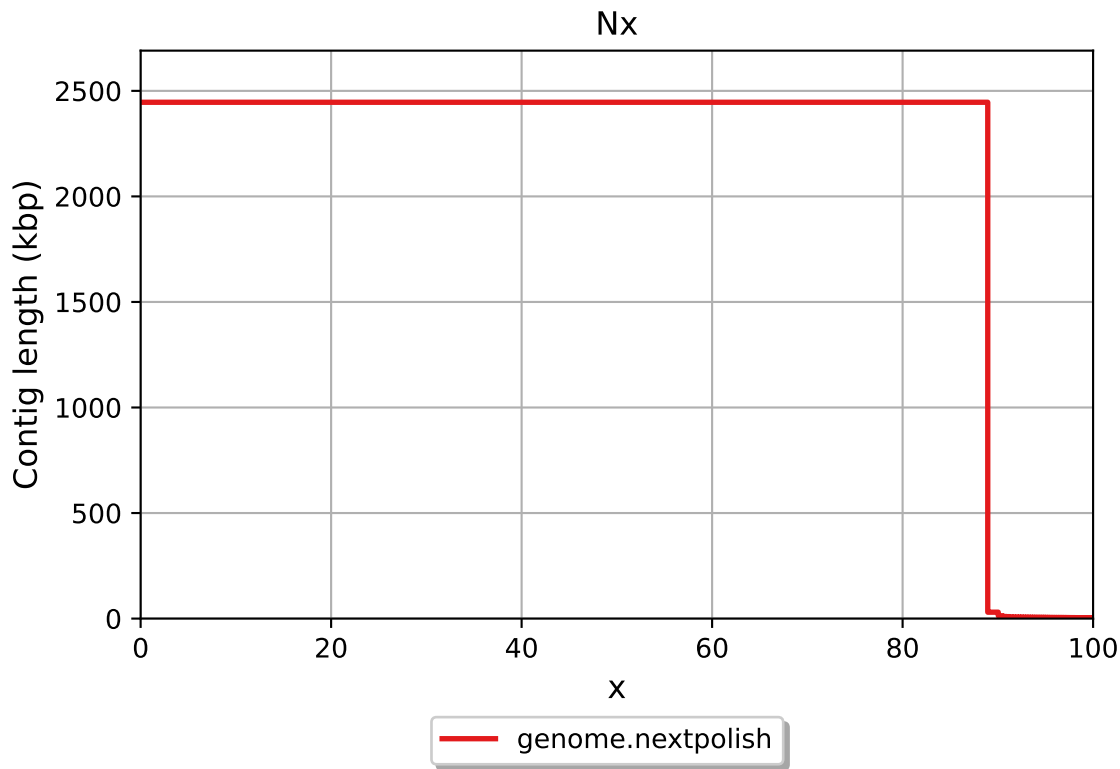


# Report

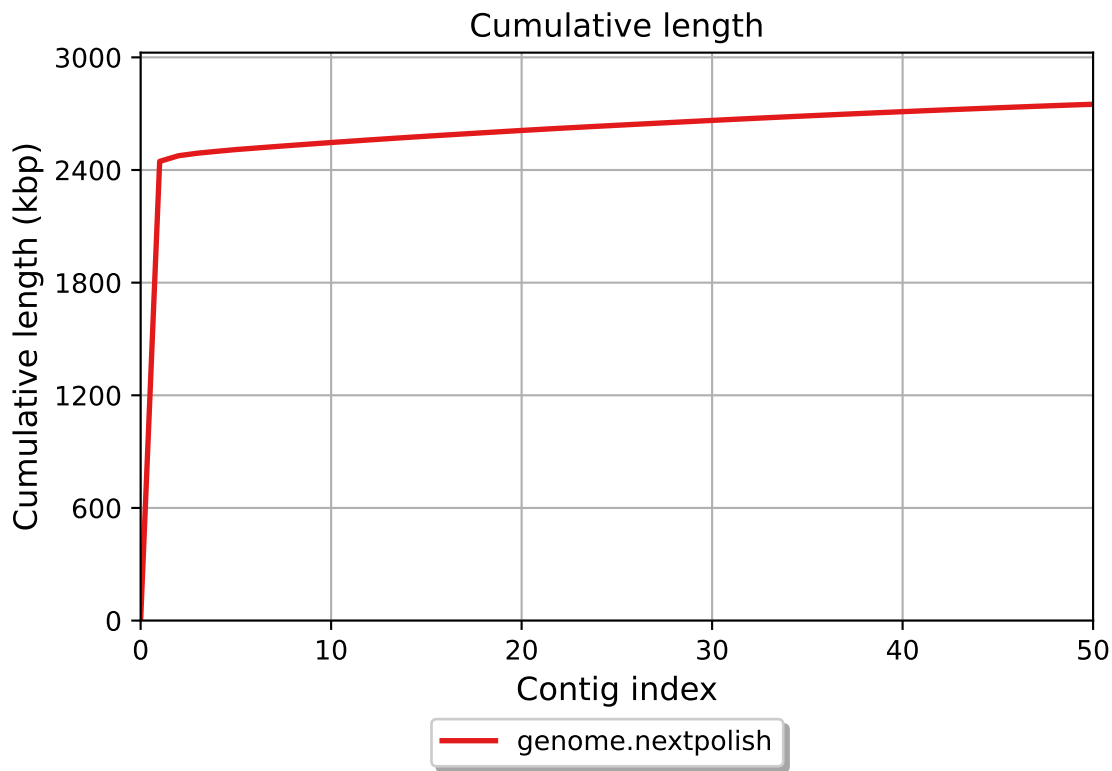
	genome.nextpolish
# contigs (>= 0 bp)	50
# contigs (>= 1000 bp)	50
# contigs (>= 5000 bp)	31
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2750140
Total length (>= 1000 bp)	2750140
Total length (>= 5000 bp)	2669183
Total length (>= 10000 bp)	2489640
Total length (>= 25000 bp)	2476297
