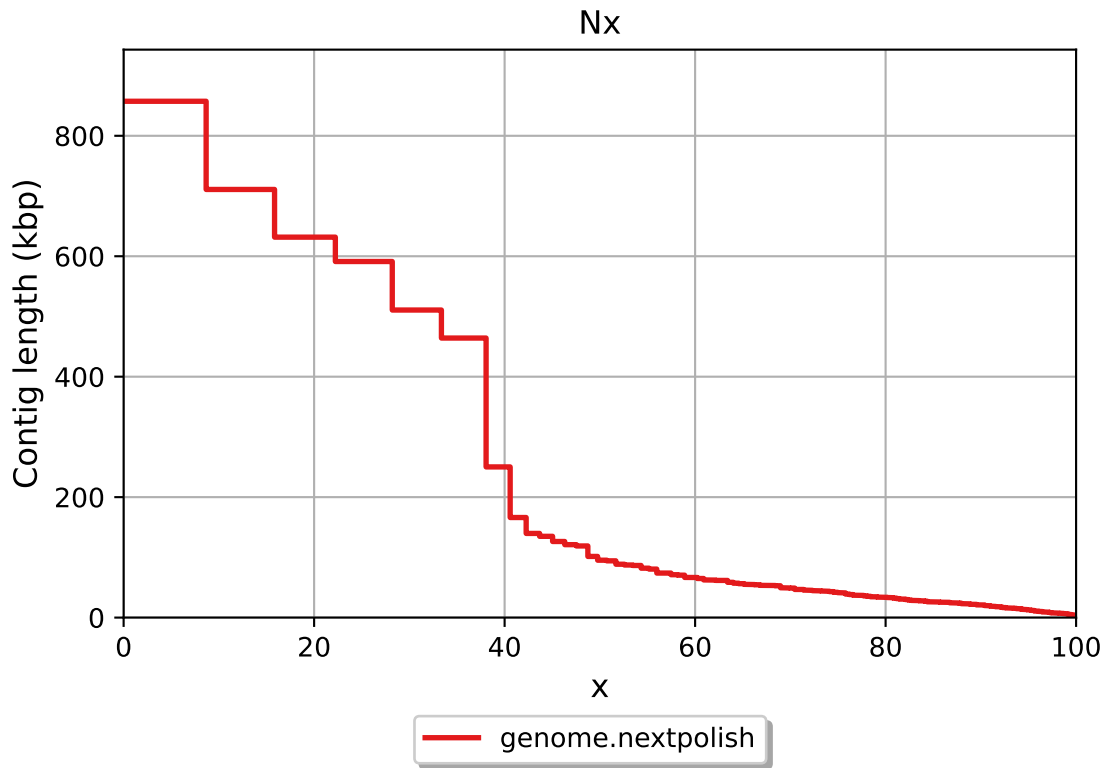


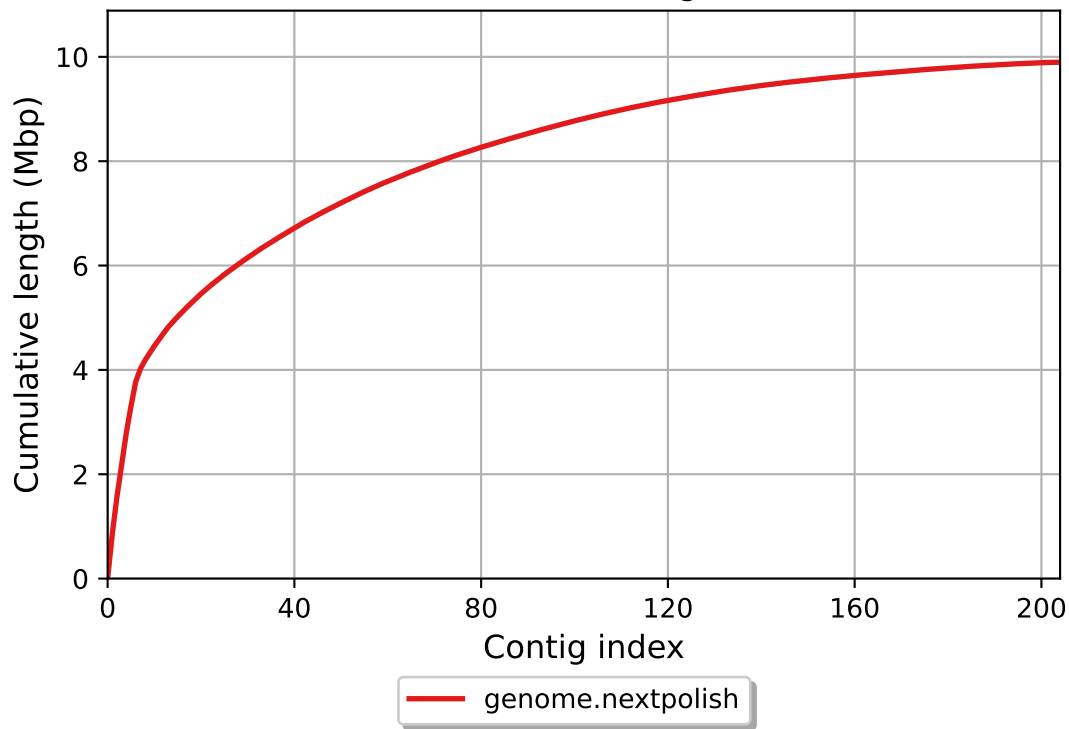
Report

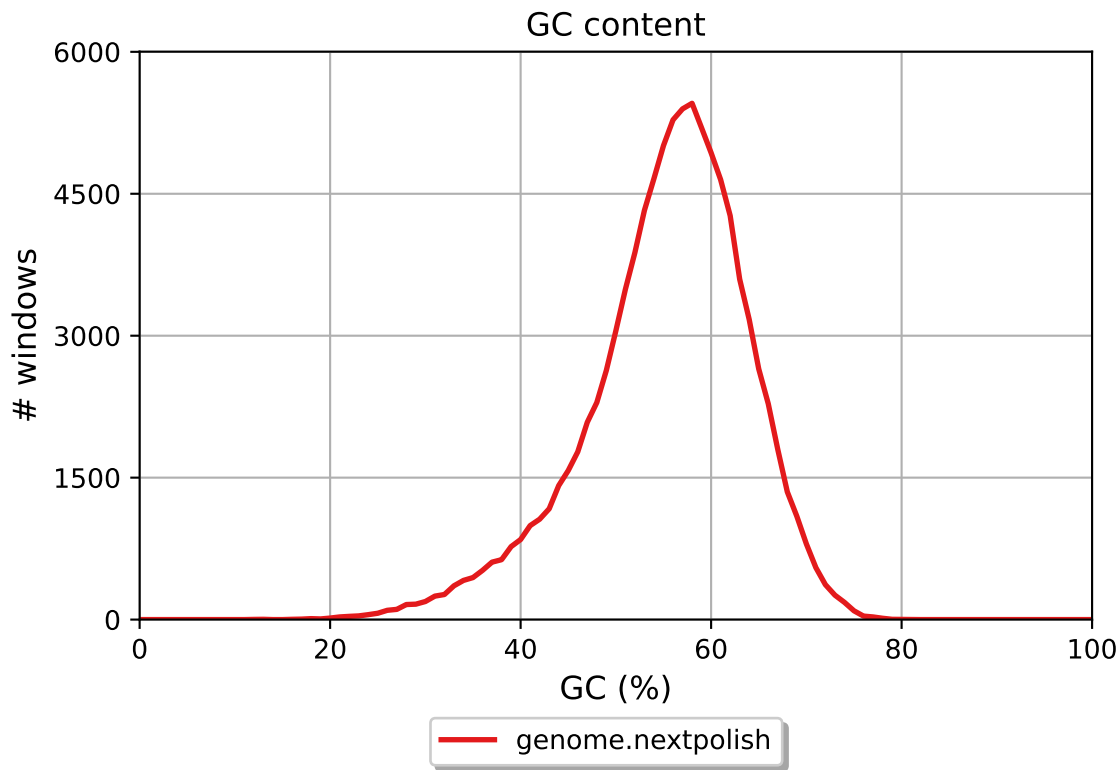
	genome.nextpolish
# contigs (>= 0 bp)	204
# contigs (>= 1000 bp)	202
# contigs (>= 5000 bp)	190
# contigs (>= 10000 bp)	149
# contigs (>= 25000 bp)	92
# contigs (>= 50000 bp)	42
Total length (>= 0 bp)	9896124
Total length (>= 1000 bp)	9894483
Total length (>= 5000 bp)	9845754
Total length (>= 10000 bp)	9546202
Total length (>= 25000 bp)	8581842
Total length (>= 50000 bp)	6824547
# contigs	204
Largest contig	857320
Total length	9896124
GC (%)	55.24
N50	95280
N90	20958
auN	289663.8
L50	15
L90	107
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 5 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

