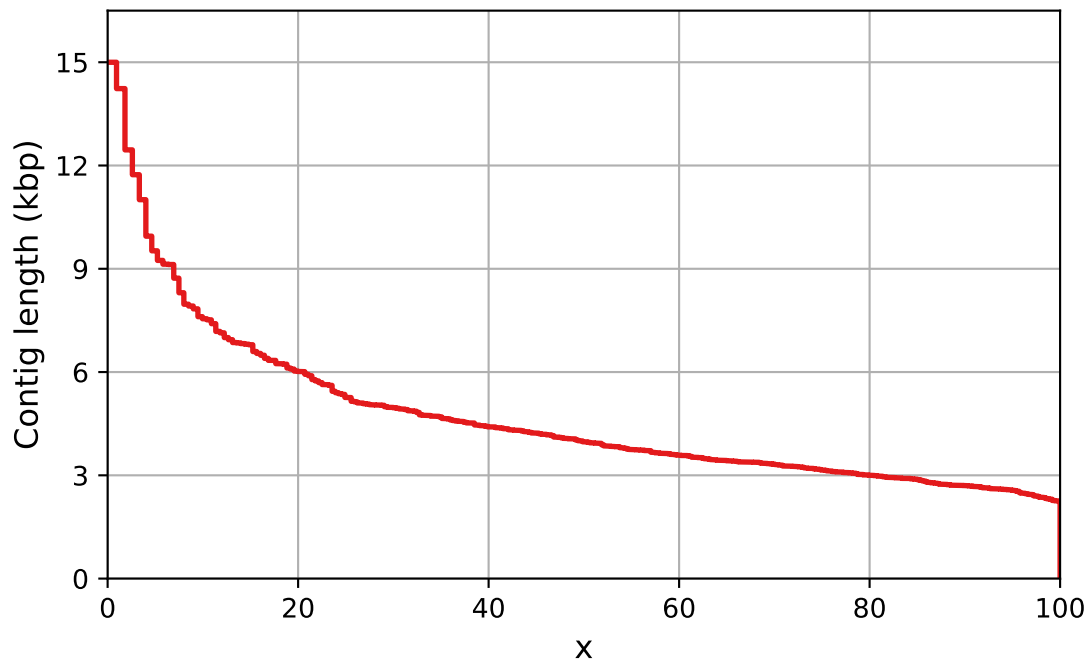


# Report

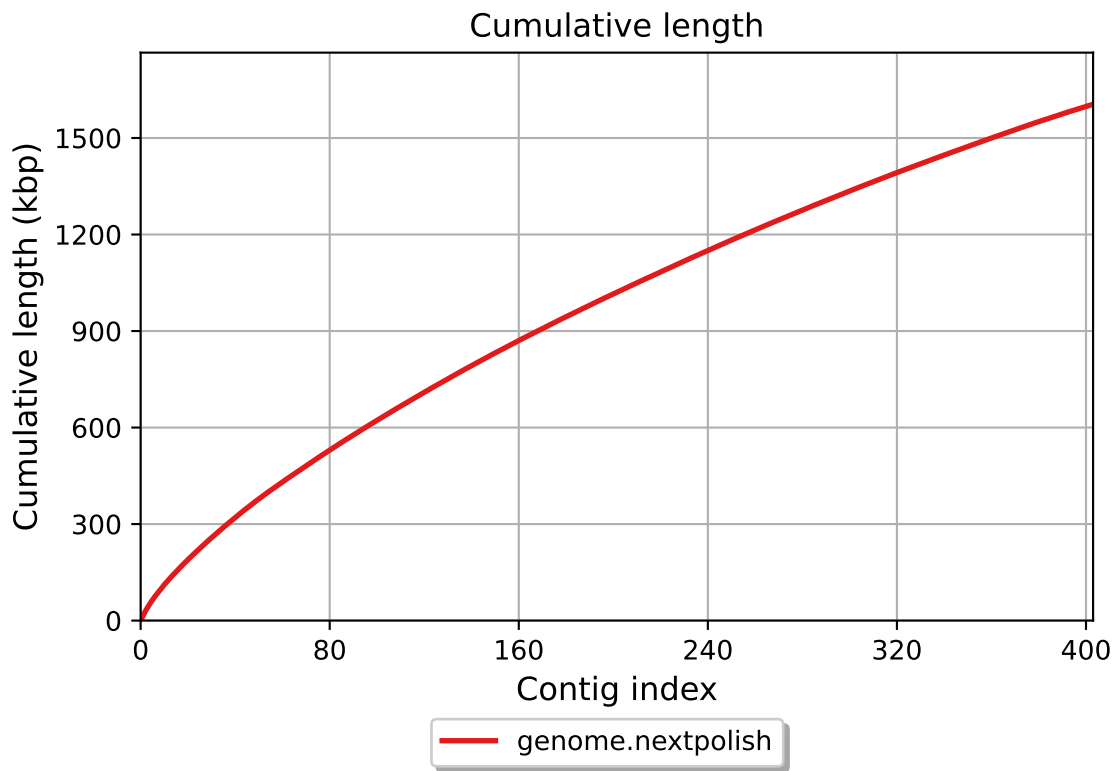
	genome.nextpolish
# contigs (>= 0 bp)	403
# contigs (>= 1000 bp)	403
# contigs (>= 5000 bp)	67
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1604850
Total length (>= 1000 bp)	1604850
Total length (>= 5000 bp)	466154
Total length (>= 10000 bp)	64427
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	403
Largest contig	14999
Total length	1604850
GC (%)	56.26
N50	3974
N90	2699
auN	4718.5
L50	143
L90	340
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

