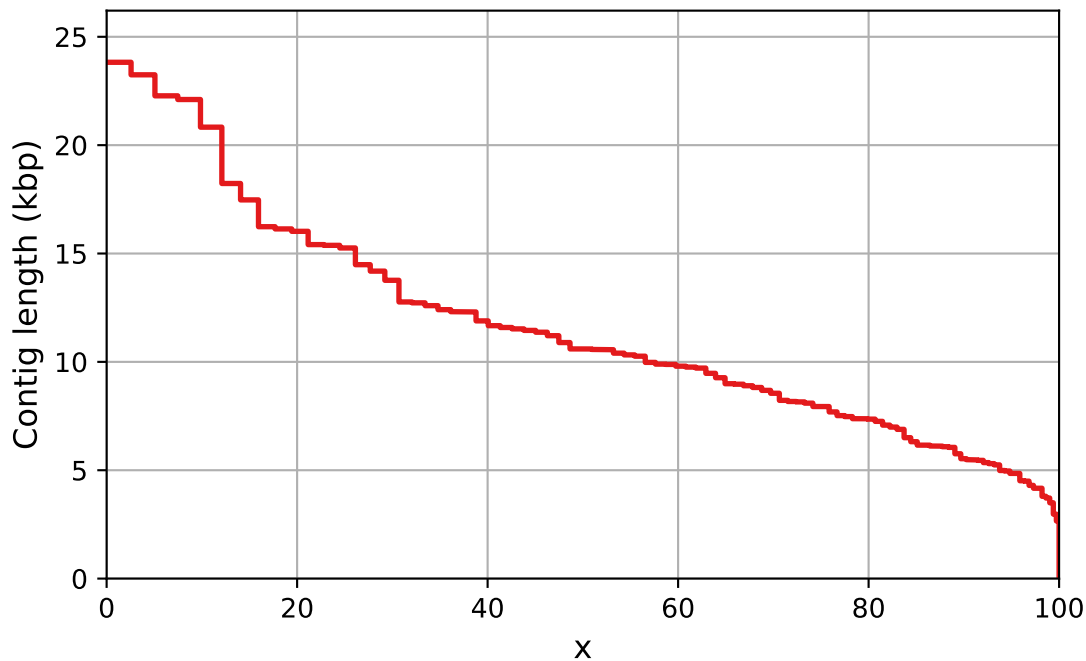


# Report

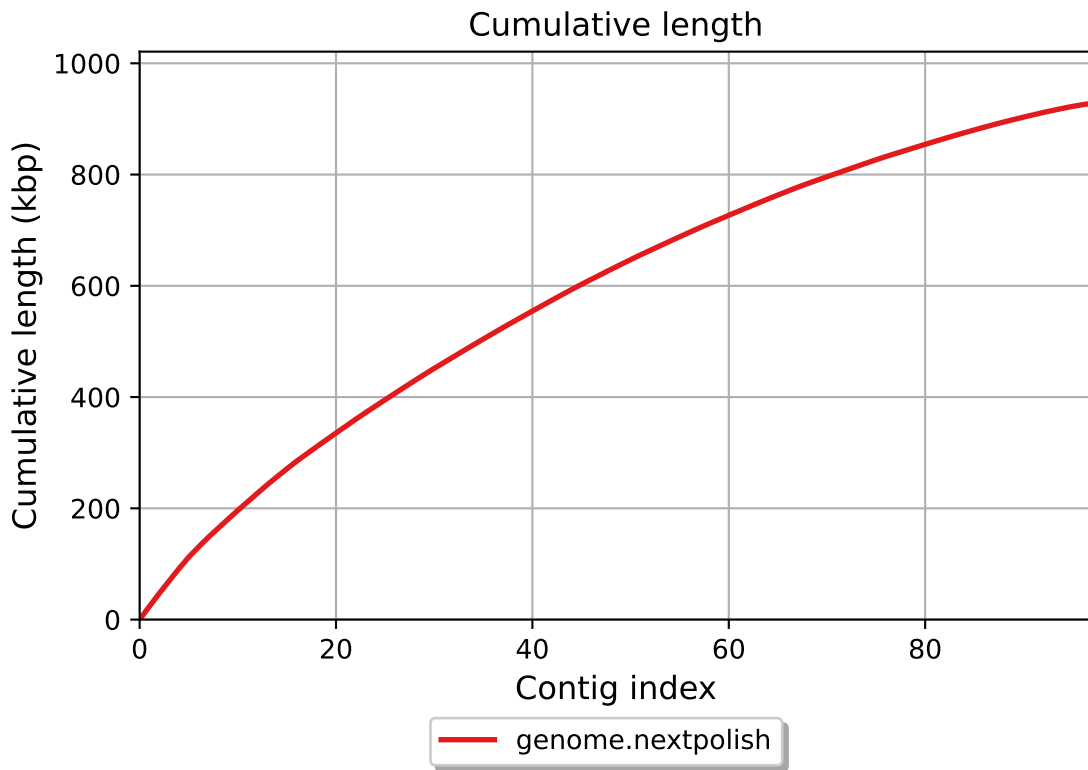
	genome.nextpolish
# contigs (>= 0 bp)	97
# contigs (>= 1000 bp)	97
# contigs (>= 5000 bp)	83
# contigs (>= 10000 bp)	37
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	928152
Total length (>= 1000 bp)	928152
Total length (>= 5000 bp)	870203
Total length (>= 10000 bp)	524875
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	97
Largest contig	23824
Total length	928152
GC (%)	54.59
N50	10595
N90	5532
auN	11798.2
L50	32
L90	77
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

