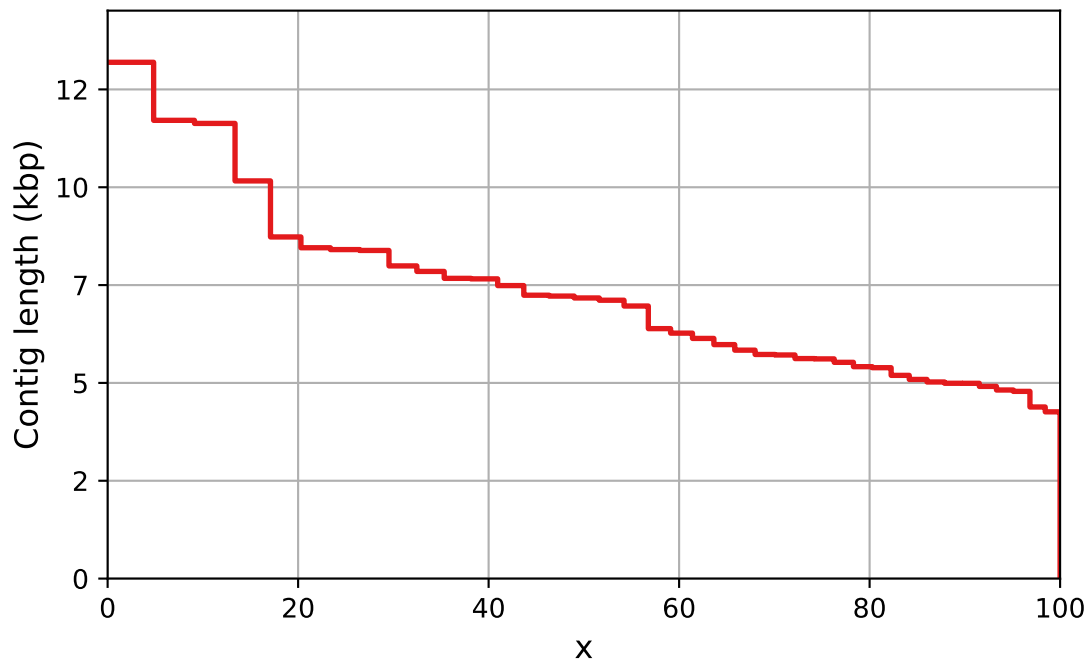


# Report

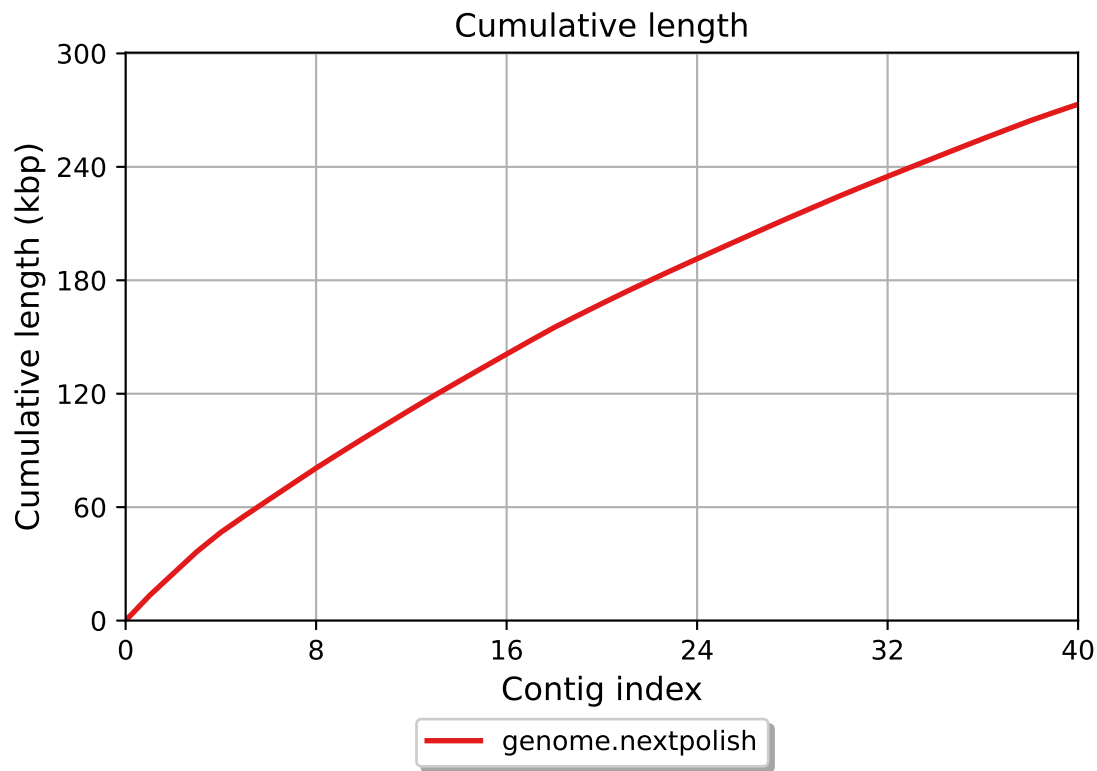
	genome.nextpolish
# contigs (>= 0 bp)	40
# contigs (>= 1000 bp)	40
# contigs (>= 5000 bp)	33
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	273104
Total length (>= 1000 bp)	273104
Total length (>= 5000 bp)	239963
Total length (>= 10000 bp)	46696
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	40
Largest contig	13193
Total length	273104
GC (%)	50.44
N50	7171
N90	4990
auN	7444.4
L50	16
L90	35
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 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).

