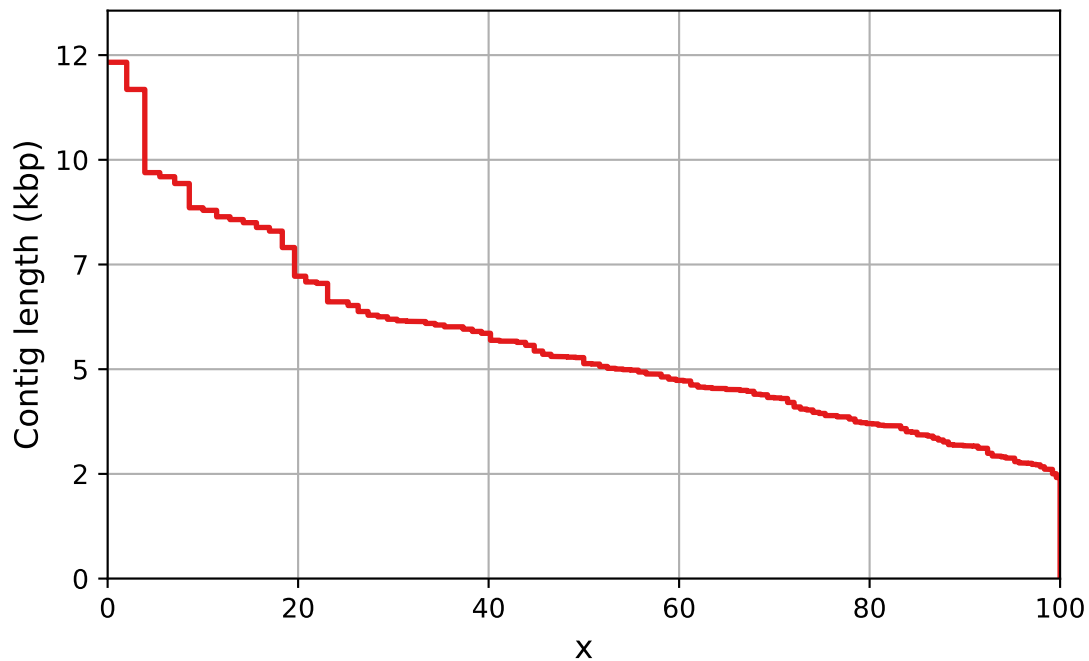


# Report

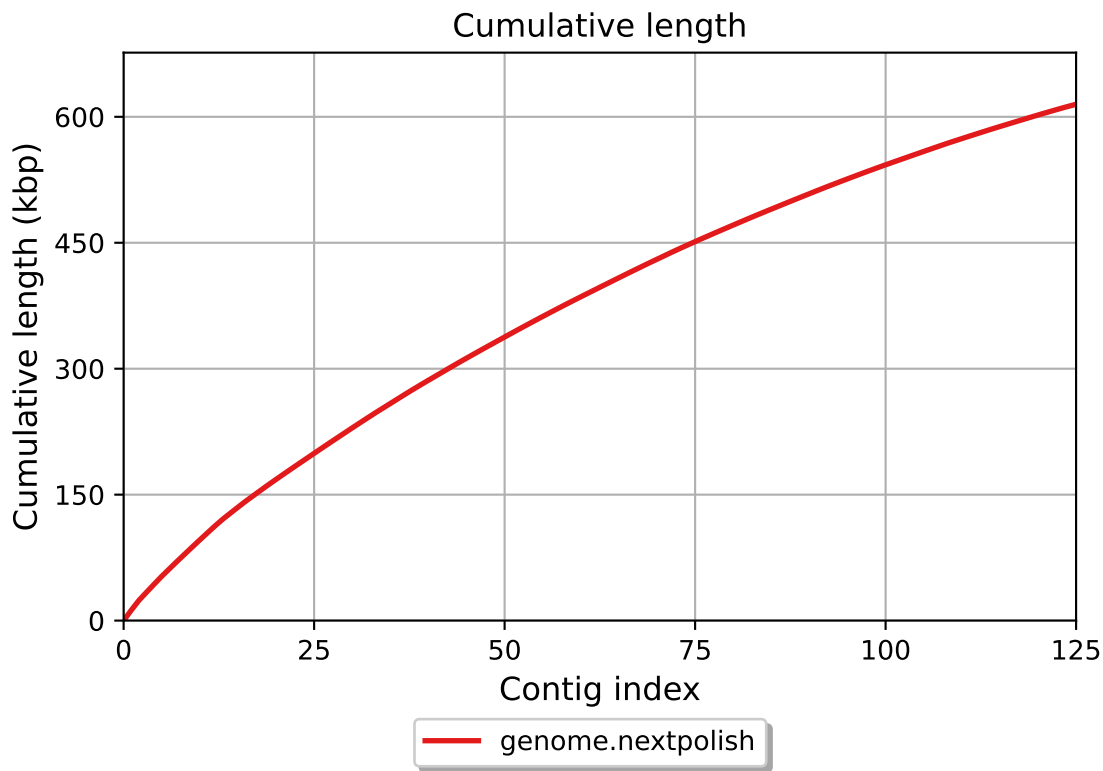
	genome.nextpolish
# contigs (>= 0 bp)	125
# contigs (>= 1000 bp)	125
# contigs (>= 5000 bp)	49
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	614945
Total length (>= 1000 bp)	614945
Total length (>= 5000 bp)	332740
Total length (>= 10000 bp)	24009
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	125
Largest contig	12329
Total length	614945
GC (%)	58.01
N50	5133
N90	3172
auN	5676.8
L50	45
L90	104
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

