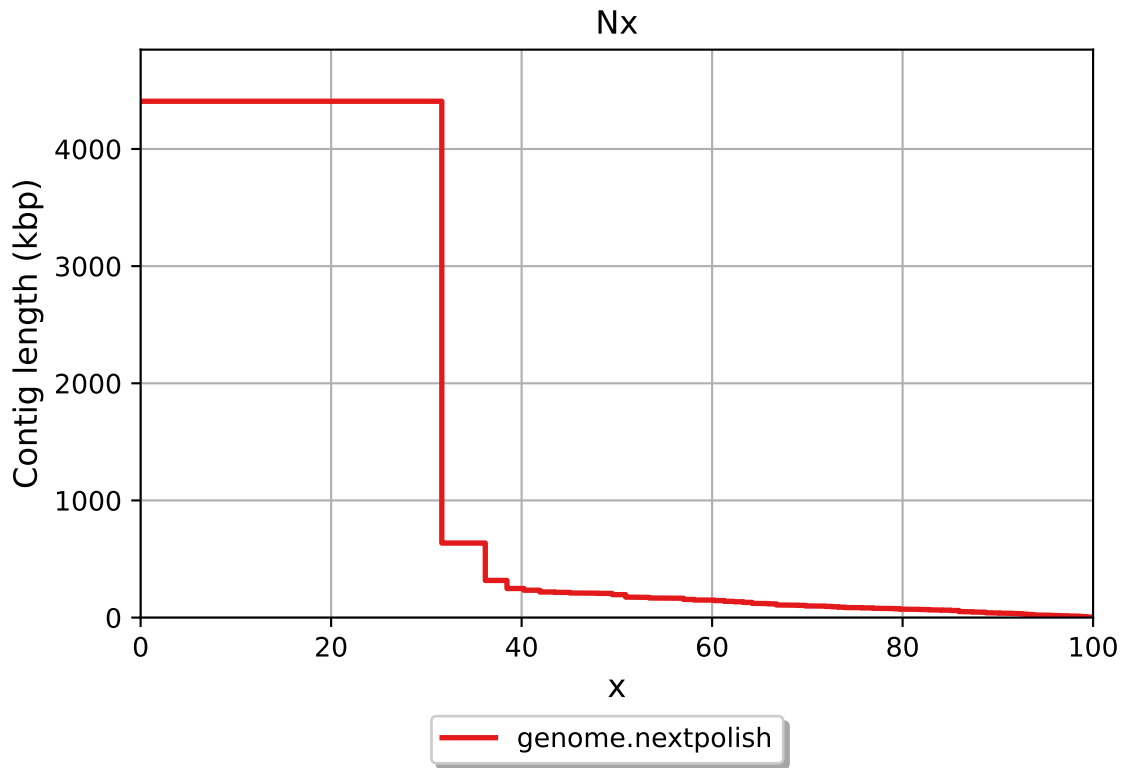


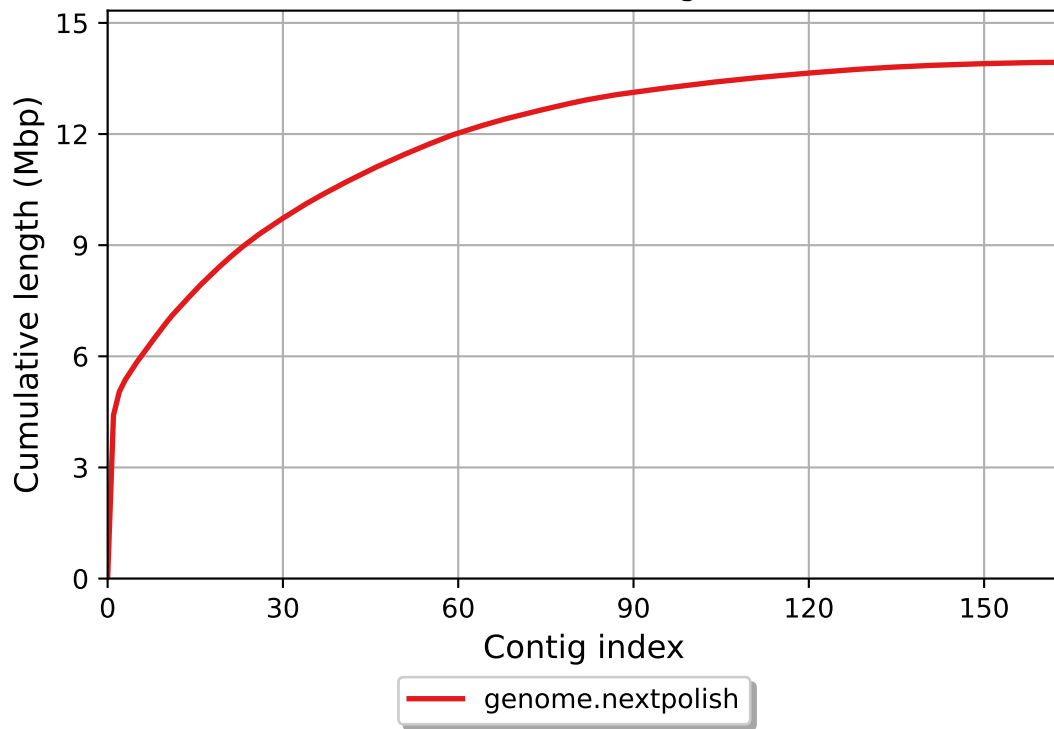
# Report

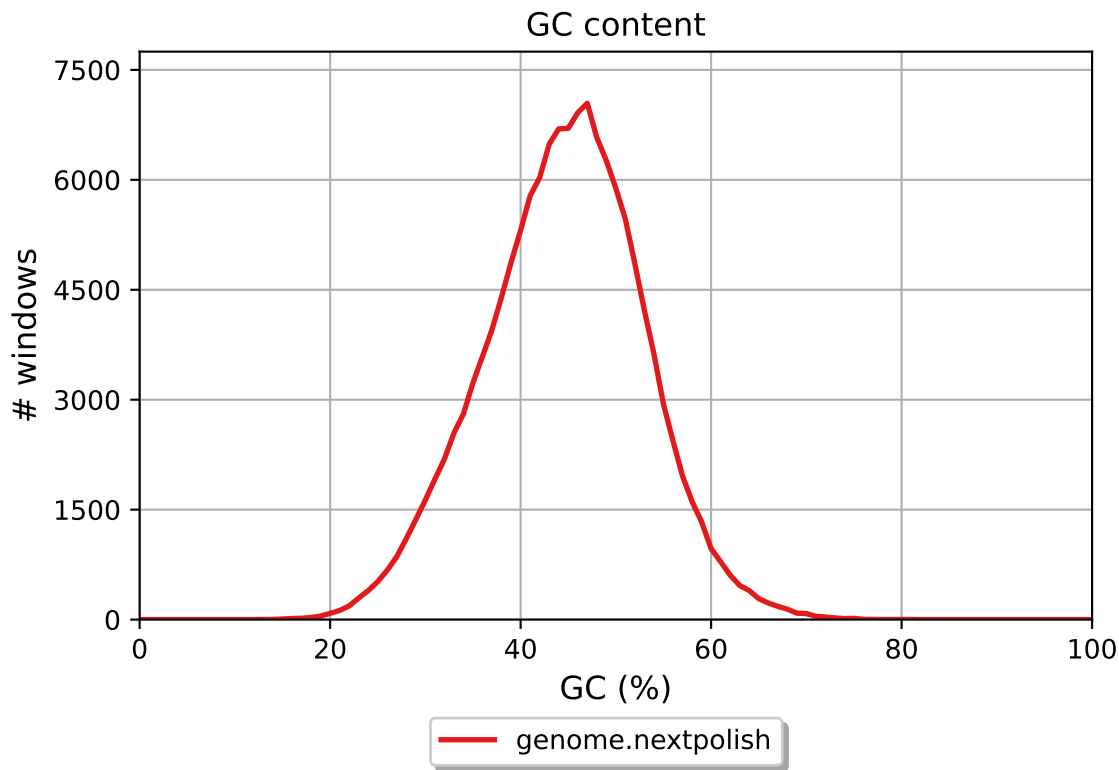
	genome.nextpolish
# contigs (>= 0 bp)	163
# contigs (>= 1000 bp)	163
# contigs (>= 5000 bp)	145
# contigs (>= 10000 bp)	131
# contigs (>= 25000 bp)	87
# contigs (>= 50000 bp)	62
Total length (>= 0 bp)	13937319
Total length (>= 1000 bp)	13937319
Total length (>= 5000 bp)	13876539
Total length (>= 10000 bp)	13775438
Total length (>= 25000 bp)	13061412
Total length (>= 50000 bp)	12128172
# contigs	163
Largest contig	4407301
Total length	13937319
GC (%)	44.54
N50	195726
N90	39038
auN	1500906.7
L50	11
L90	72
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	38 + 0 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

