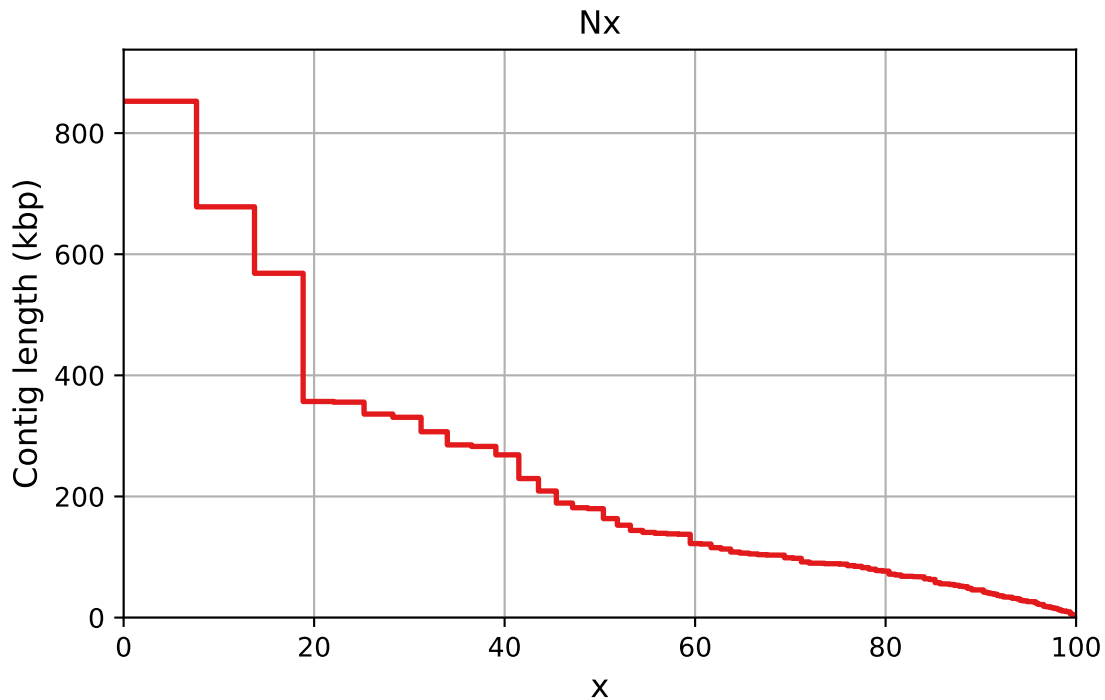


Report

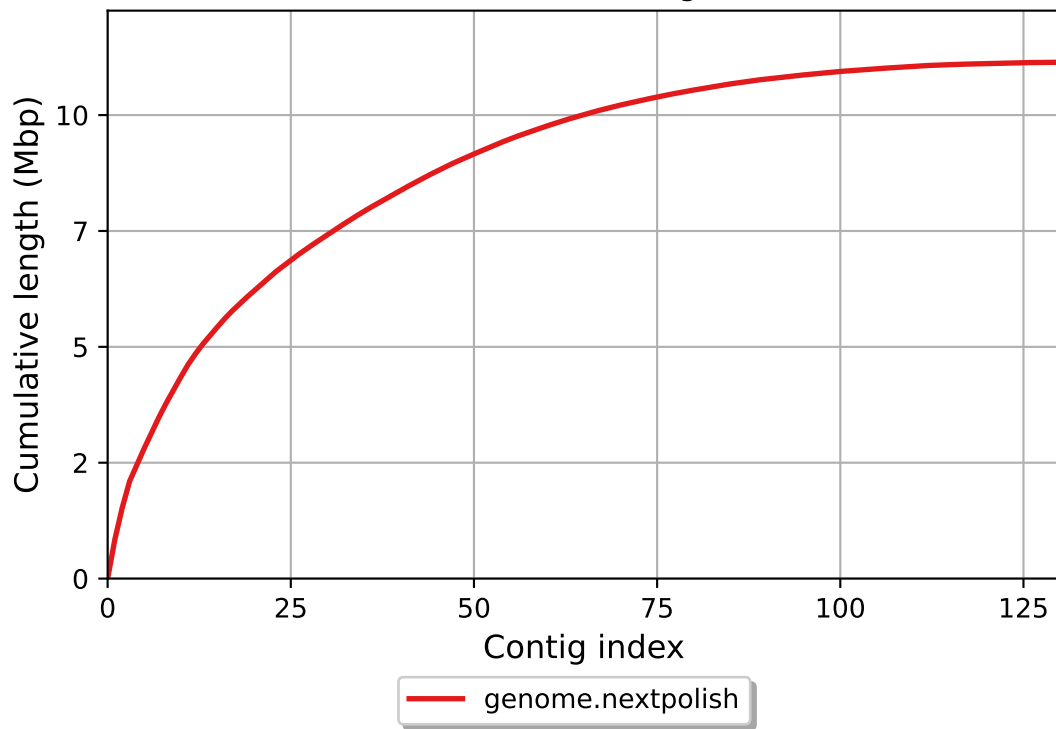
	genome.nextpolish
# contigs (>= 0 bp)	130
# contigs (>= 1000 bp)	128
# contigs (>= 5000 bp)	116
# contigs (>= 10000 bp)	108
# contigs (>= 25000 bp)	85
# contigs (>= 50000 bp)	62
Total length (>= 0 bp)	11139778
Total length (>= 1000 bp)	11138255
Total length (>= 5000 bp)	11094804
Total length (>= 10000 bp)	11032019
Total length (>= 25000 bp)	10673163
Total length (>= 50000 bp)	9870844
# contigs	130
Largest contig	852602
Total length	11139778
GC (%)	53.31
N50	179870
N90	45628
auN	267647.2
L50	16
L90	66
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 1 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).

