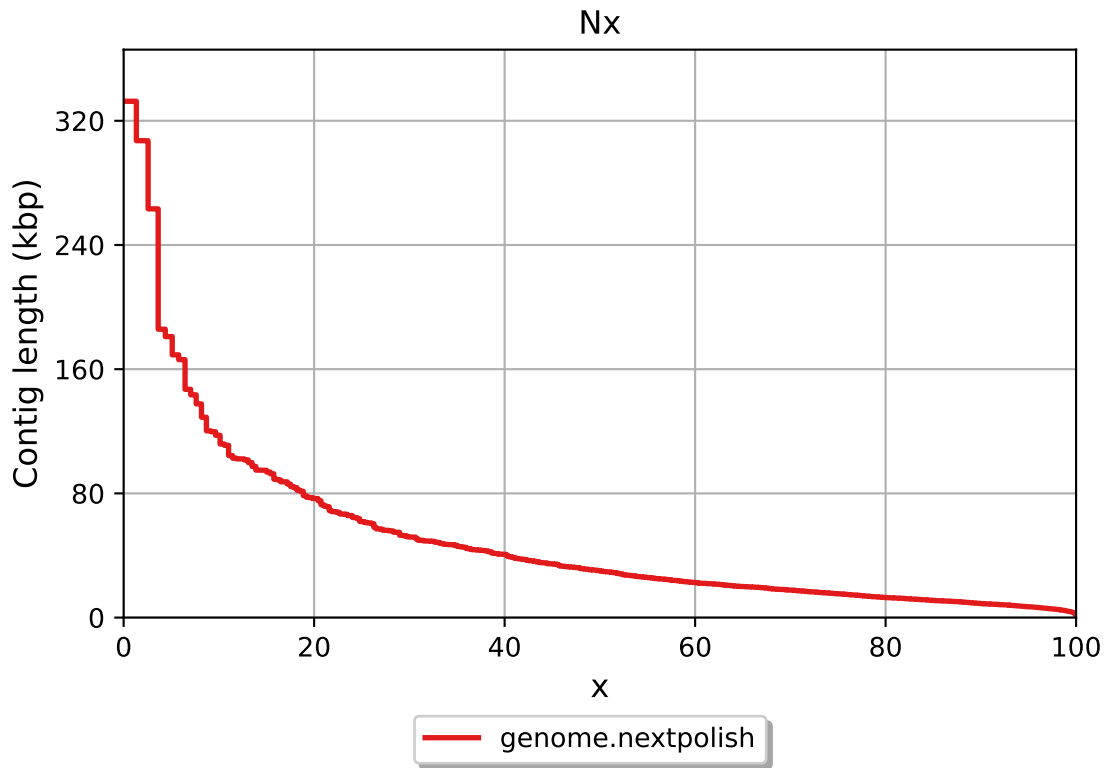


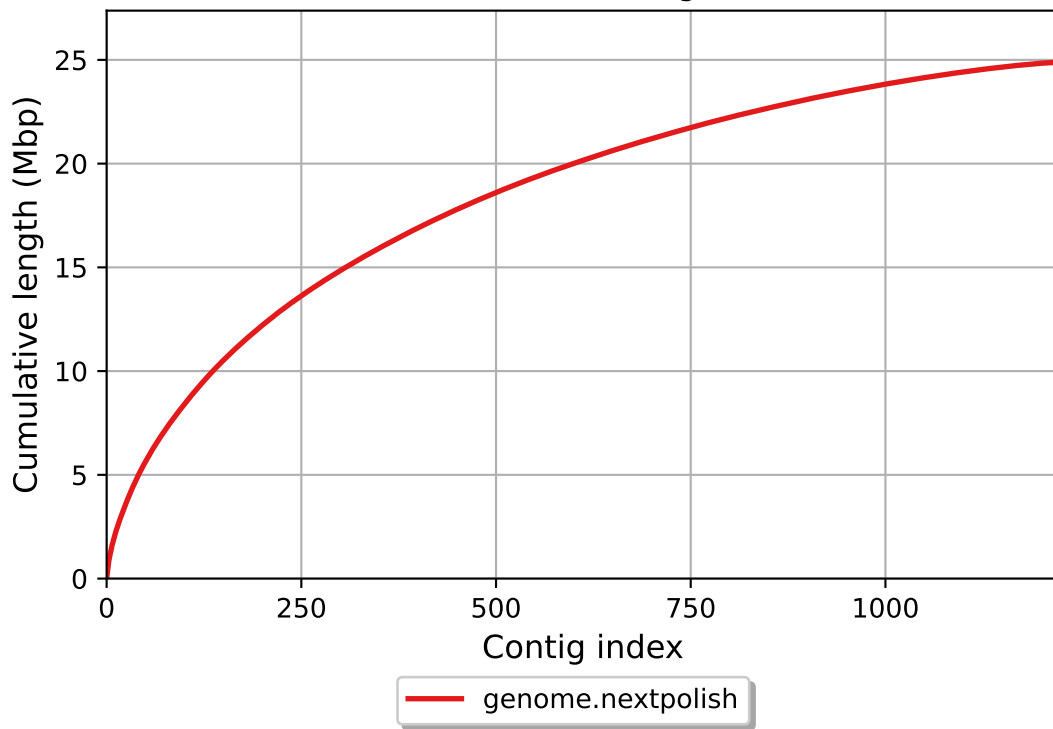
Report

	genome.nextpolish
# contigs (>= 0 bp)	1223
# contigs (>= 1000 bp)	1218
# contigs (>= 5000 bp)	1109
# contigs (>= 10000 bp)	767
# contigs (>= 25000 bp)	264
# contigs (>= 50000 bp)	85
Total length (>= 0 bp)	24884733
Total length (>= 1000 bp)	24881280
Total length (>= 5000 bp)	24465414
