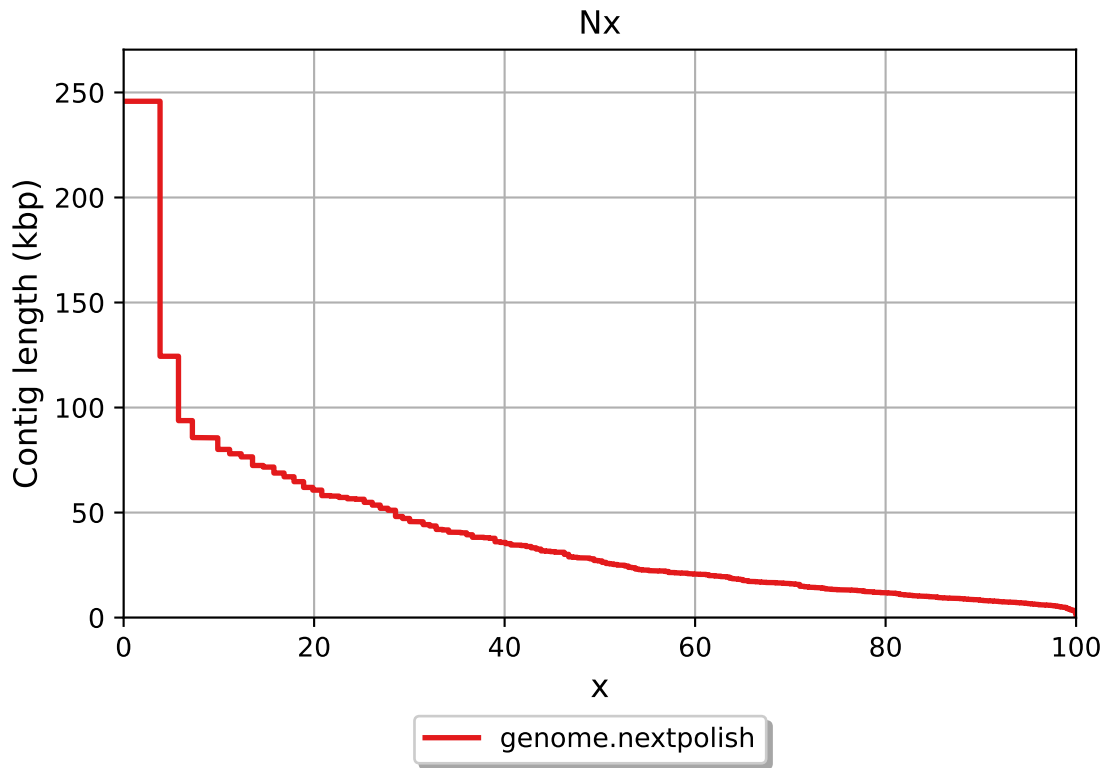


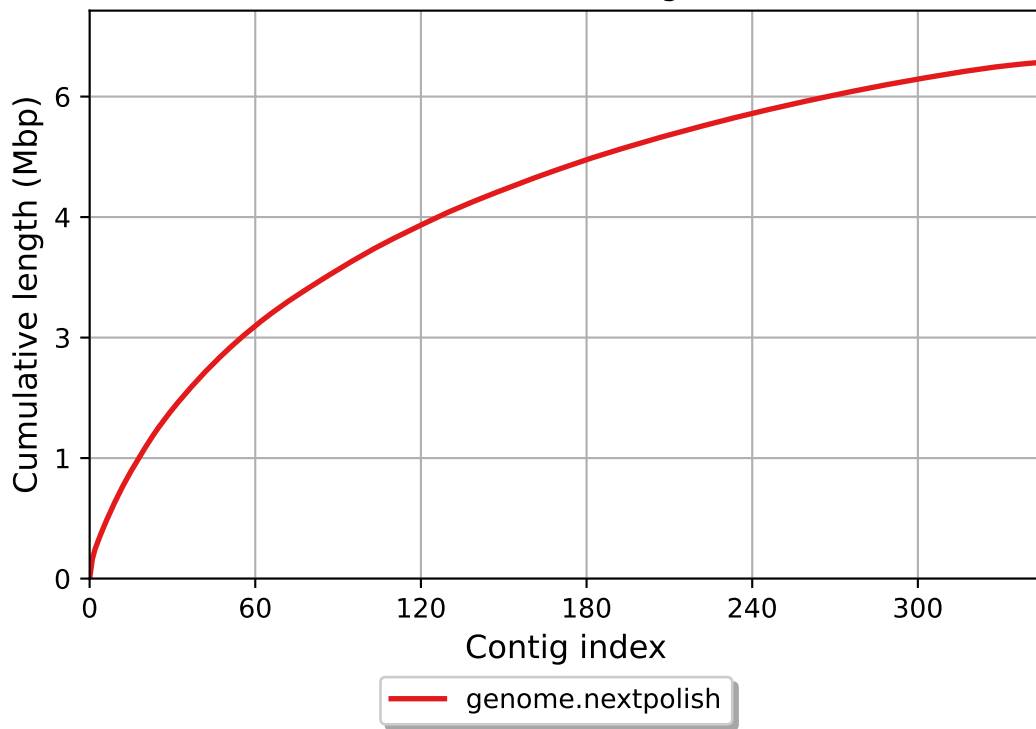
Report

	genome.nextpolish
# contigs (>= 0 bp)	345
# contigs (>= 1000 bp)	345
# contigs (>= 5000 bp)	320
# contigs (>= 10000 bp)	201
# contigs (>= 25000 bp)	67
# contigs (>= 50000 bp)	24
Total length (>= 0 bp)	6427065
Total length (>= 1000 bp)	6427065
Total length (>= 5000 bp)	6329769
