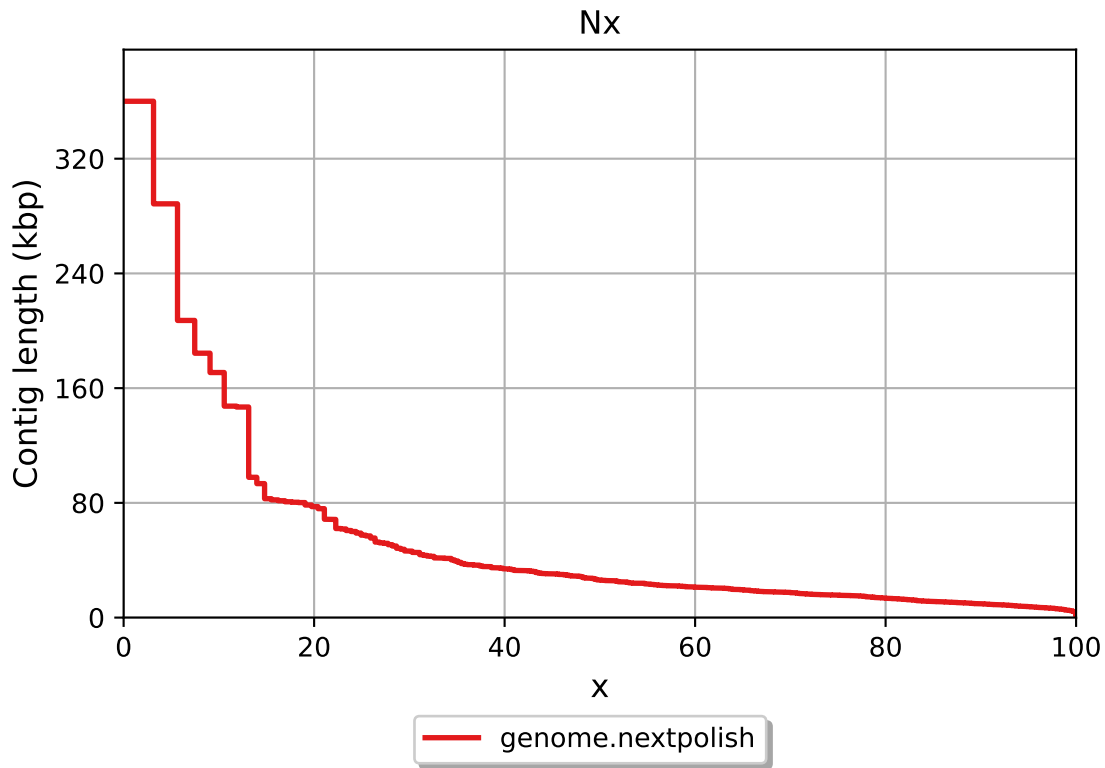


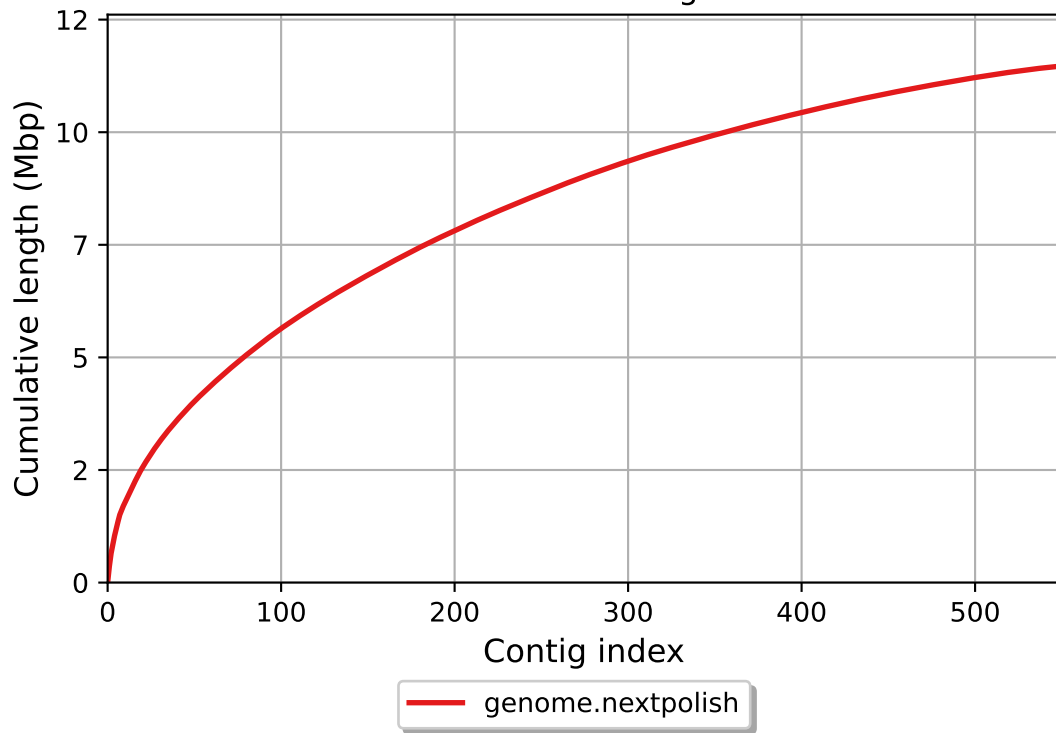
Report

	genome.nextpolish
# contigs (>= 0 bp)	549
# contigs (>= 1000 bp)	549
# contigs (>= 5000 bp)	531
# contigs (>= 10000 bp)	375
# contigs (>= 25000 bp)	112
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	11464058
Total length (>= 1000 bp)	11464058
Total length (>= 5000 bp)	11388917