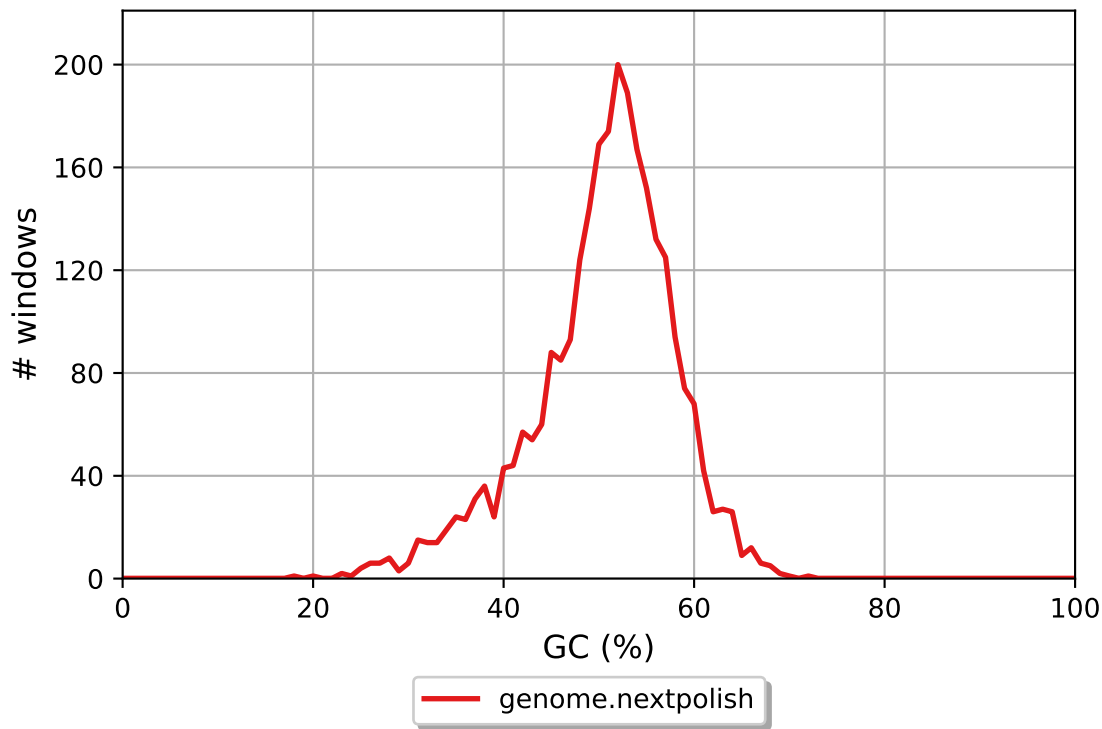
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

