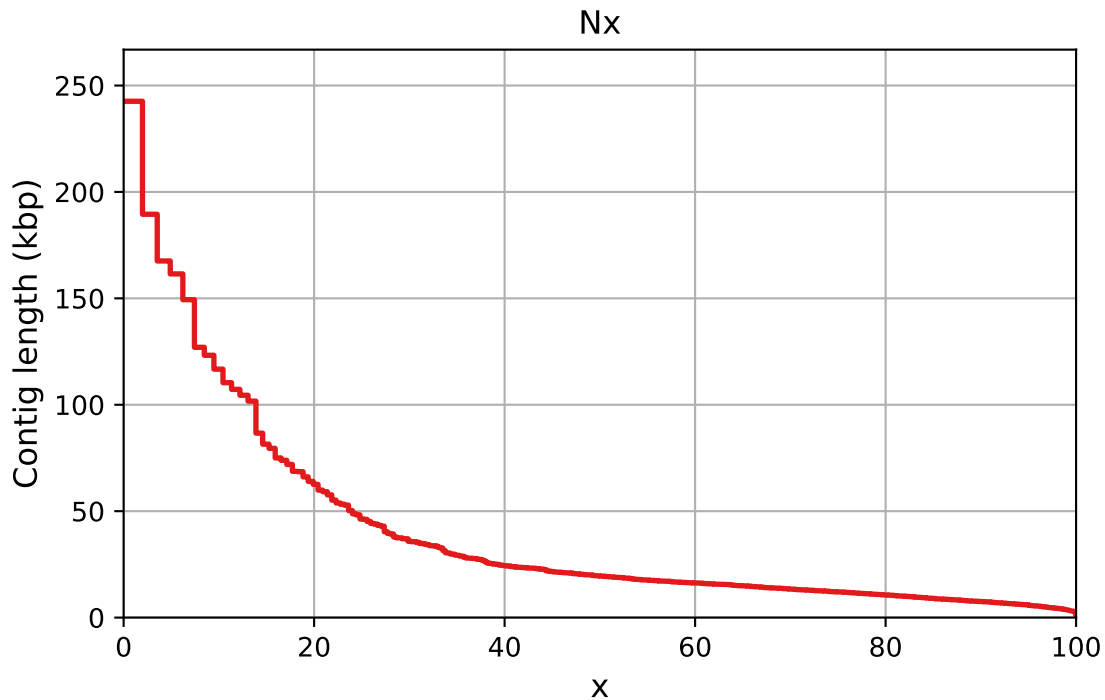


Report

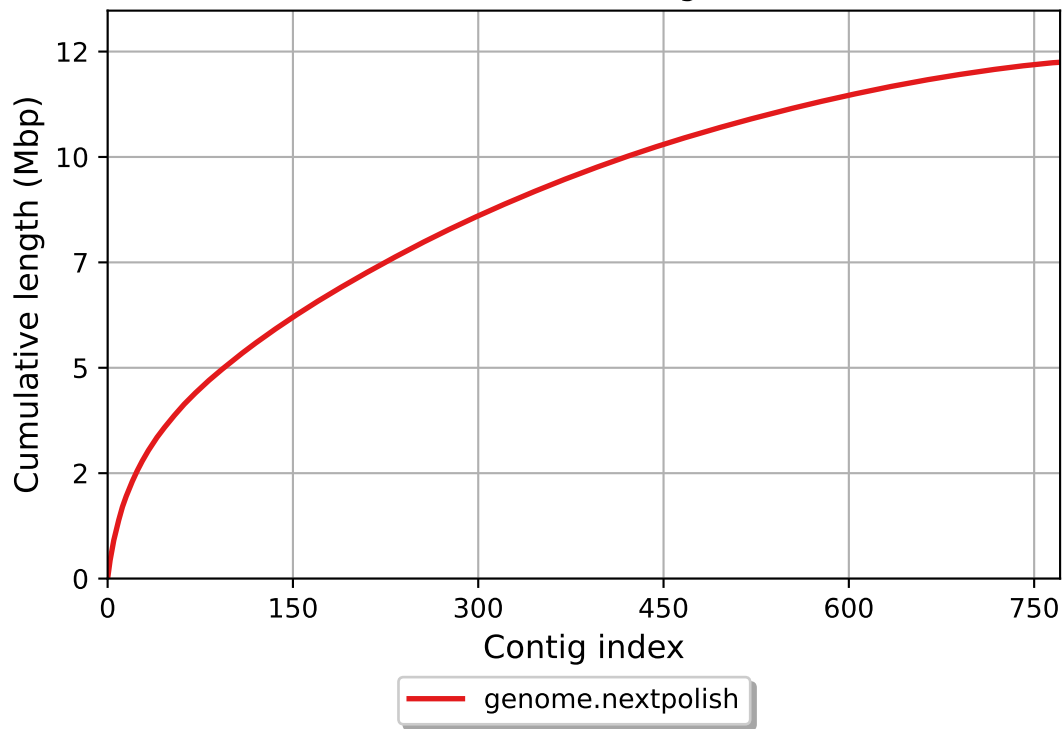
	genome.nextpolish
# contigs (>= 0 bp)	771
# contigs (>= 1000 bp)	771
# contigs (>= 5000 bp)	667
# contigs (>= 10000 bp)	427
# contigs (>= 25000 bp)	86
# contigs (>= 50000 bp)	31
Total length (>= 0 bp)	12246050
Total length (>= 1000 bp)	12246050
Total length (>= 5000 bp)	11848414
Total length (>= 10000 bp)	10077983
Total length (>= 25000 bp)	4801849
Total length (>= 50000 bp)	2941871
# contigs	771
Largest contig	242600
Total length	12246050
GC (%)	49.69
N50	19520
N90	7525
auN	41762.9
L50	147
L90	537
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 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).

