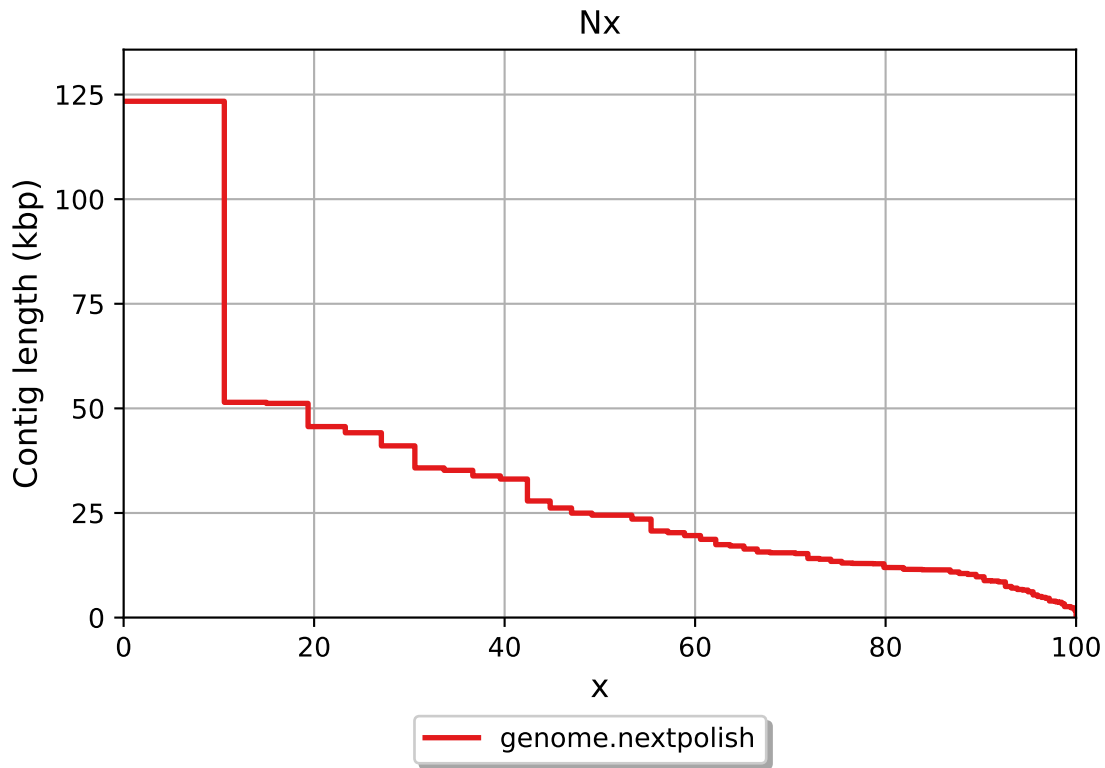
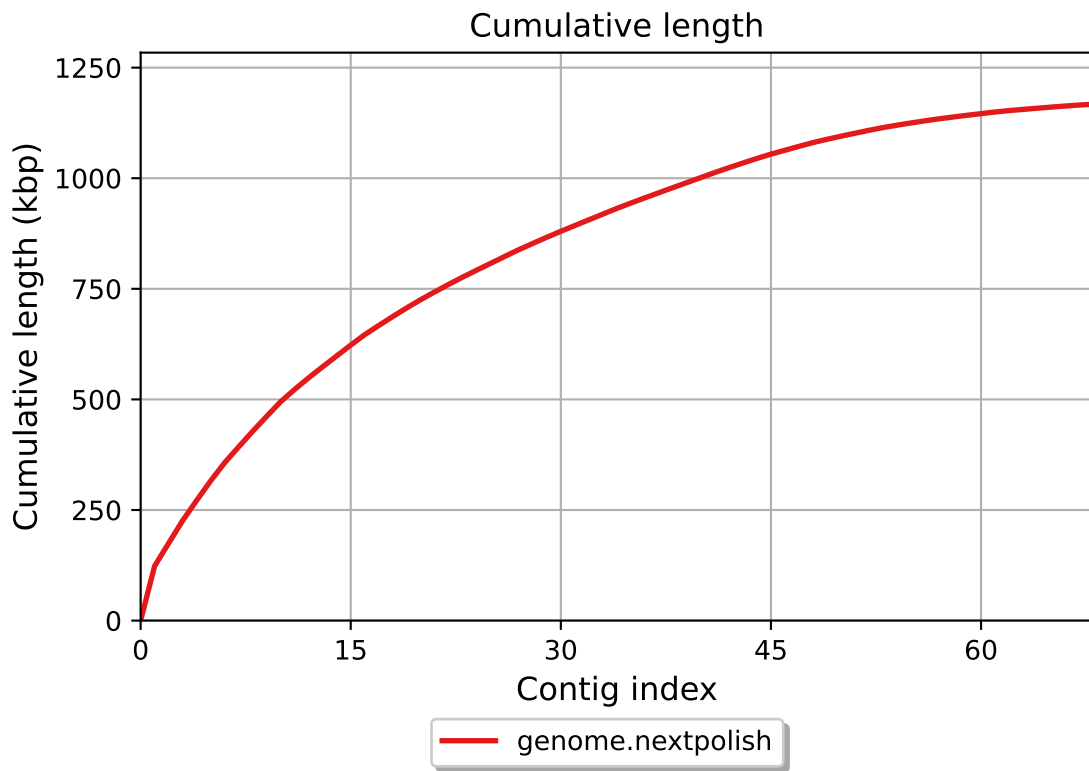


