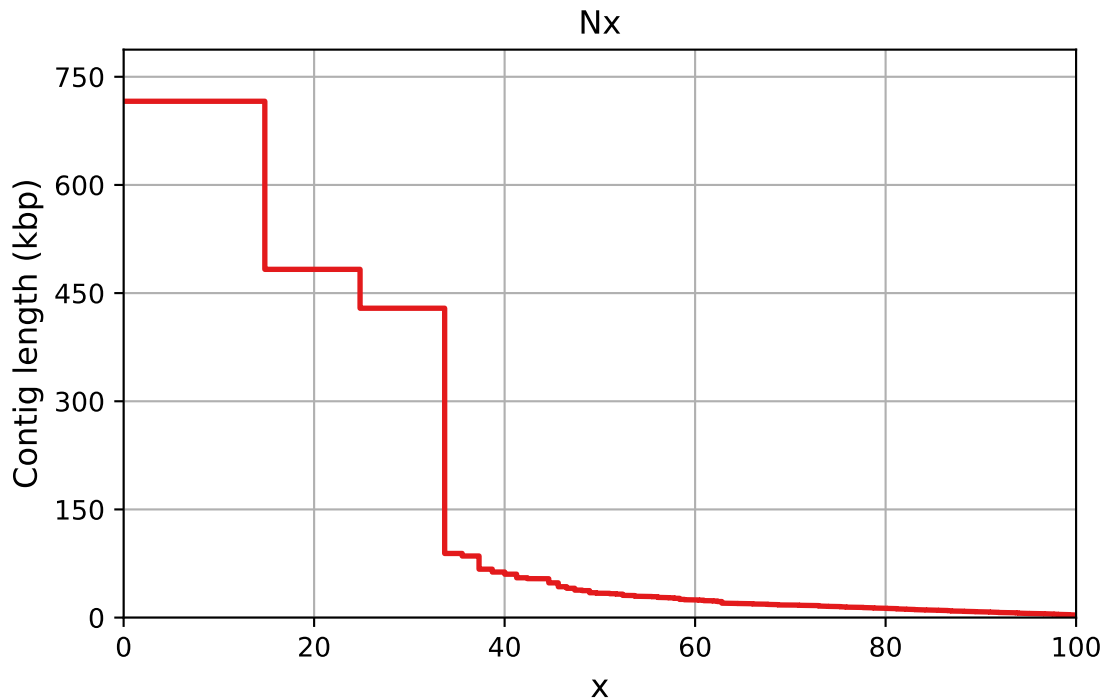


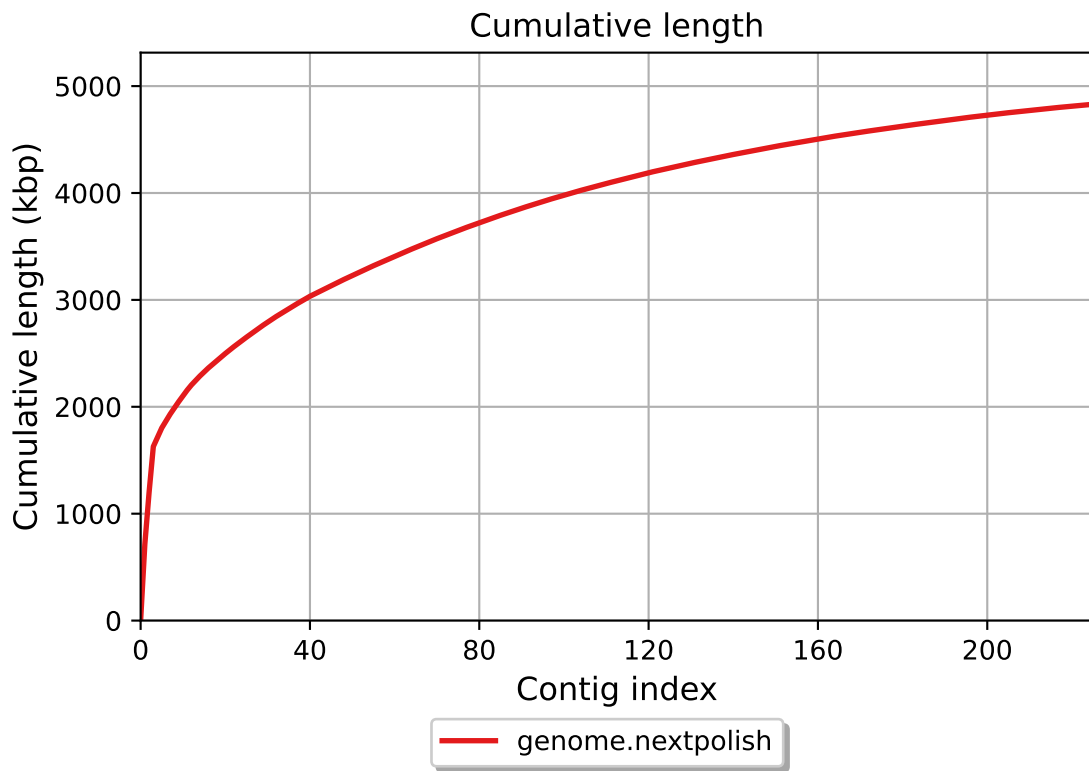
# Report

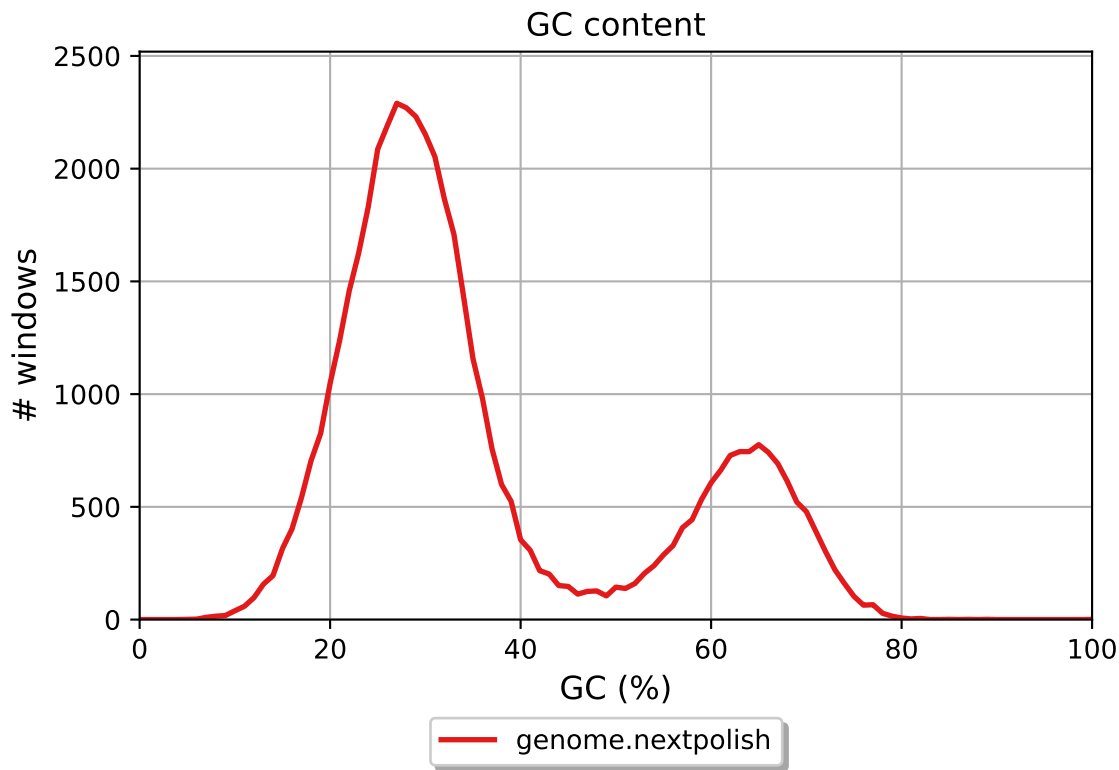
	genome.nextpolish
# contigs (>= 0 bp)	225
# contigs (>= 1000 bp)	225
# contigs (>= 5000 bp)	194
# contigs (>= 10000 bp)	118
# contigs (>= 25000 bp)	32
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	4830025
Total length (>= 1000 bp)	4830025
Total length (>= 5000 bp)	4699288
Total length (>= 10000 bp)	4169818
Total length (>= 25000 bp)	2843957
Total length (>= 50000 bp)	2155945
# contigs	225
Largest contig	716038
Total length	4830025
GC (%)	36.66
N50	33777
N90	7871
auN	210271.5
L50	18
L90	139
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 6 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).



— genome.nextpolish





genome.nextpolish GC content

