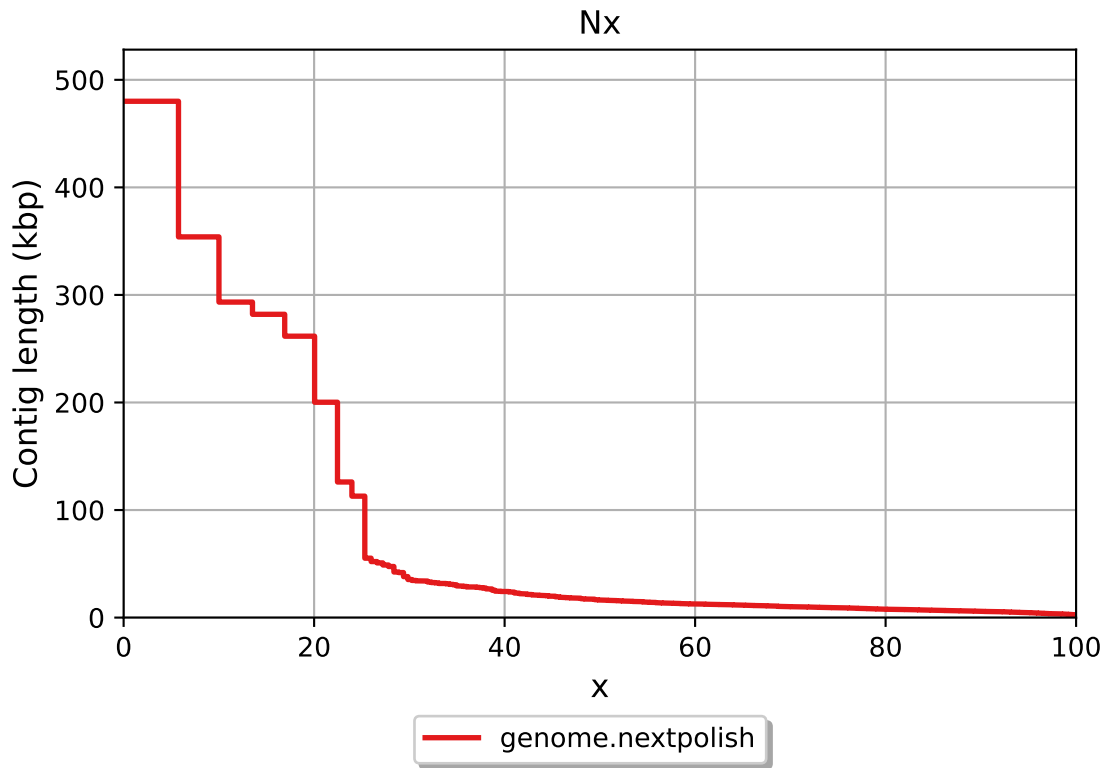


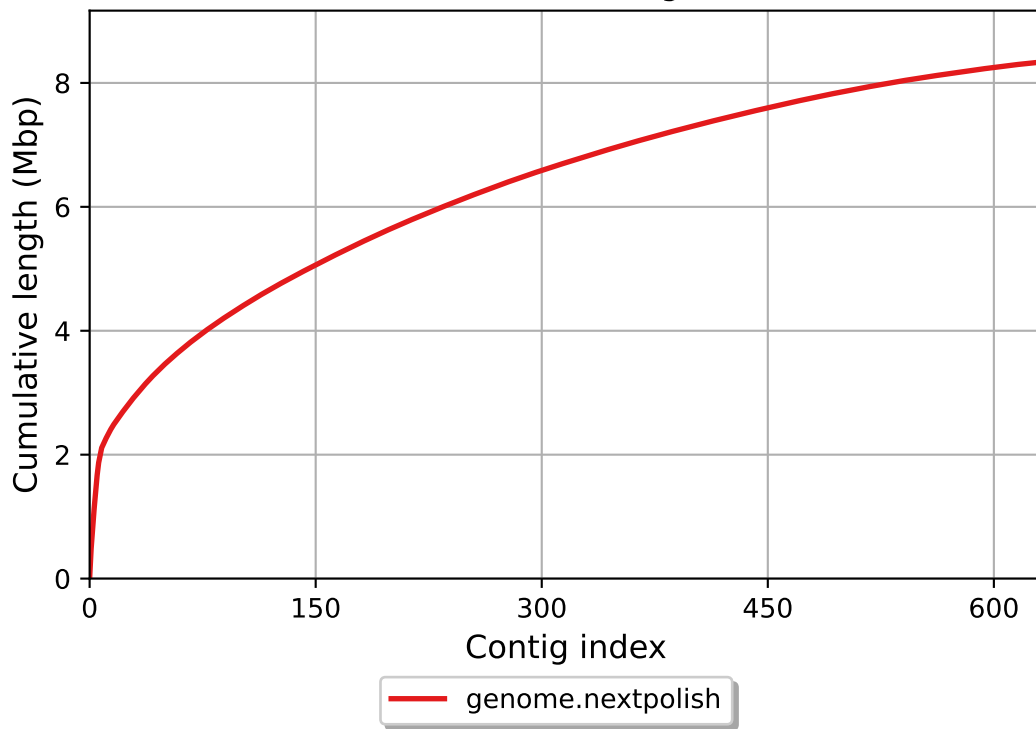
Report

	genome.nextpolish
# contigs (>= 0 bp)	632
# contigs (>= 1000 bp)	629
# contigs (>= 5000 bp)	487
# contigs (>= 10000 bp)	225
# contigs (>= 25000 bp)	41
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	8333430
Total length (>= 1000 bp)	8331031
Total length (>= 5000 bp)	7796038
