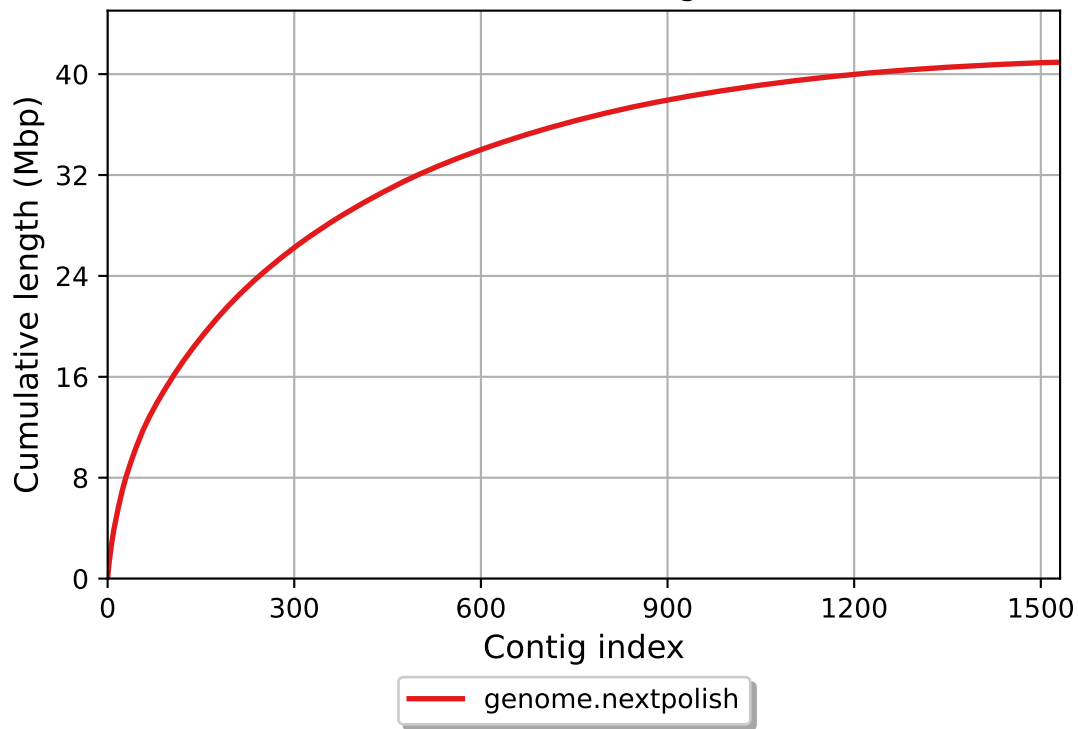
Report

	genome.nextpolish
# contigs (>= 0 bp)	1533
# contigs (>= 1000 bp)	1518
# contigs (>= 5000 bp)	1176
# contigs (>= 10000 bp)	863
# contigs (>= 25000 bp)	463
# contigs (>= 50000 bp)	205
Total length (>= 0 bp)	40941032
Total length (>= 1000 bp)	40931605
Total length (>= 5000 bp)	39867261
Total length (>= 10000 bp)	37591728
Total length (>= 25000 bp)	31160697
Total length (>= 50000 bp)	22189964
# contigs	1531
Largest contig	609936
Total length	40940398
GC (%)	44.99
N50	57710
N90	11784
auN	103479.2
L50	173
L90	796
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	88 + 1 part

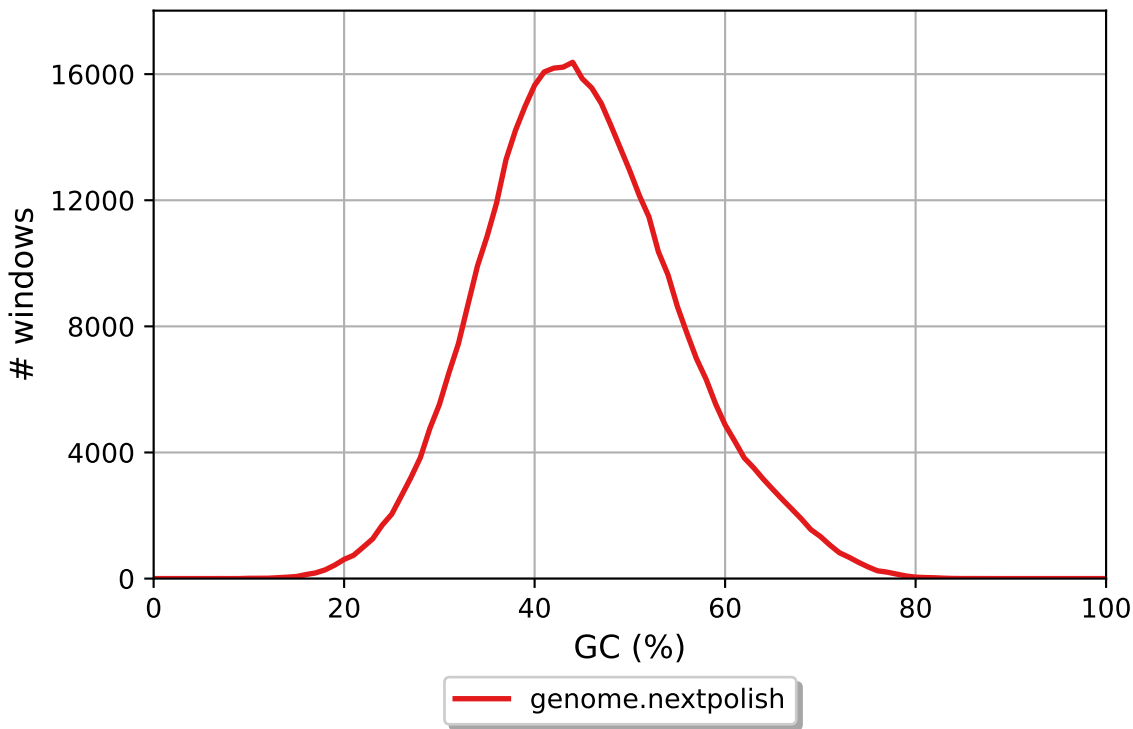
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