Total length (>= 10000 bp)	5435943
Total length (>= 25000 bp)	3331009
Total length (>= 50000 bp)	1834659
# contigs	345
Largest contig	245808
Total length	6427065
GC (%)	49.35
N50	26990
N90	8261
auN	42500.3
L50	63
L90	240
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 1 part

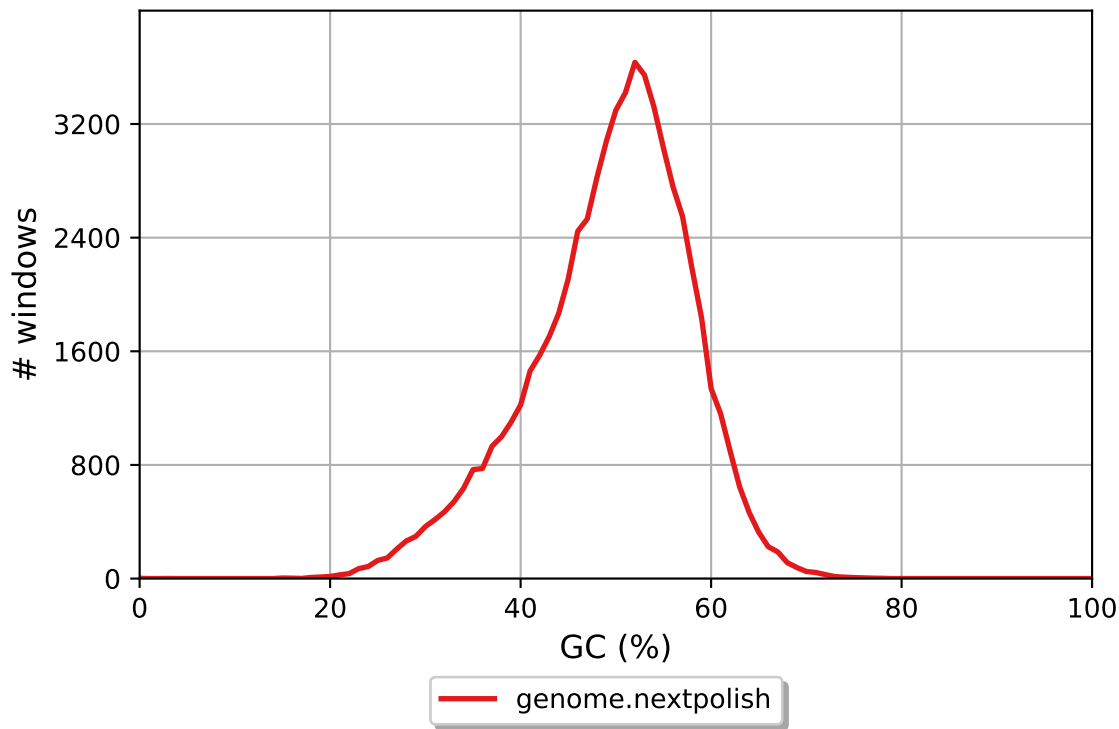
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

