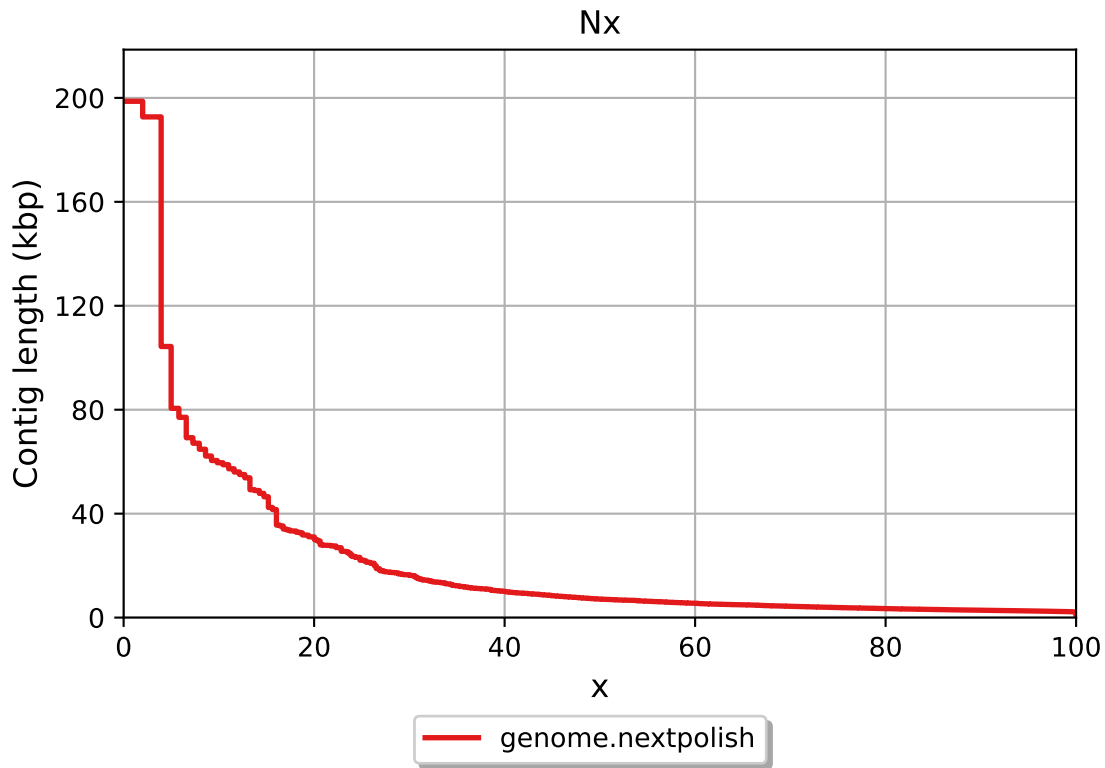


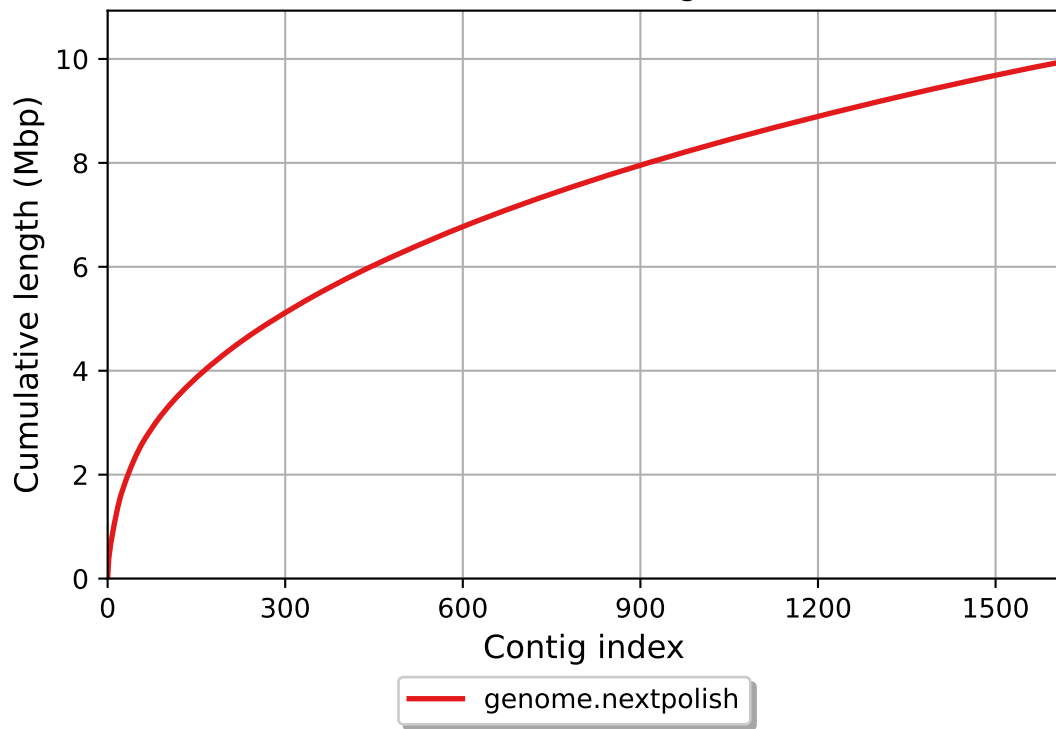
Report

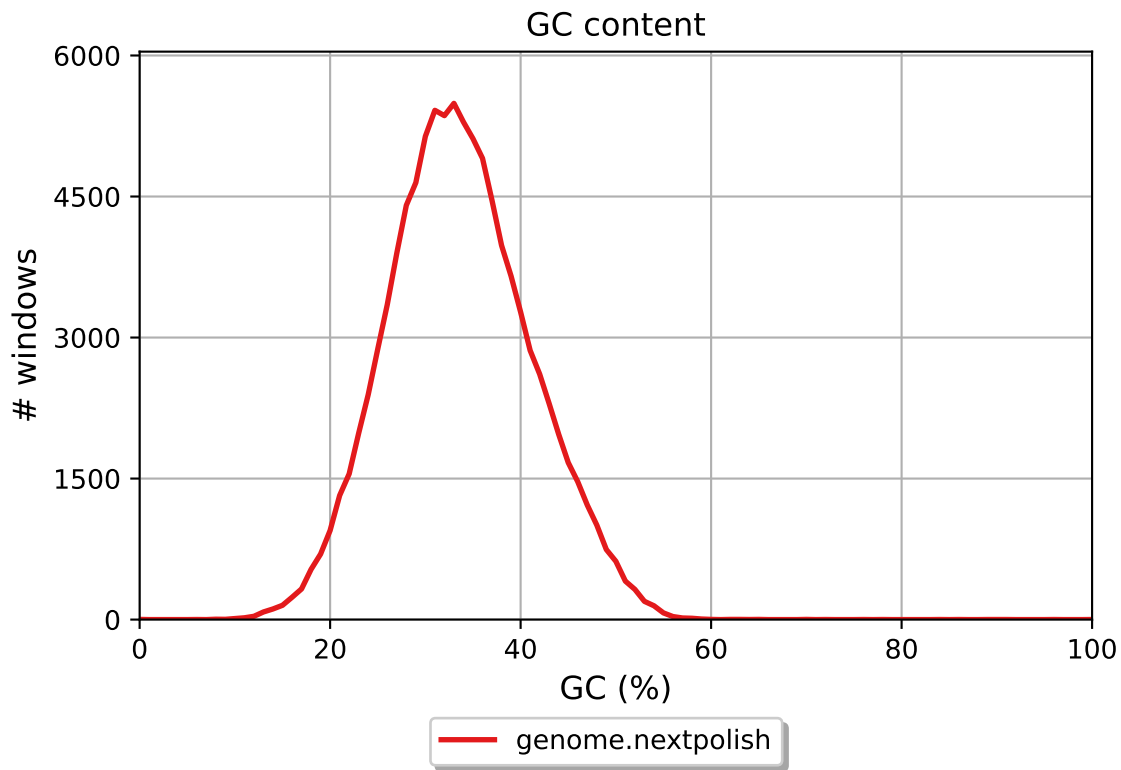
	genome.nextpolish
# contigs (>= 0 bp)	1609
# contigs (>= 1000 bp)	1608
# contigs (>= 5000 bp)	513
# contigs (>= 10000 bp)	161
# contigs (>= 25000 bp)	47
# contigs (>= 50000 bp)	16
Total length (>= 0 bp)	9936071
Total length (>= 1000 bp)	9935503
Total length (>= 5000 bp)	6356684
Total length (>= 10000 bp)	3980134
Total length (>= 25000 bp)	2346592
Total length (>= 50000 bp)	1317627
# contigs	1609
Largest contig	198701
Total length	9936071
GC (%)	33.58
N50	7117
N90	2840
auN	23144.0
L50	279
L90	1217
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 2 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