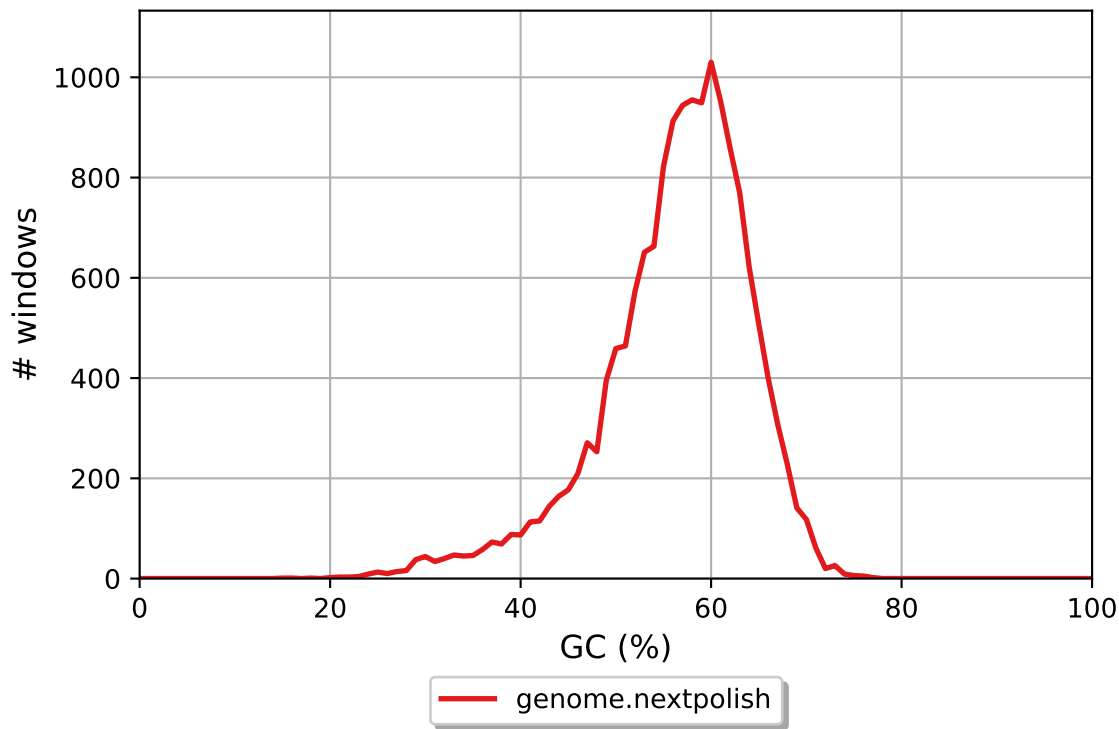
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

