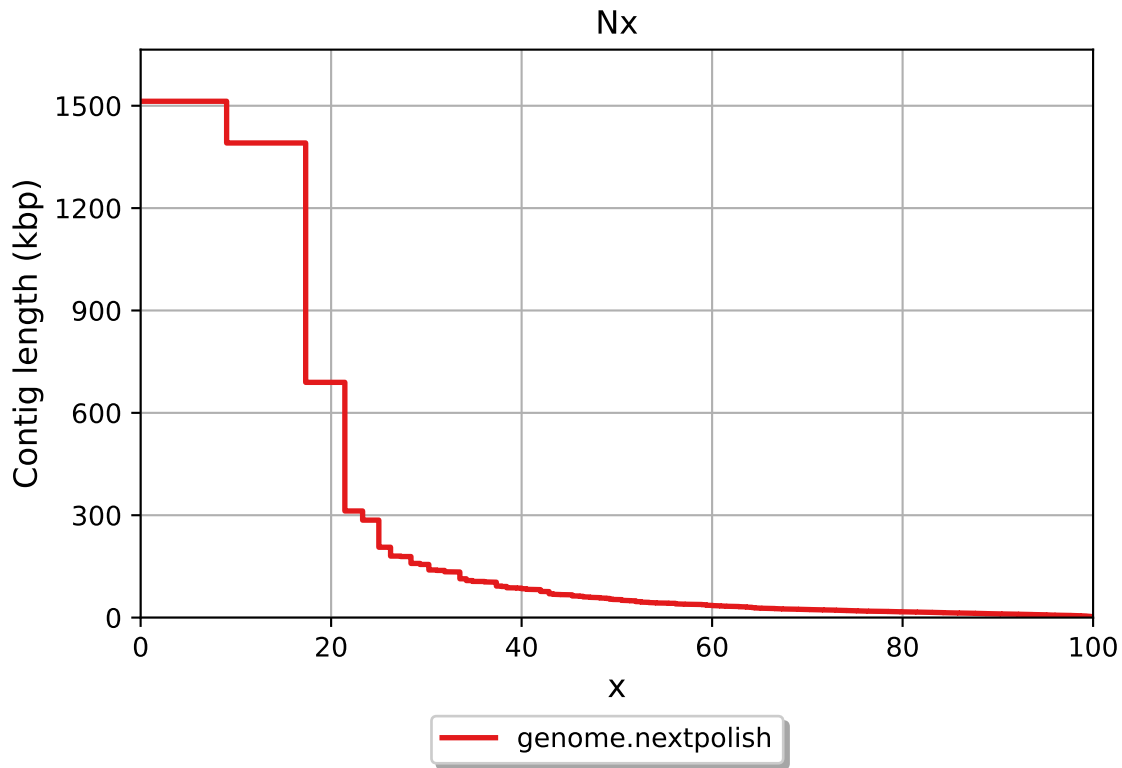


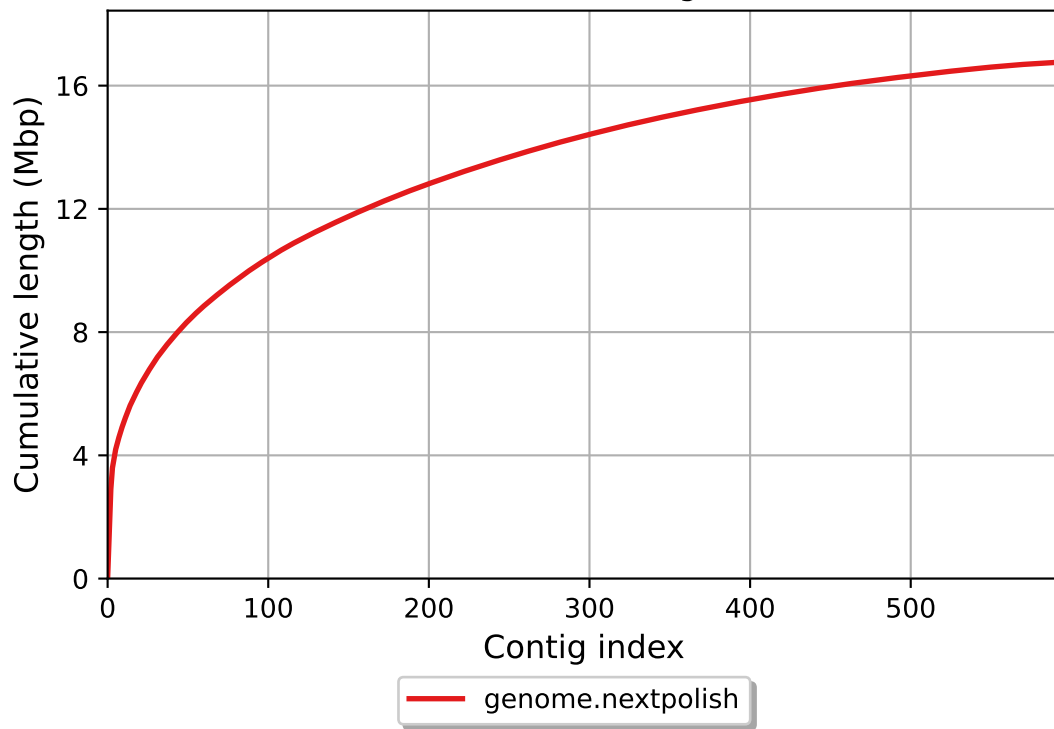
# Report

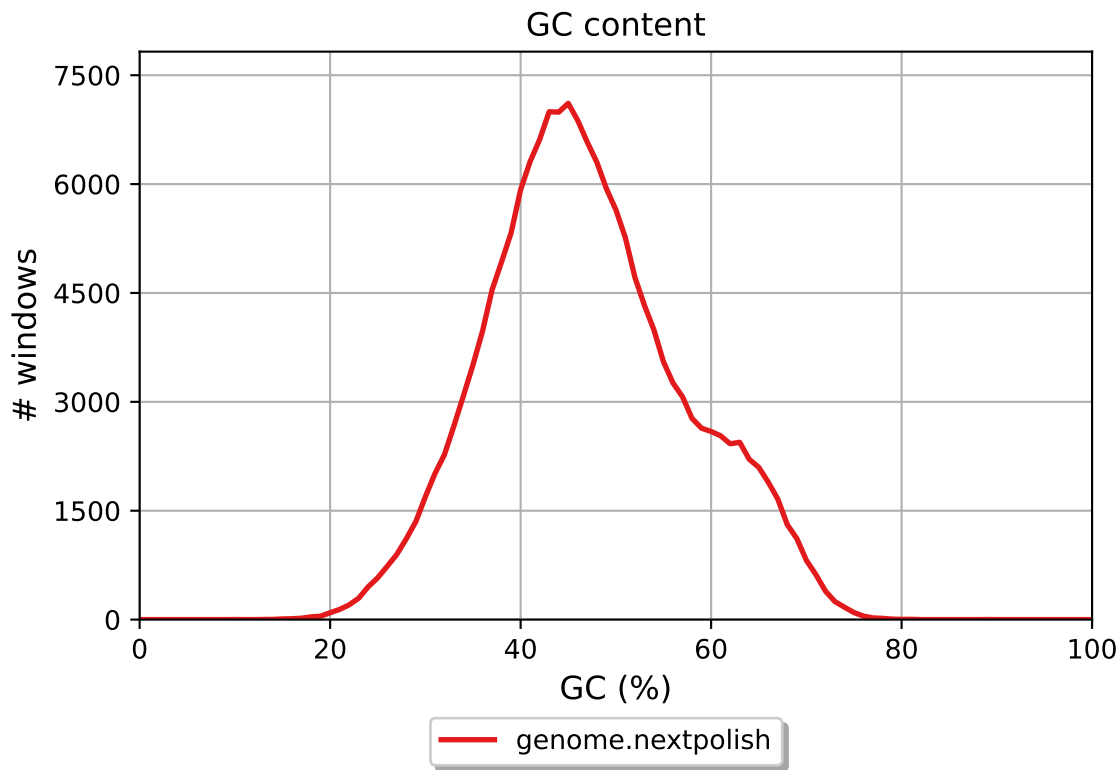
	genome.nextpolish
# contigs (>= 0 bp)	593
# contigs (>= 1000 bp)	593
# contigs (>= 5000 bp)	548
# contigs (>= 10000 bp)	385
# contigs (>= 25000 bp)	133
# contigs (>= 50000 bp)	53
Total length (>= 0 bp)	16758414
Total length (>= 1000 bp)	16758414
Total length (>= 5000 bp)	16591791
Total length (>= 10000 bp)	15397384
Total length (>= 25000 bp)	11341981
Total length (>= 50000 bp)	8514123
# contigs	593
Largest contig	1513132
Total length	16758414
GC (%)	46.93
N50	52918
N90	10955
auN	329450.7
L50	51
L90	355
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	41 + 3 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