Total length (>= 10000 bp)	10198328
Total length (>= 25000 bp)	5953155
Total length (>= 50000 bp)	3234107
# contigs	549
Largest contig	360044
Total length	11464058
GC (%)	54.29
N50	26181
N90	9610
auN	58004.9
L50	104
L90	388
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	43 + 3 part

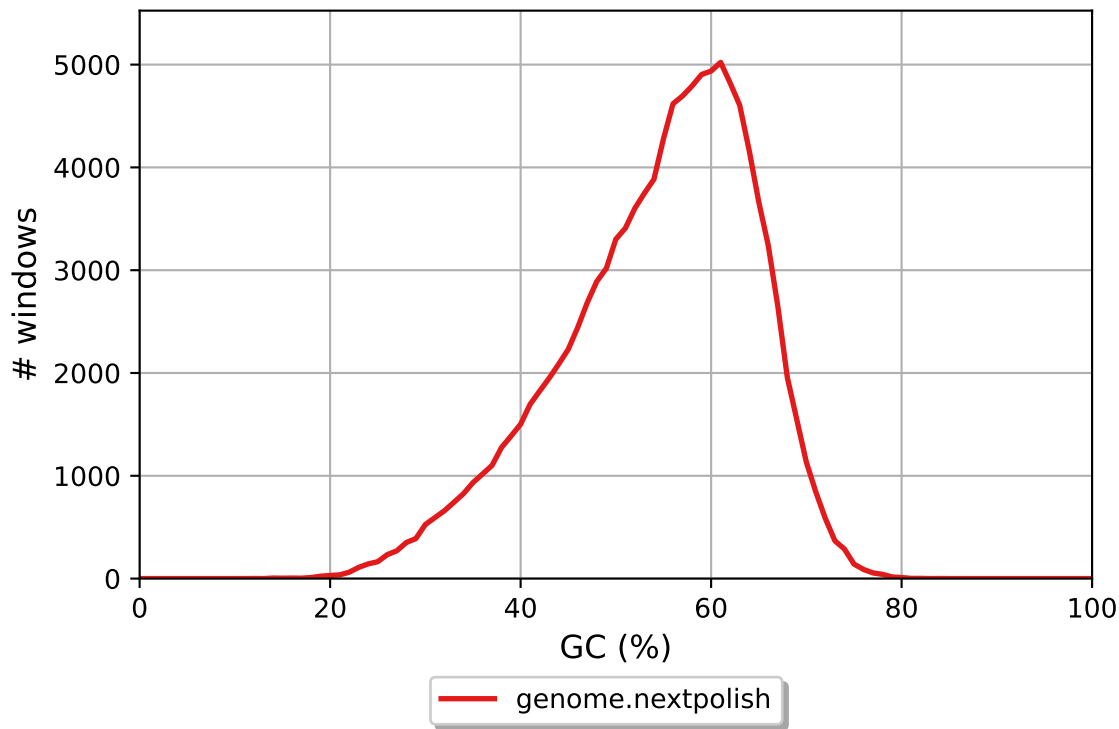
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