Total length (>= 10000 bp)	21911333
Total length (>= 25000 bp)	13985516
Total length (>= 50000 bp)	7680253
# contigs	1223
Largest contig	332568
Total length	24884733
GC (%)	53.16
N50	30181
N90	9066
auN	52299.5
L50	208
L90	818
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	49 + 8 part

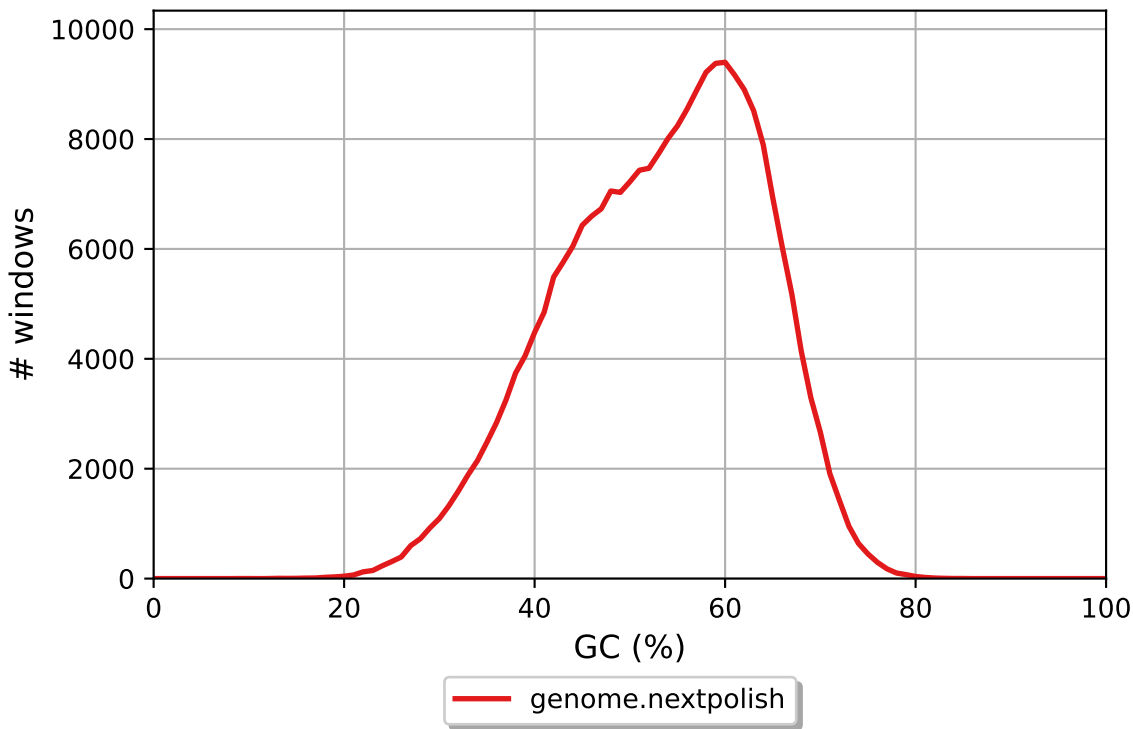
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

