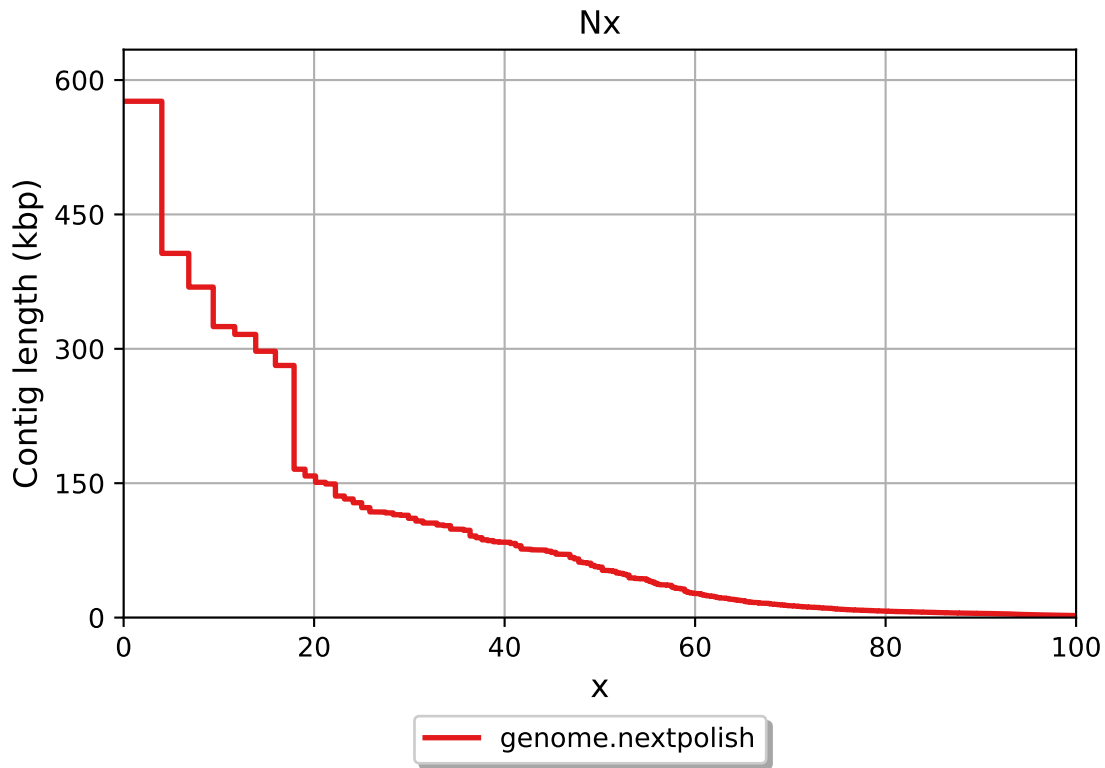


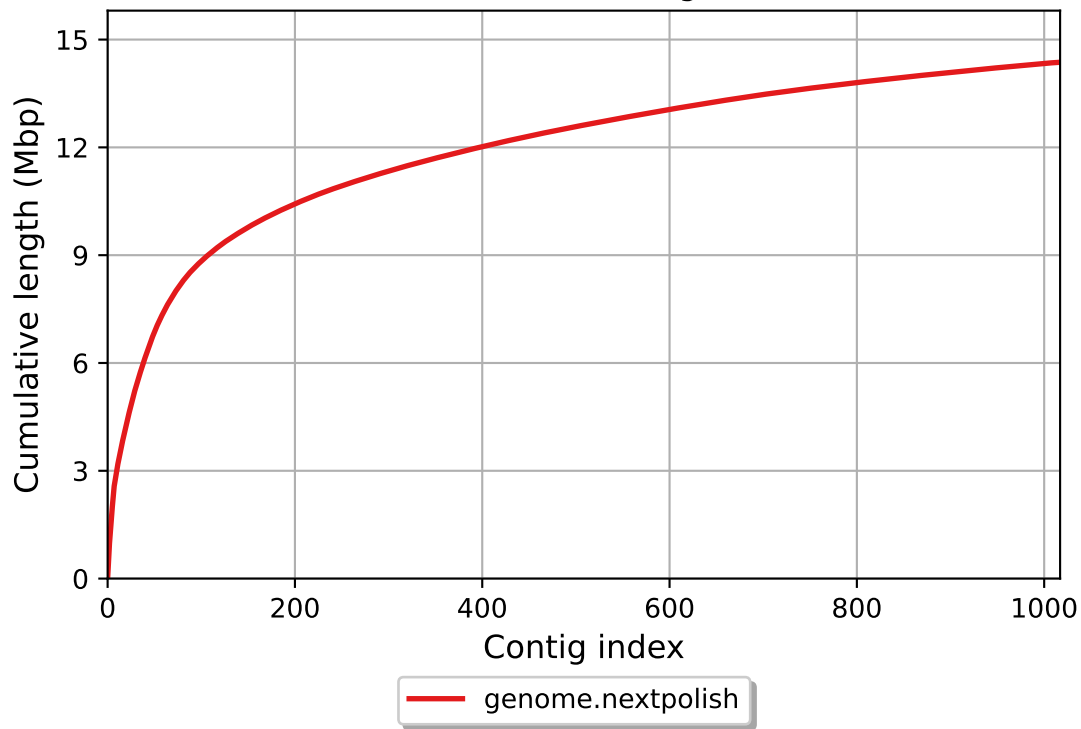
Report

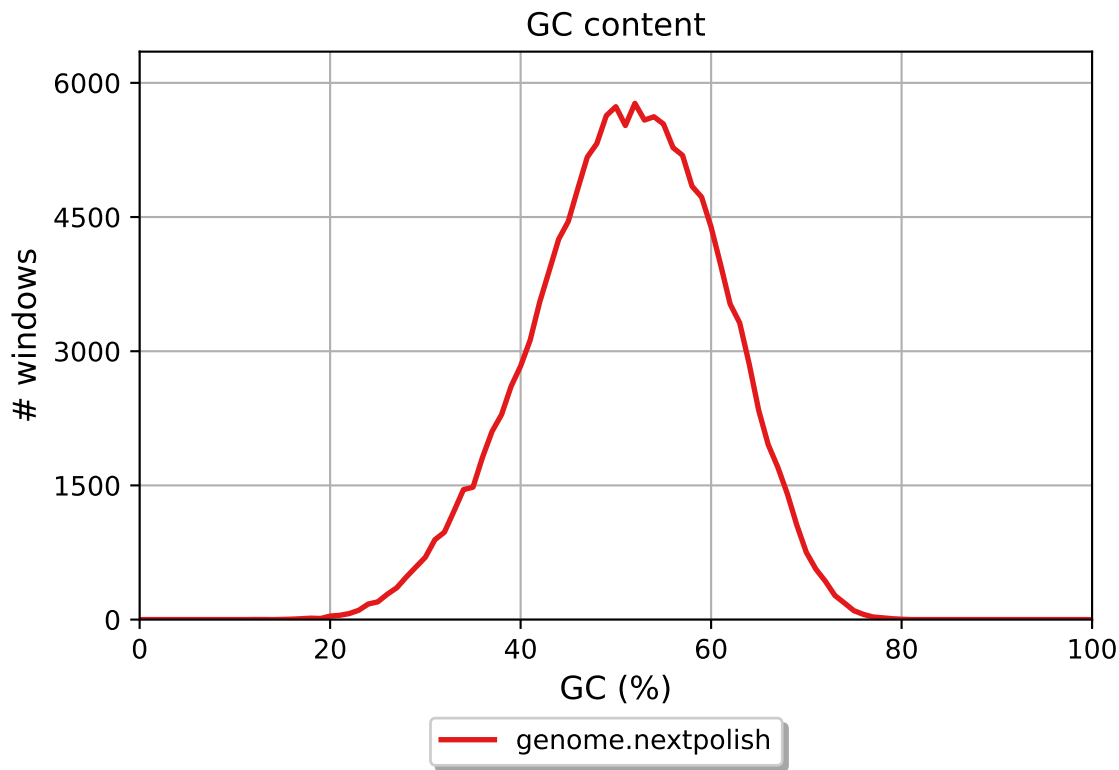
	genome.nextpolish
# contigs (>= 0 bp)	1017
# contigs (>= 1000 bp)	1015
# contigs (>= 5000 bp)	510
# contigs (>= 10000 bp)	226
# contigs (>= 25000 bp)	98
# contigs (>= 50000 bp)	60
Total length (>= 0 bp)	14368382
Total length (>= 1000 bp)	14367203
Total length (>= 5000 bp)	12628828
Total length (>= 10000 bp)	10706201
Total length (>= 25000 bp)	8786222
Total length (>= 50000 bp)	7429797
# contigs	1017
Largest contig	576313
Total length	14368382
GC (%)	51.04
N50	56285
N90	4652
auN	111190.0
L50	56
L90	573
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	46 + 10 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





genome.nextpolish GC content

