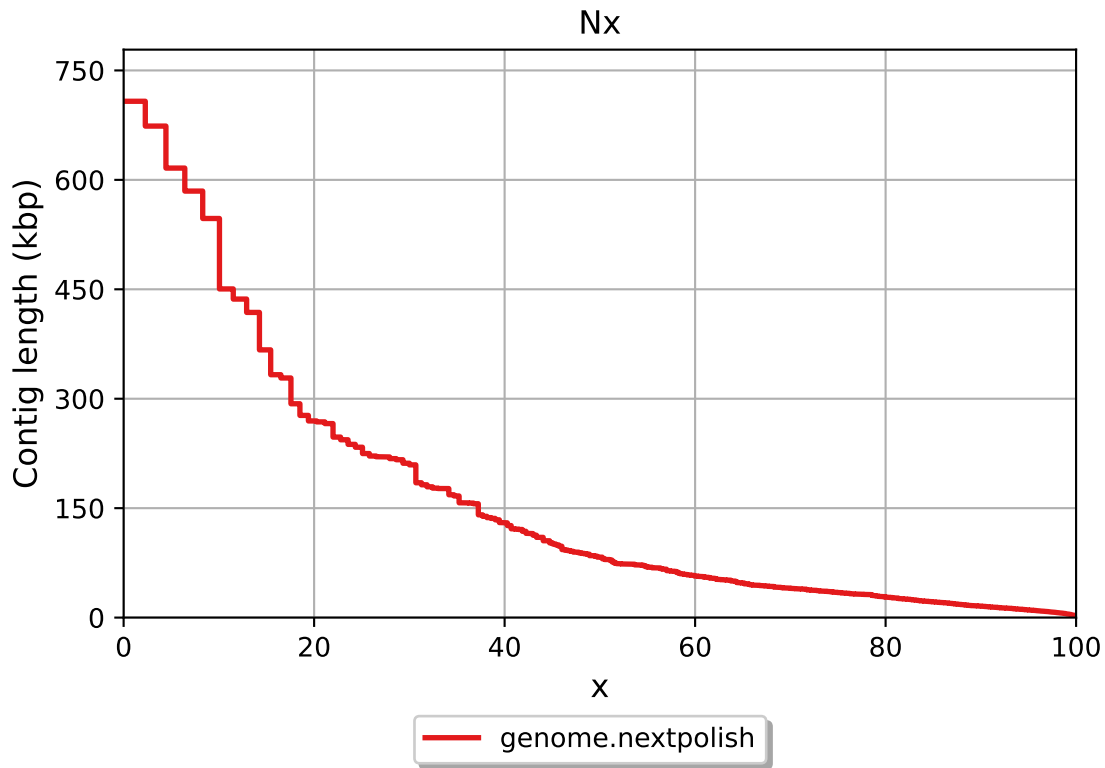


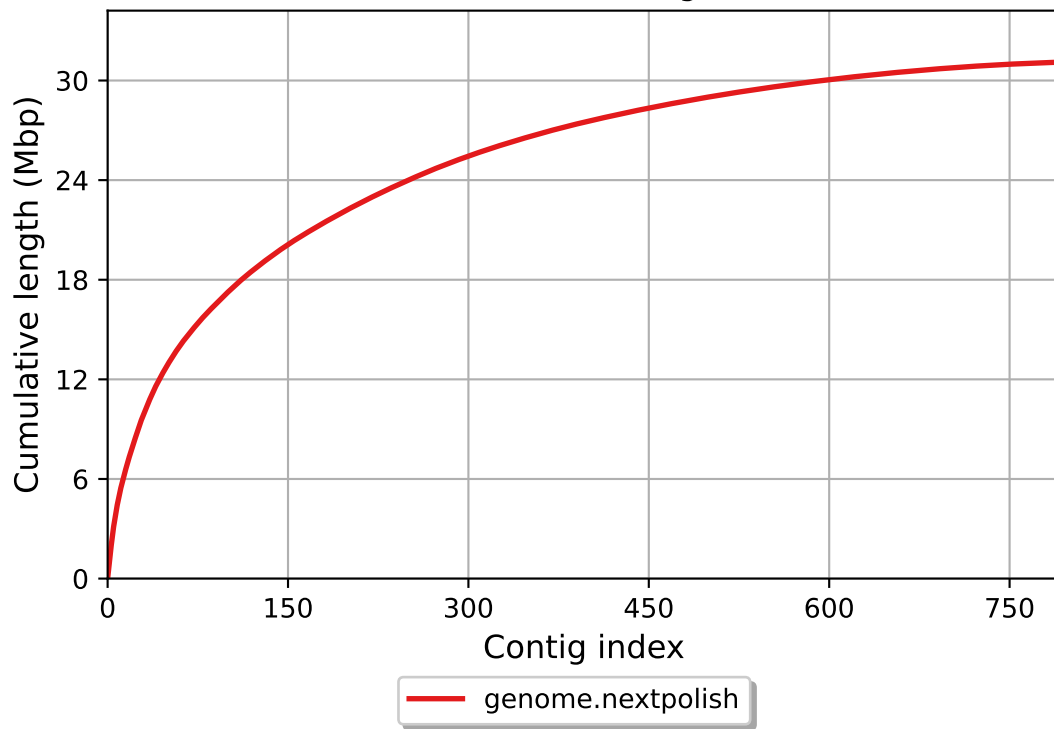
Report

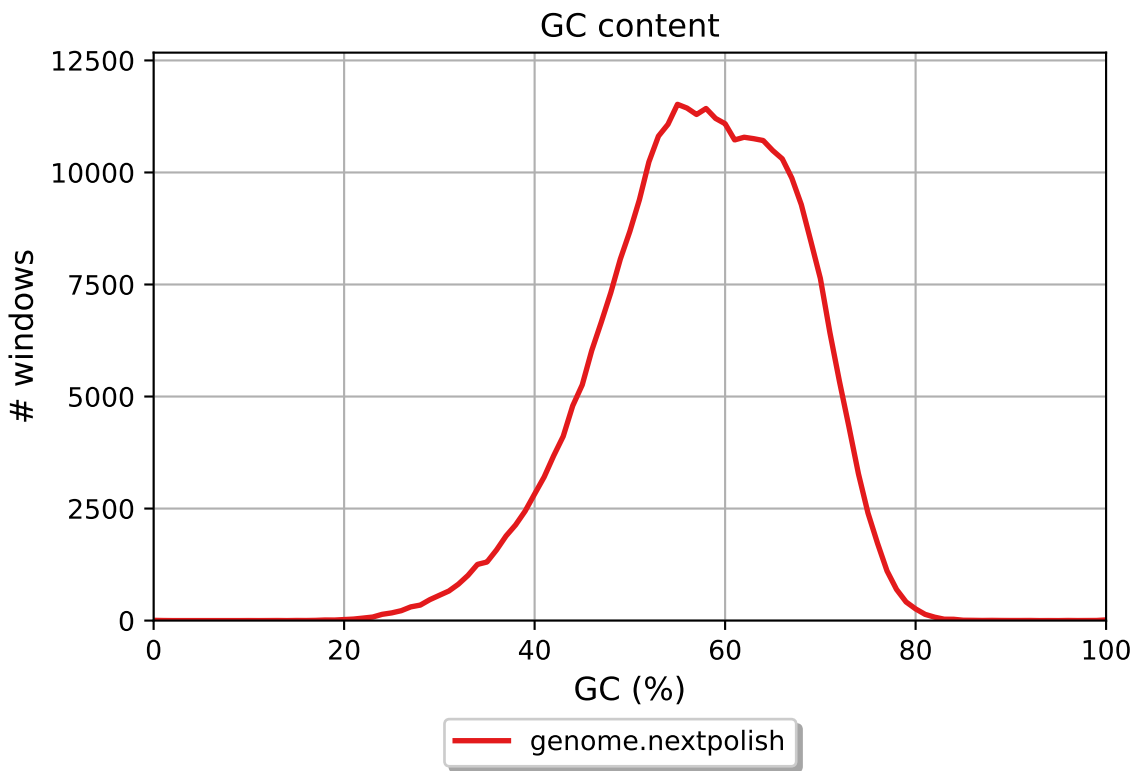
	genome.nextpolish
# contigs (>= 0 bp)	792
# contigs (>= 1000 bp)	789
# contigs (>= 5000 bp)	709
# contigs (>= 10000 bp)	562
# contigs (>= 25000 bp)	308
# contigs (>= 50000 bp)	145
Total length (>= 0 bp)	31099004
Total length (>= 1000 bp)	31096626
Total length (>= 5000 bp)	30802474
Total length (>= 10000 bp)	29699131
Total length (>= 25000 bp)	25649873
Total length (>= 50000 bp)	19874947
# contigs	792
Largest contig	707734
Total length	31099004
GC (%)	57.28
N50	82811
N90	15649
auN	168593.3
L50	78
L90	427
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	66 + 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

