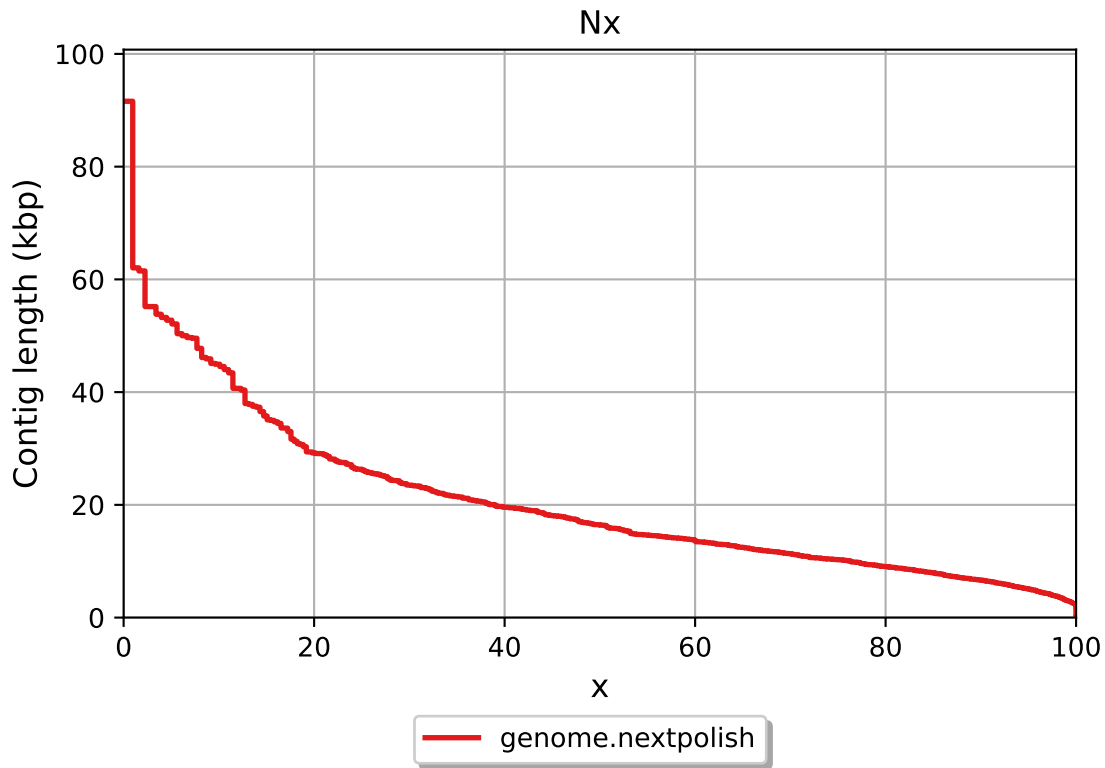


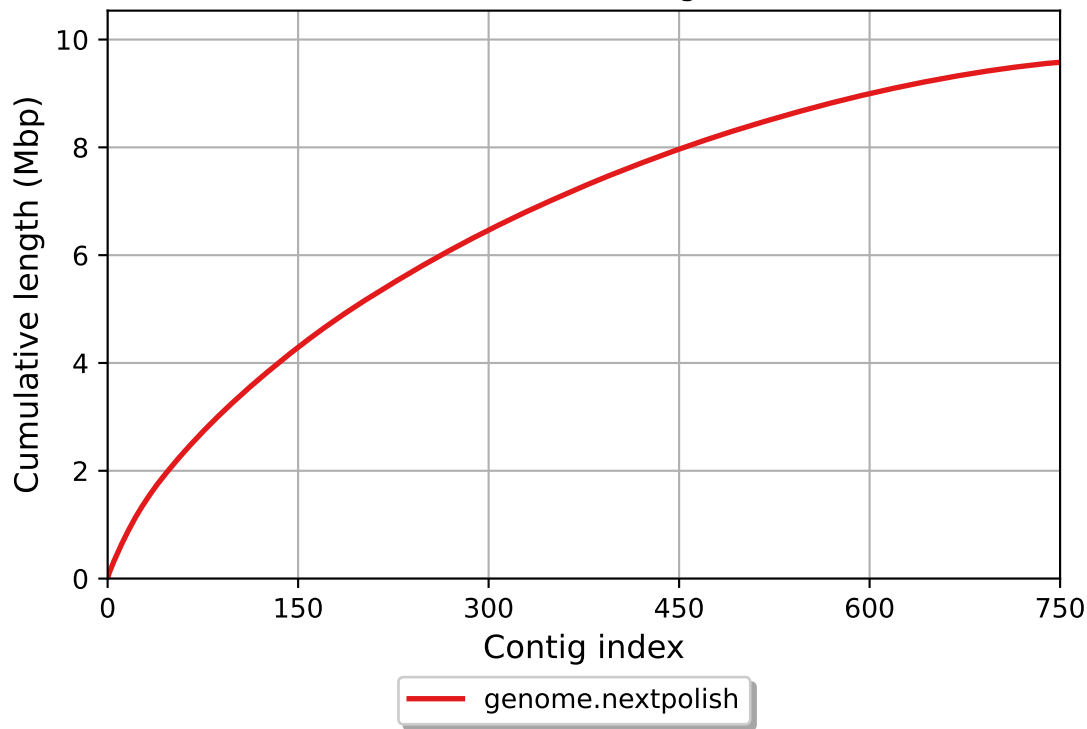
# Report

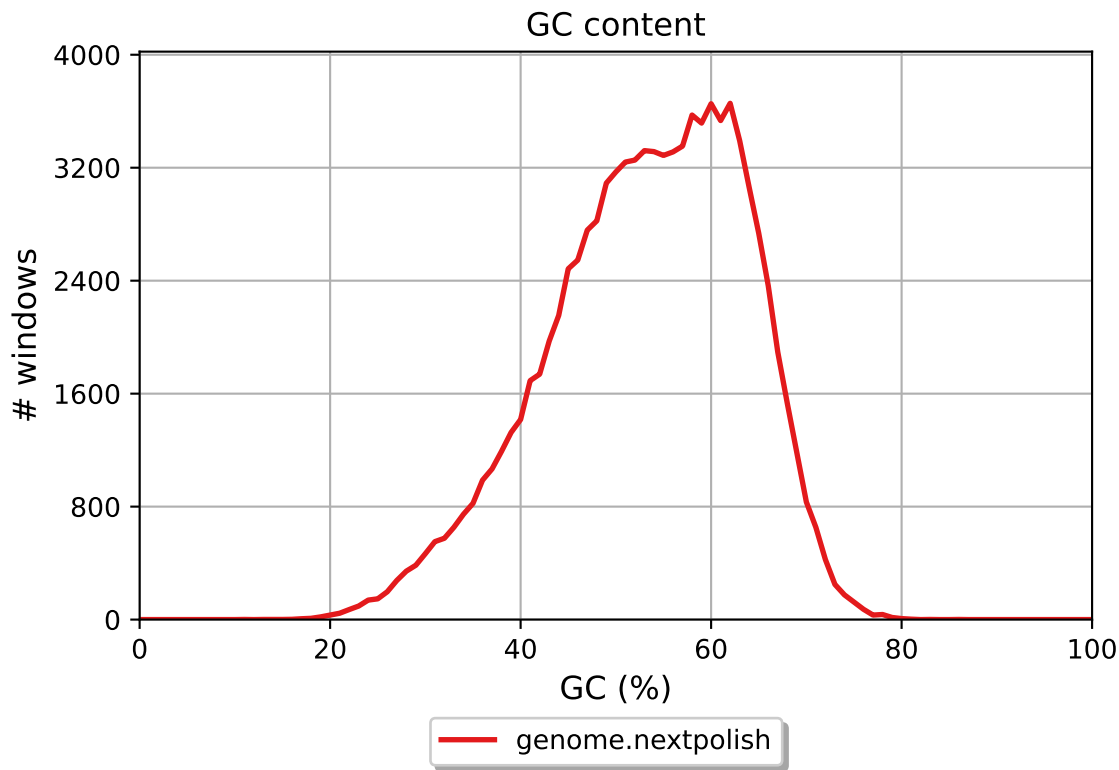
	genome.nextpolish
# contigs (>= 0 bp)	750
# contigs (>= 1000 bp)	748
# contigs (>= 5000 bp)	624
# contigs (>= 10000 bp)	378
# contigs (>= 25000 bp)	72
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	9578496
Total length (>= 1000 bp)	9577329
Total length (>= 5000 bp)	9121793
Total length (>= 10000 bp)	7311101
Total length (>= 25000 bp)	2652563
Total length (>= 50000 bp)	637781
# contigs	750
Largest contig	91599
Total length	9578496
GC (%)	53.17
N50	16460
N90	6662
auN	20828.3
L50	179
L90	538
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 5 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

