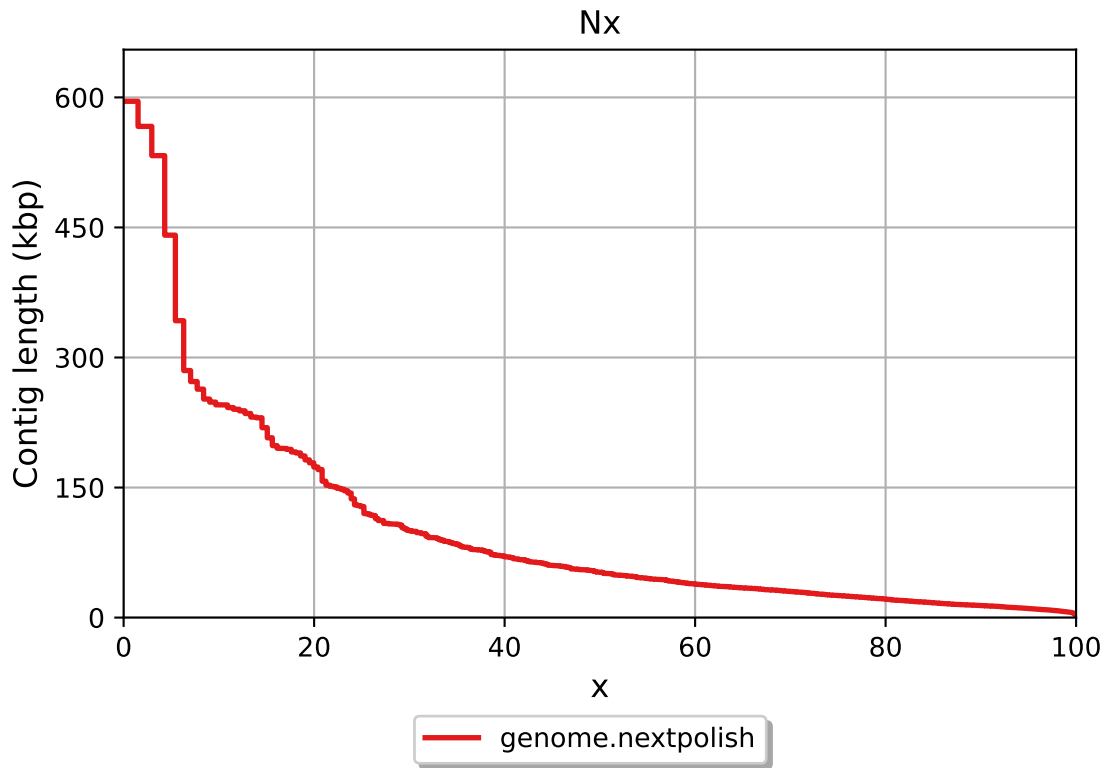


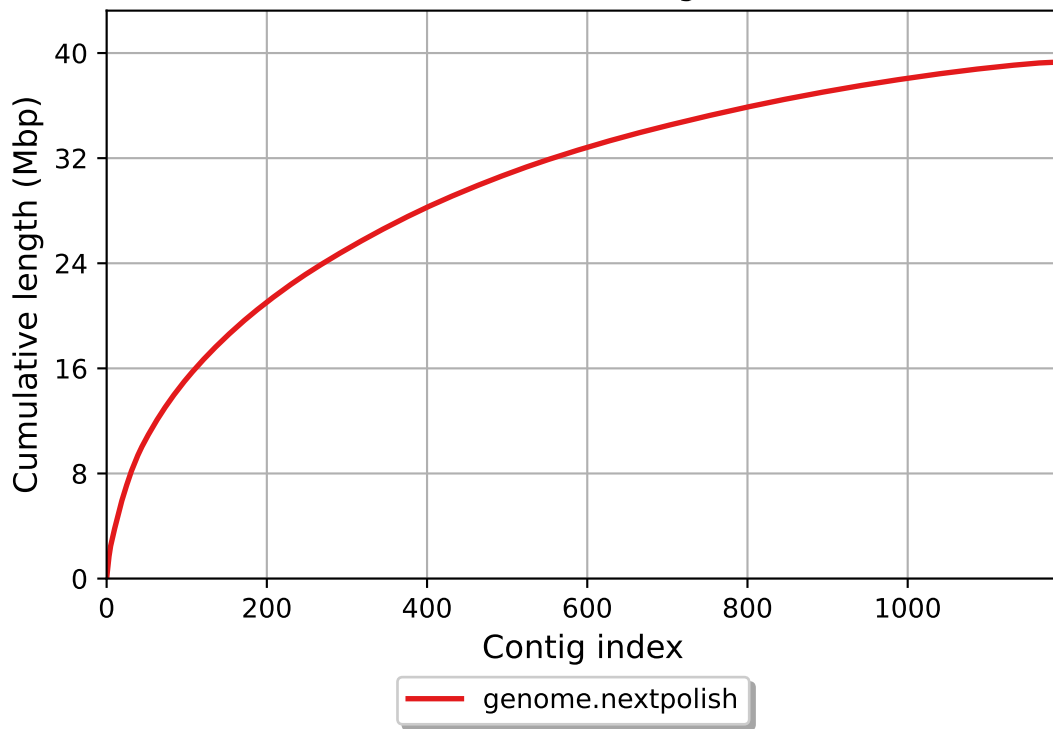
# Report

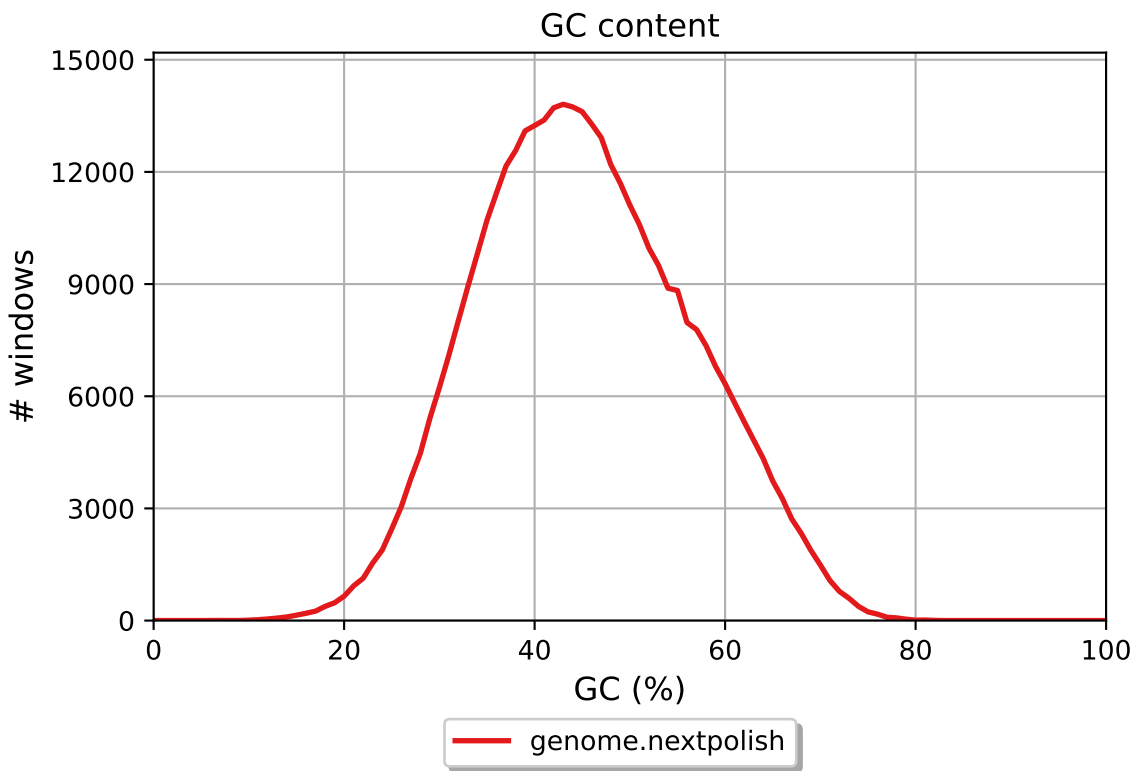
	genome.nextpolish
# contigs (>= 0 bp)	1189
# contigs (>= 1000 bp)	1182
# contigs (>= 5000 bp)	1154
# contigs (>= 10000 bp)	945
# contigs (>= 25000 bp)	456
# contigs (>= 50000 bp)	183
Total length (>= 0 bp)	39298162
Total length (>= 1000 bp)	39293659
Total length (>= 5000 bp)	39197318
Total length (>= 10000 bp)	37562621
Total length (>= 25000 bp)	29744868
Total length (>= 50000 bp)	20218580
# contigs	1189
Largest contig	595536
Total length	39298162
GC (%)	45.25
N50	52287
N90	13856
auN	104401.2
L50	172
L90	762
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	79 + 7 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

