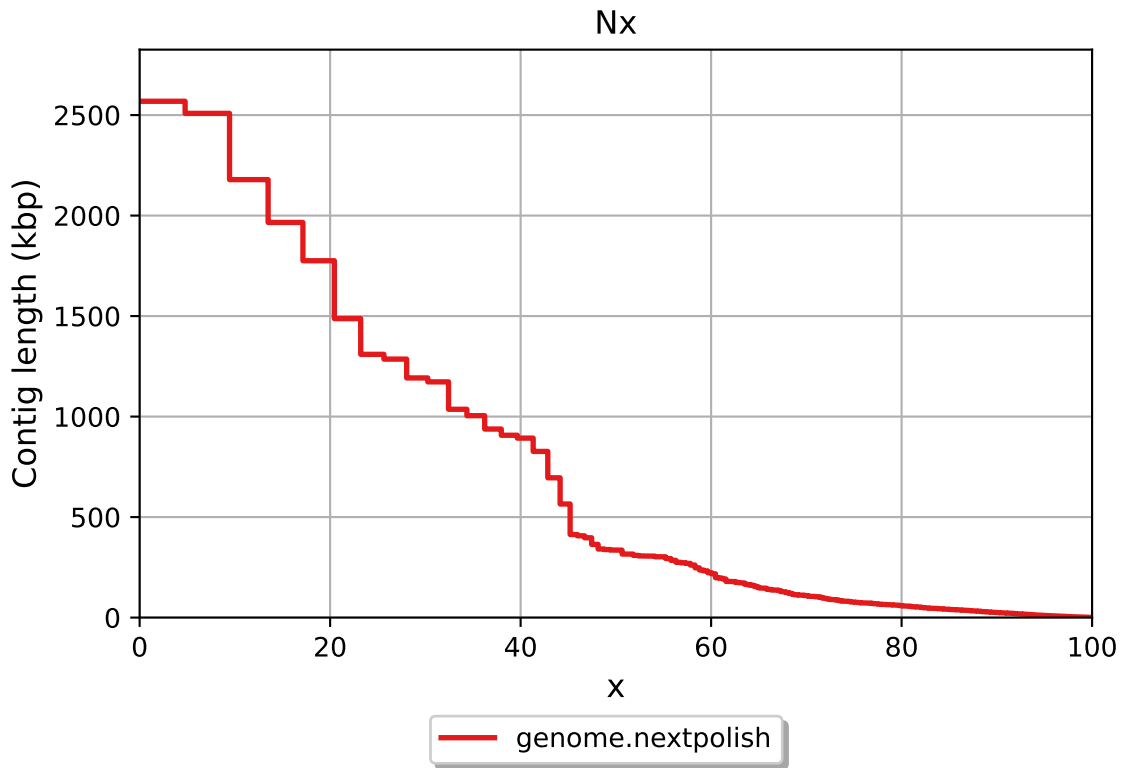


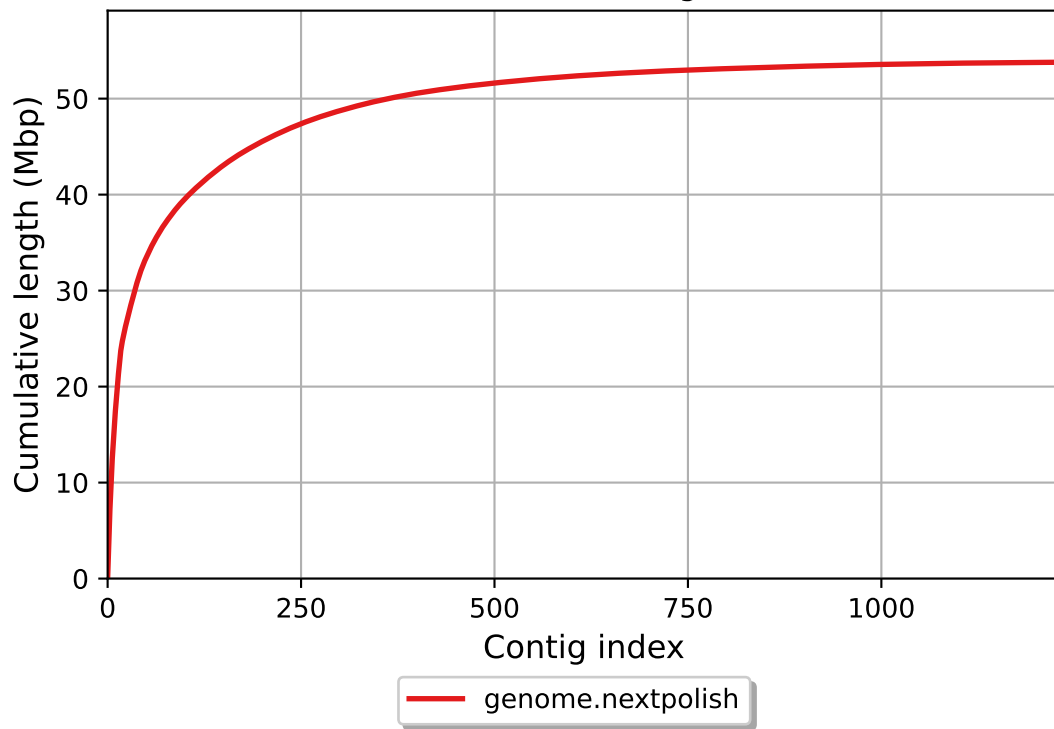
Report

	genome.nextpolish
# contigs (>= 0 bp)	1245
# contigs (>= 1000 bp)	1111
# contigs (>= 5000 bp)	630
# contigs (>= 10000 bp)	457
# contigs (>= 25000 bp)	286
# contigs (>= 50000 bp)	170
Total length (>= 0 bp)	53792956
Total length (>= 1000 bp)	53704761
Total length (>= 5000 bp)	52497066
