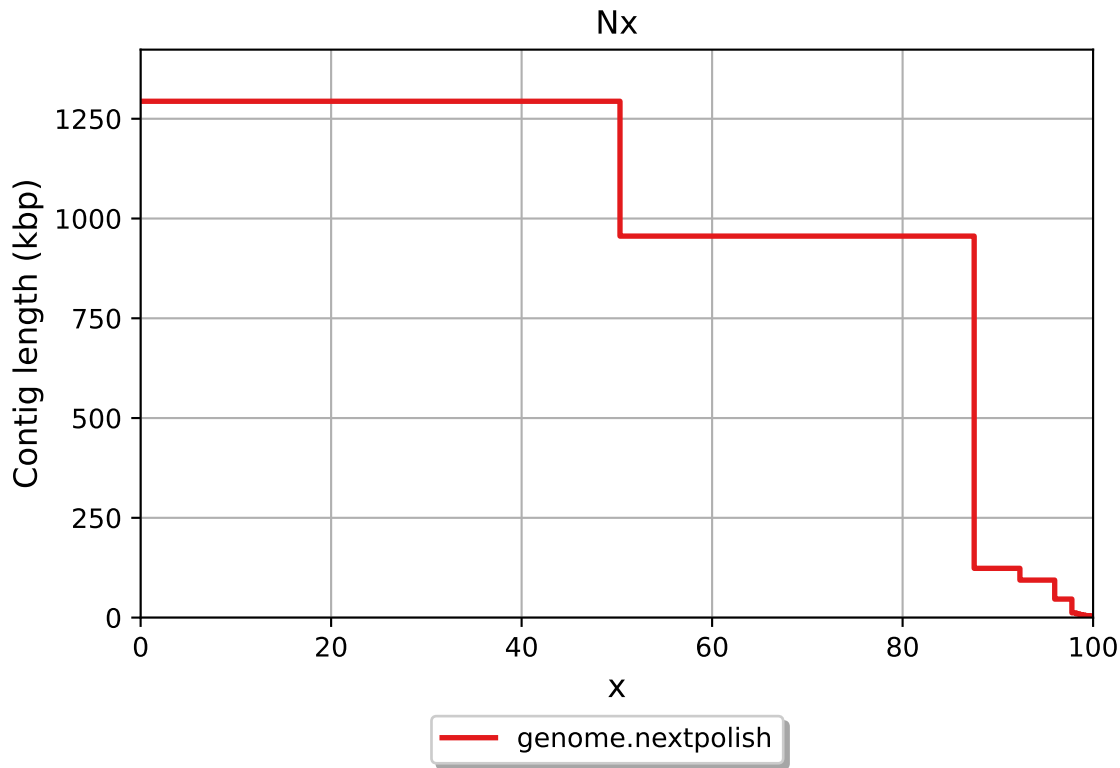
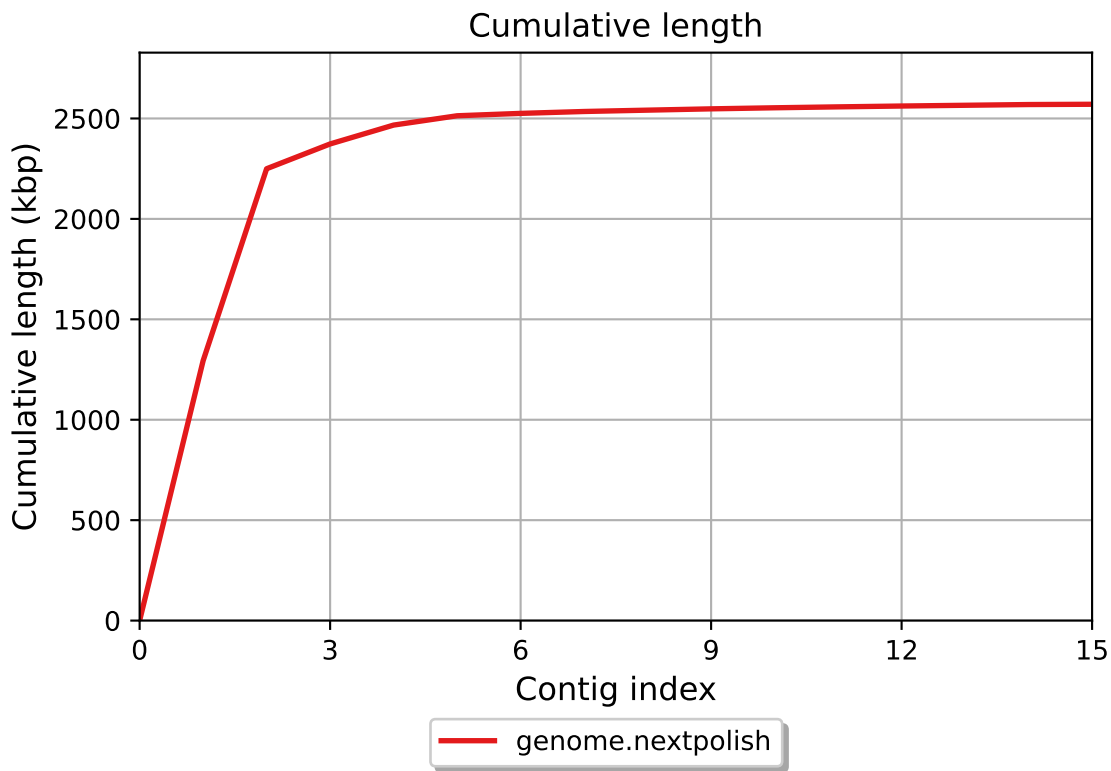


Report

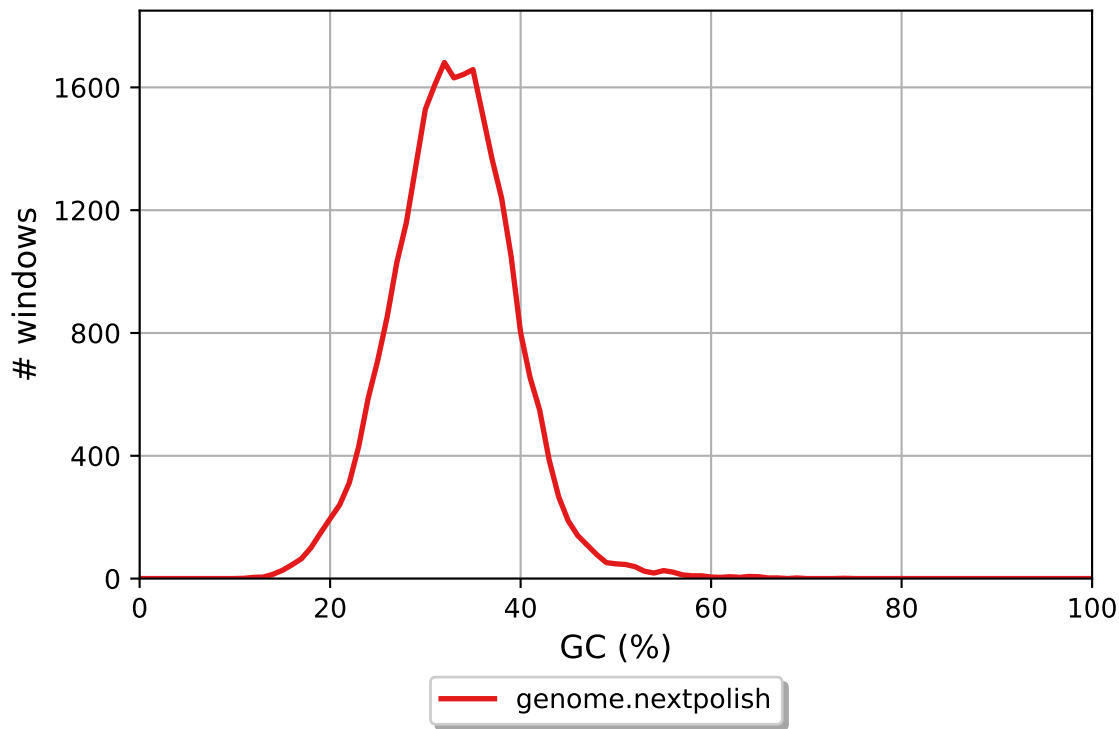
	genome.nextpolish
# contigs (>= 0 bp)	15
# contigs (>= 1000 bp)	15
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2570713
Total length (>= 1000 bp)	2570713
Total length (>= 5000 bp)	2553210
Total length (>= 10000 bp)	2525325
Total length (>= 25000 bp)	2513453
Total length (>= 50000 bp)	2467083
# contigs	15
Largest contig	1293807
Total length	2570713
GC (%)	33.02
N50	1293807
N90	123425
auN	1016974.4
L50	1
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 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