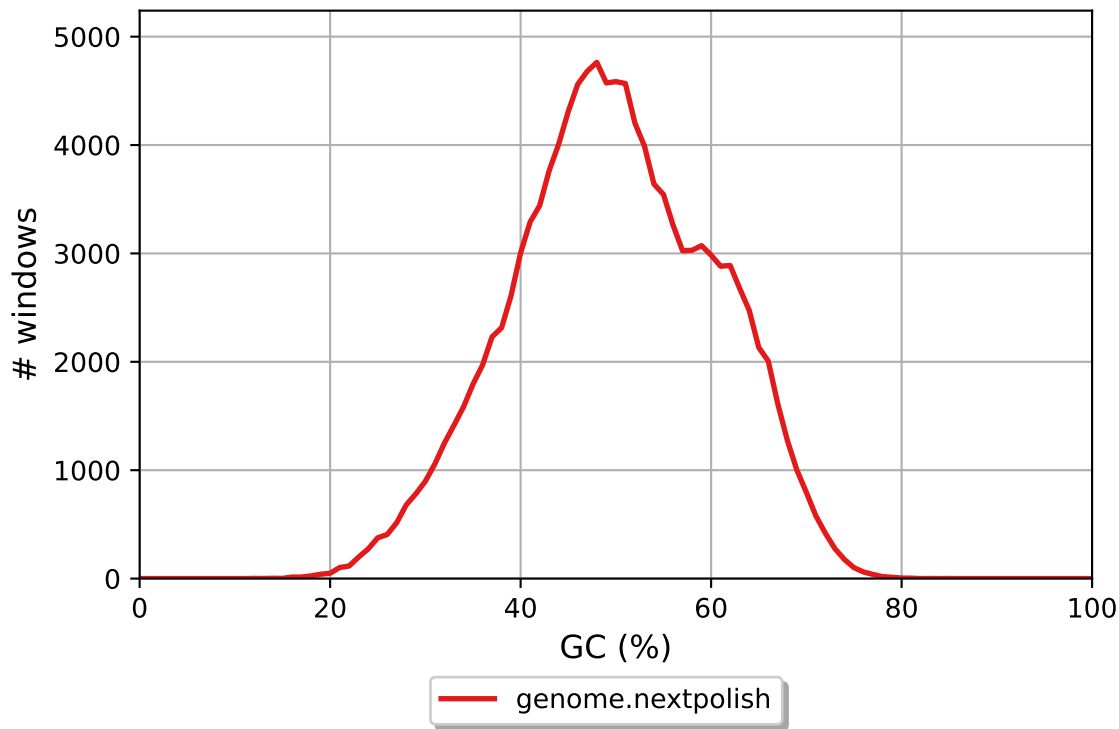


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

