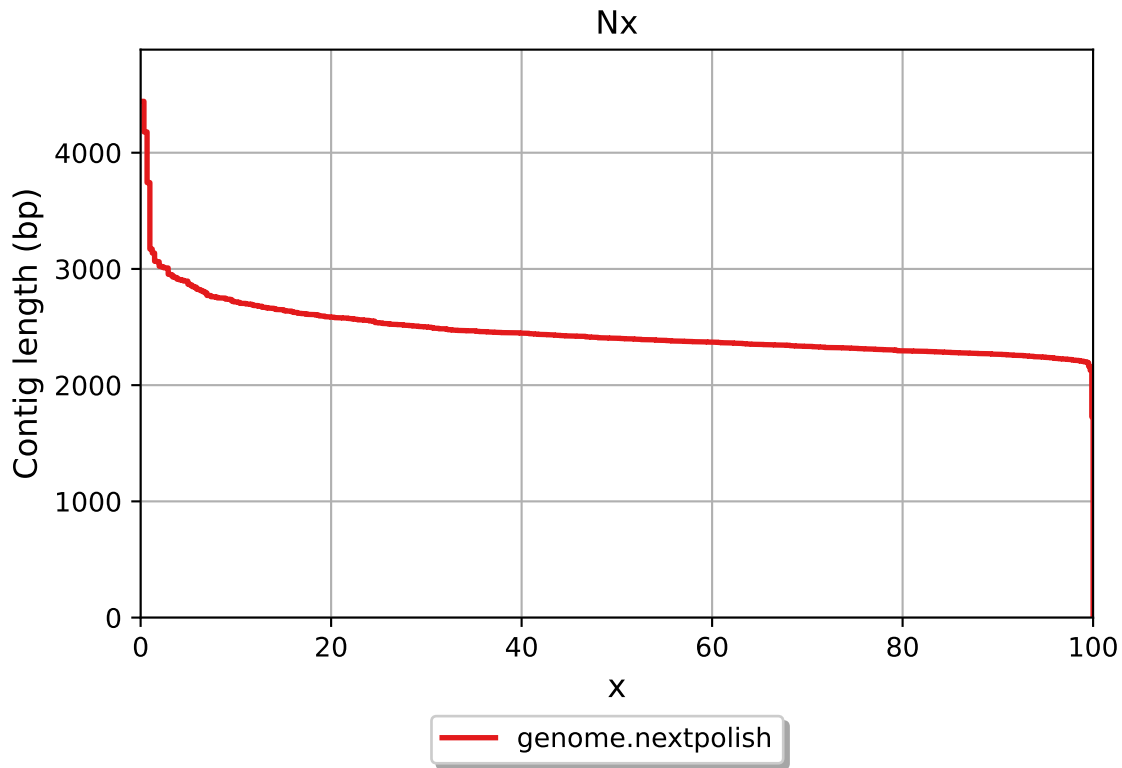
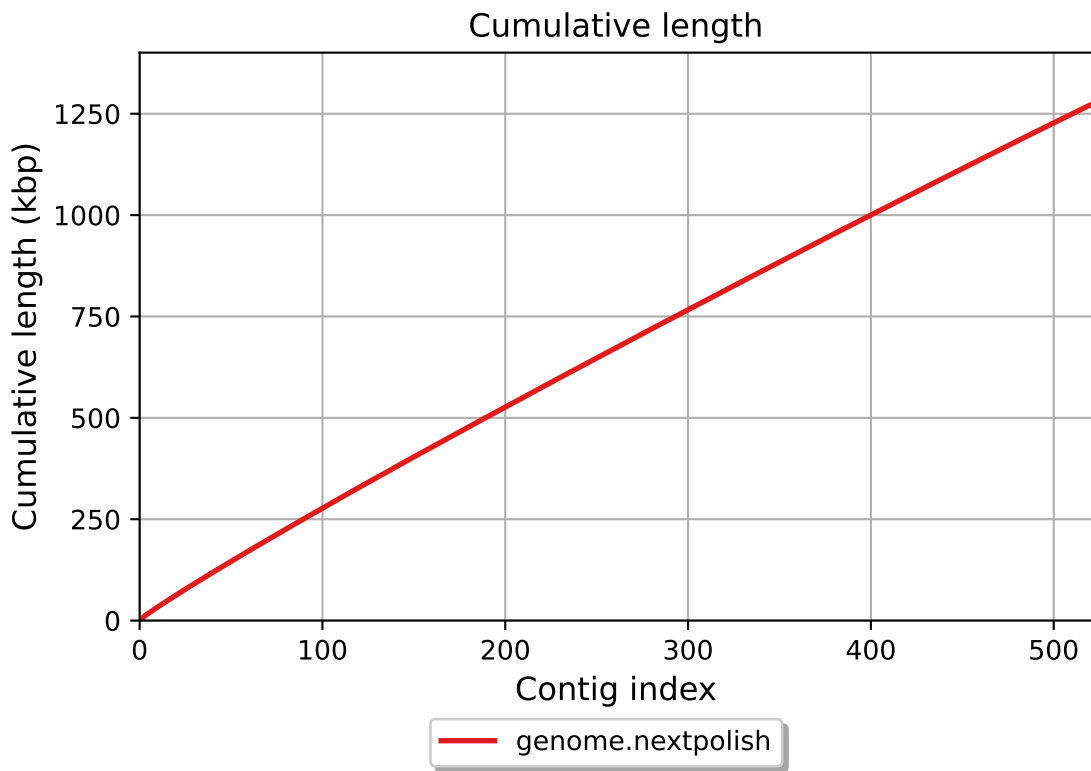