Total length (>= 10000 bp)	51230486
Total length (>= 25000 bp)	48404172
Total length (>= 50000 bp)	44177124
# contigs	1231
Largest contig	2568676
Total length	53787934
GC (%)	43.24
N50	336119
N90	24973
auN	802489.6
L50	25
L90	287
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	128 + 8 part

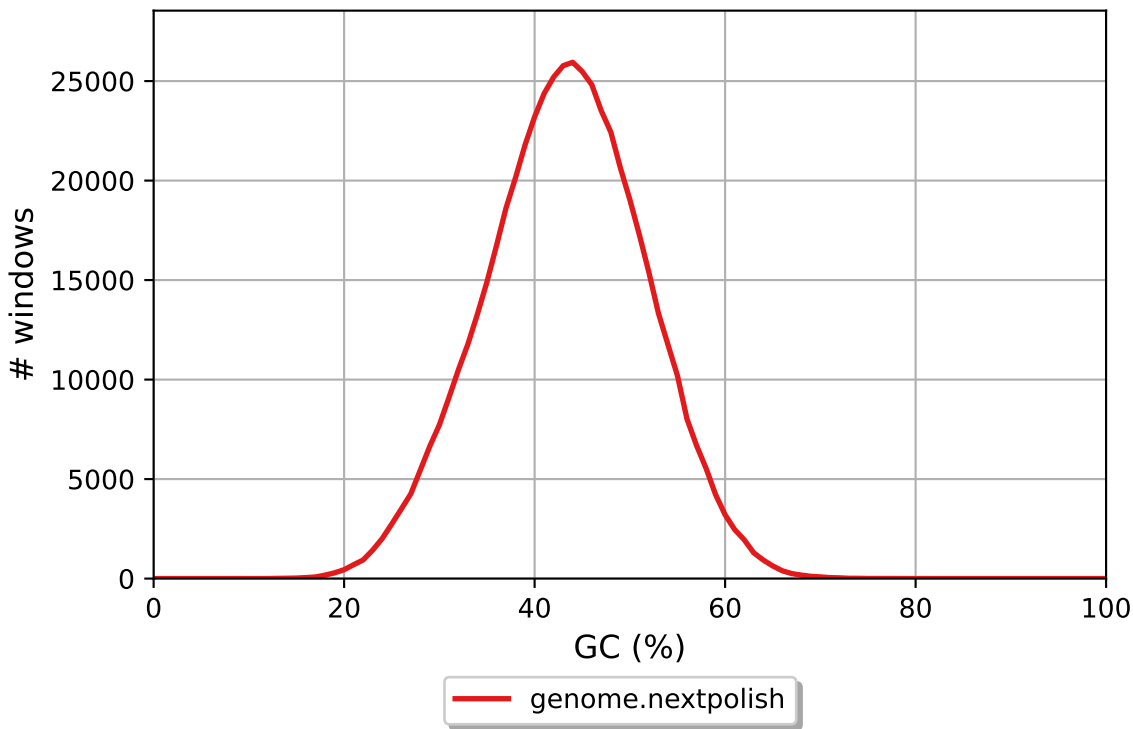
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