Total length (>= 50000 bp)	2445978
# contigs	50
Largest contig	2445978
Total length	2750140
GC (%)	34.12
N50	2445978
N90	30319
auN	2176415.2
L50	1
L90	2
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 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).



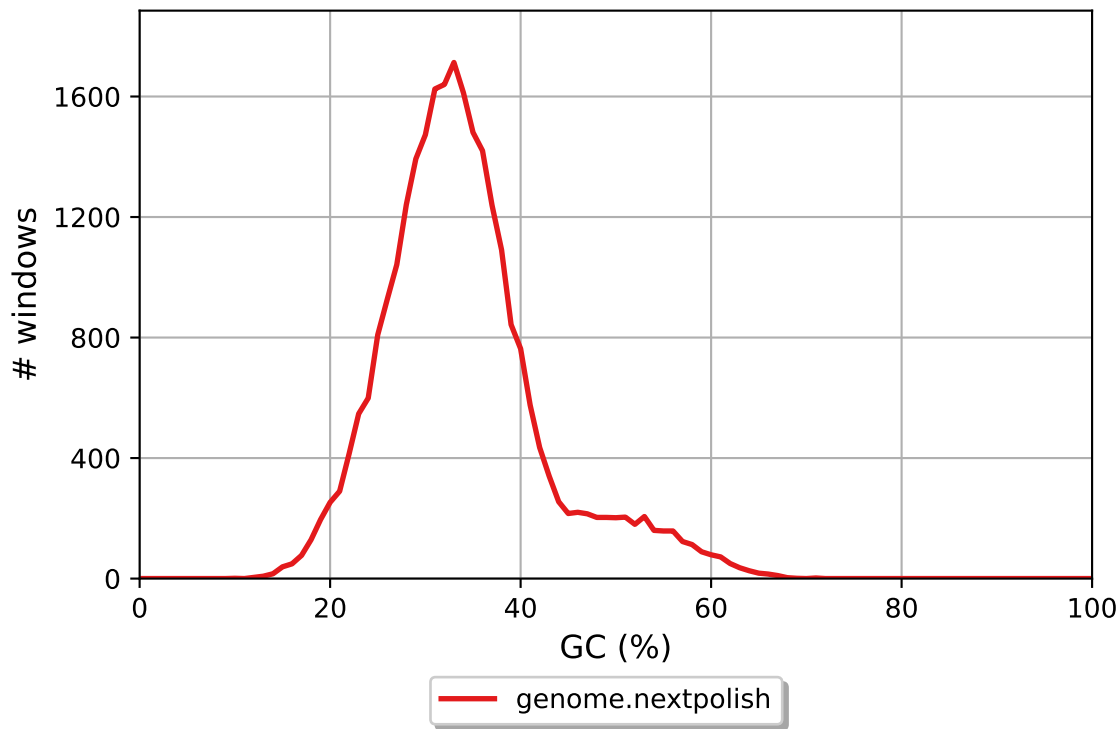








## GC content





genome.nextpolish GC content