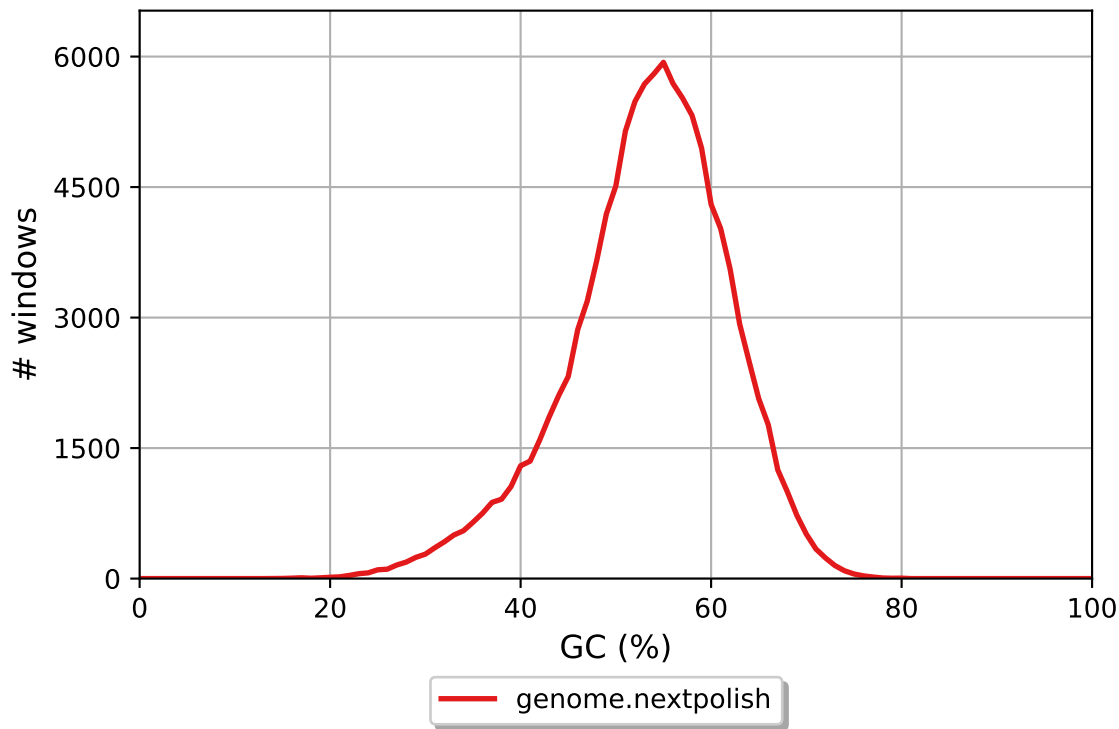


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