Total length (>= 10000 bp)	5907713
Total length (>= 25000 bp)	3248599
Total length (>= 50000 bp)	2268355
# contigs	632
Largest contig	480061
Total length	8333430
GC (%)	43.57
N50	16376
N90	5806
auN	90857.5
L50	87
L90	433
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	27 + 5 part

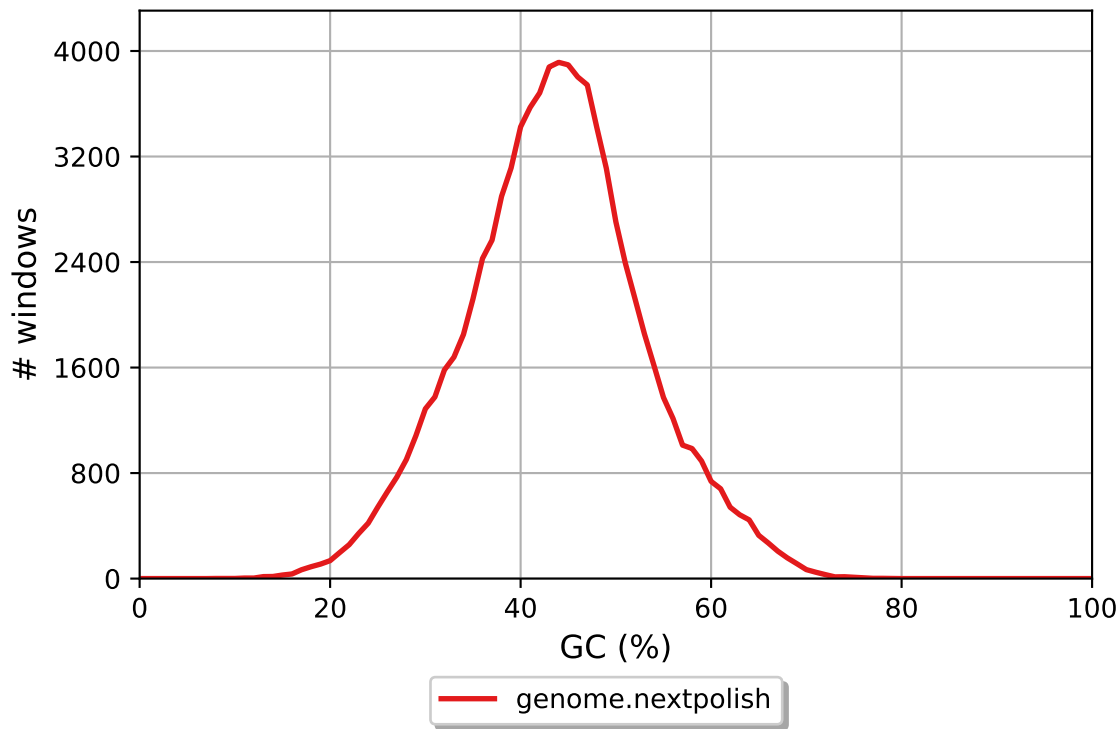
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