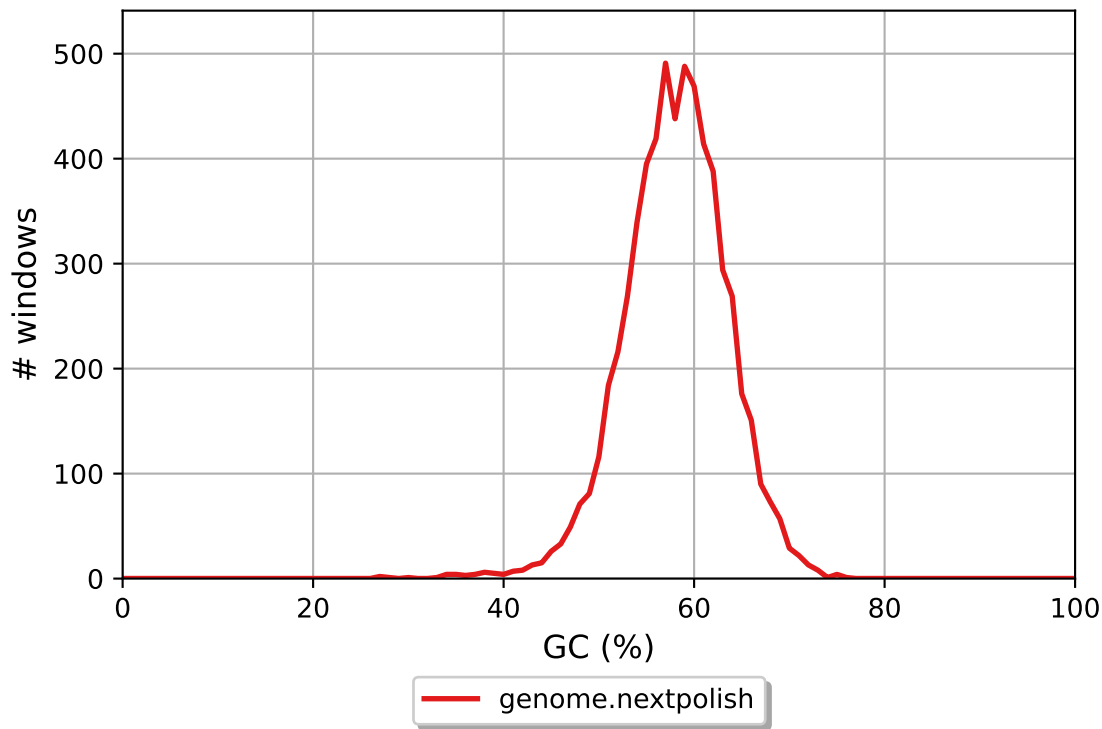
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