# Report

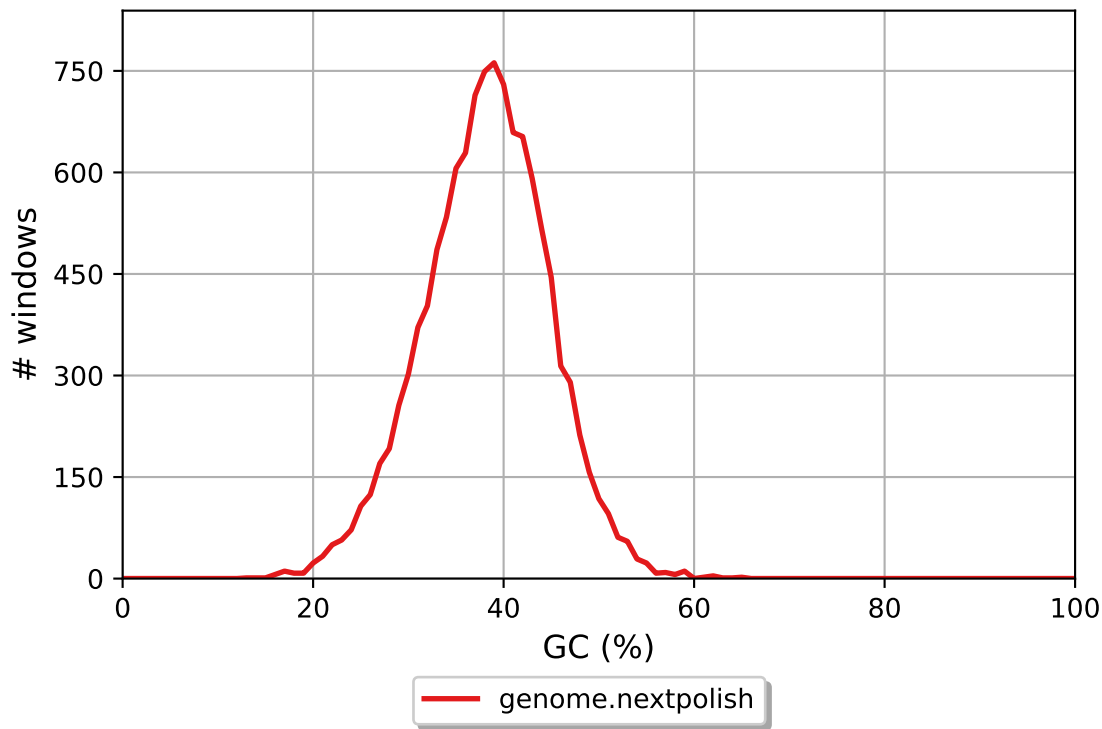
	genome.nextpolish
# contigs (>= 0 bp)	68
# contigs (>= 1000 bp)	68
# contigs (>= 5000 bp)	55
# contigs (>= 10000 bp)	44
# contigs (>= 25000 bp)	12
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	1167185
Total length (>= 1000 bp)	1167185
Total length (>= 5000 bp)	1124938
Total length (>= 10000 bp)	1044689
Total length (>= 25000 bp)	548891
Total length (>= 50000 bp)	226034
# contigs	68
Largest contig	123396
Total length	1167185
GC (%)	38.17
N50	24477
N90	9751
auN	35546.0
L50	14
L90	45
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).





## GC content



genome.nextpolish GC content

