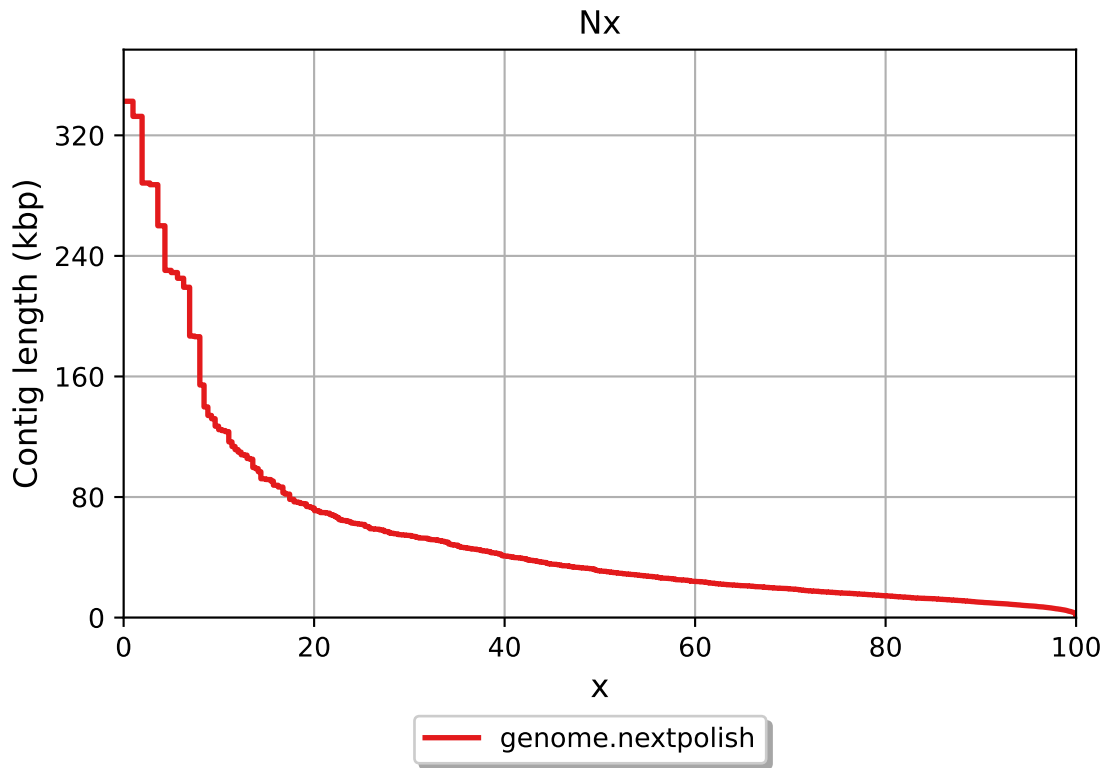


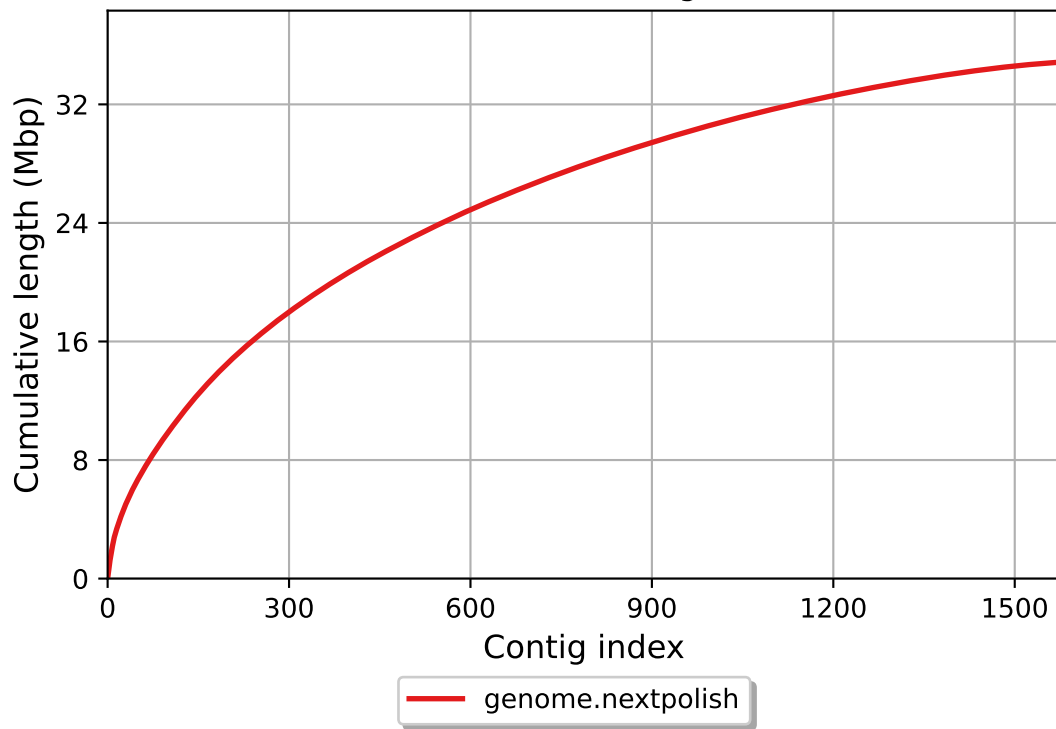
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1575
# contigs (>= 1000 bp)	1575
# contigs (>= 5000 bp)	1465
# contigs (>= 10000 bp)	1079
# contigs (>= 25000 bp)	390
# contigs (>= 50000 bp)	137
Total length (>= 0 bp)	34841472
Total length (>= 1000 bp)	34841472
Total length (>= 5000 bp)	34428442