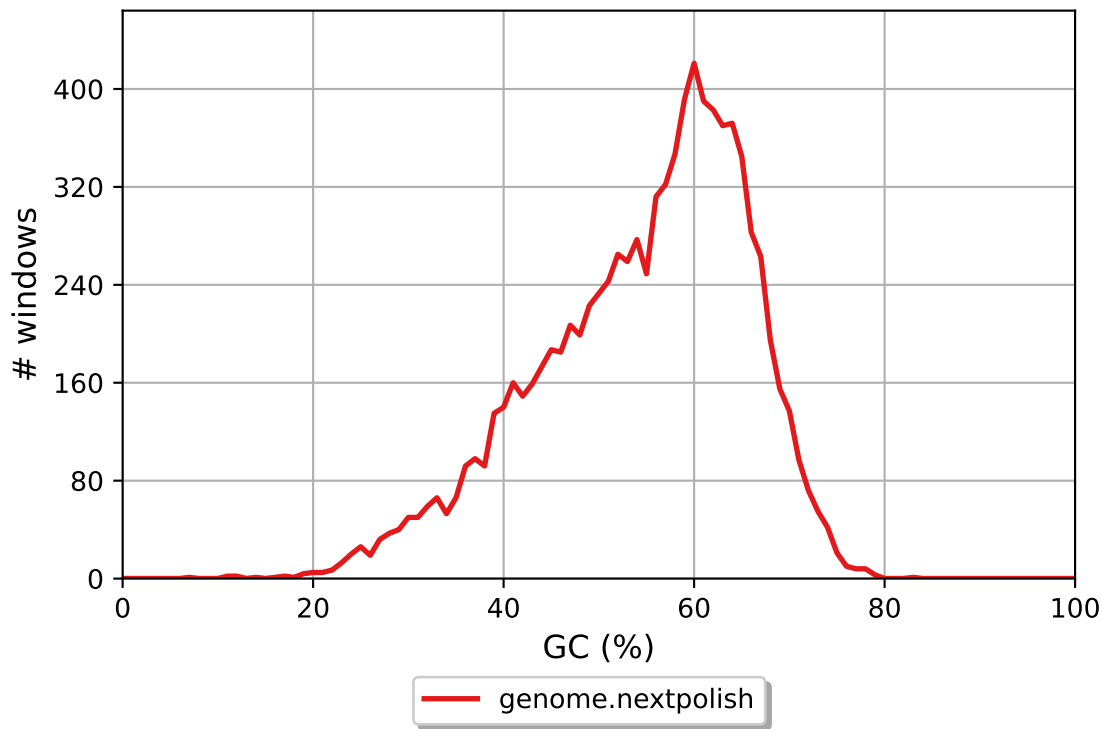
Nx



genome.nextpolish



## GC content



genome.nextpolish GC content