Report

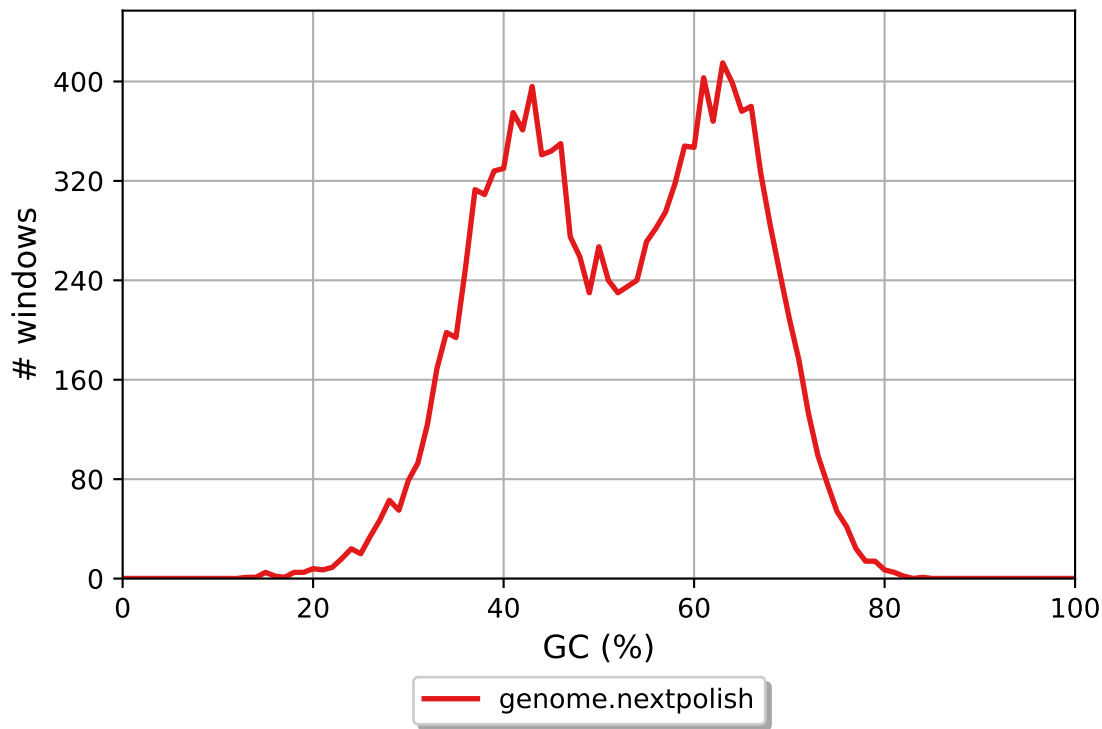
	genome.nextpolish
# contigs (≥ 0 bp)	521
# contigs (≥ 1000 bp)	521
# contigs (≥ 5000 bp)	0
# contigs (≥ 10000 bp)	0
# contigs (≥ 25000 bp)	0
# contigs (≥ 50000 bp)	0
Total length (≥ 0 bp)	1273375
Total length (≥ 1000 bp)	1273375
Total length (≥ 5000 bp)	0
Total length (≥ 10000 bp)	0
Total length (≥ 25000 bp)	0
Total length (≥ 50000 bp)	0
# contigs	521
Largest contig	4442
Total length	1273375
GC (%)	52.12
N50	2404
N90	2266
auN	2463.8
L50	246
L90	464
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 4 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).





GC content



genome.nextpolish GC content