Total length (>= 10000 bp)	31468203
Total length (>= 25000 bp)	20447820
Total length (>= 50000 bp)	11833655
# contigs	1575
Largest contig	342604
Total length	34841472
GC (%)	46.05
N50	30957
N90	10175
auN	56461.1
L50	282
L90	1069
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	93 + 8 part

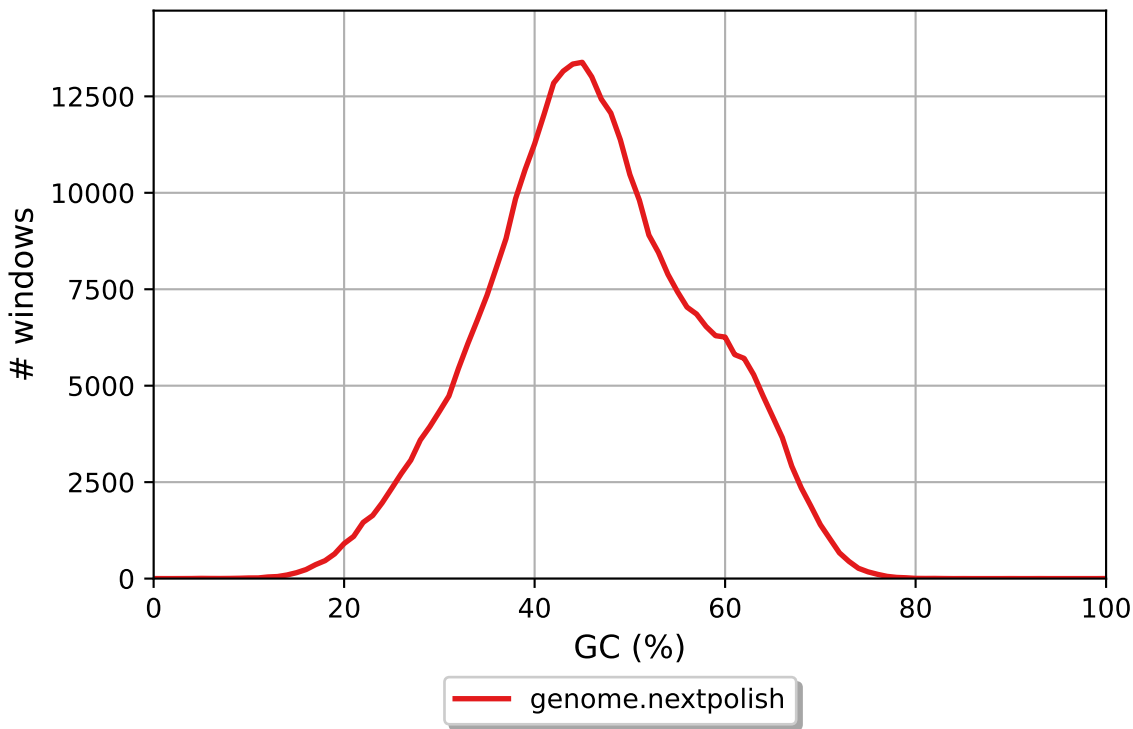
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

