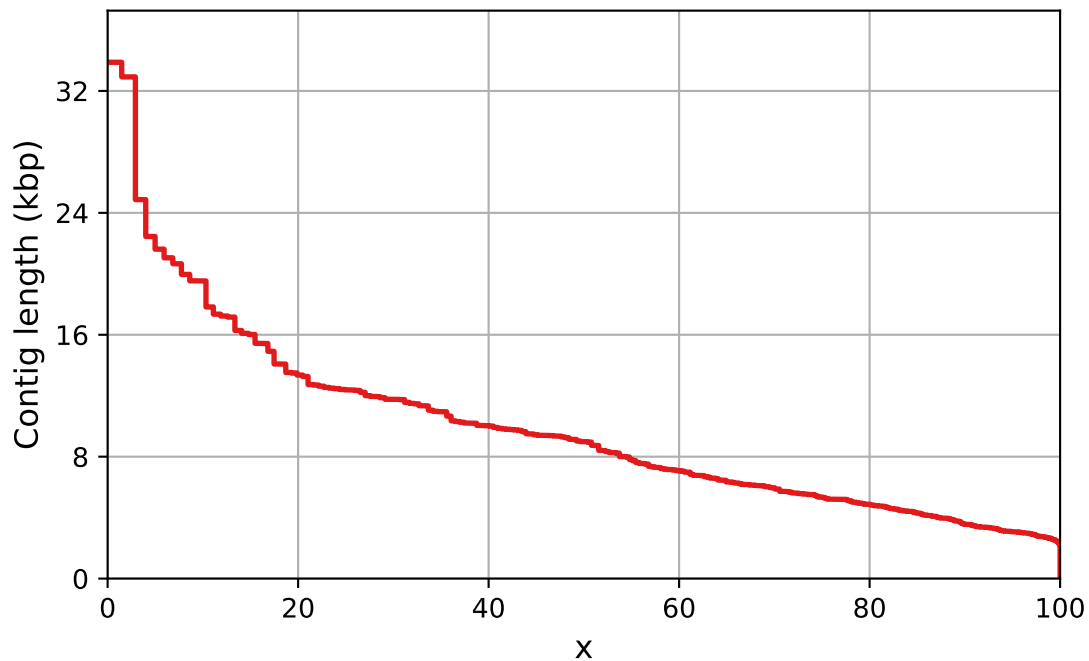


# Report

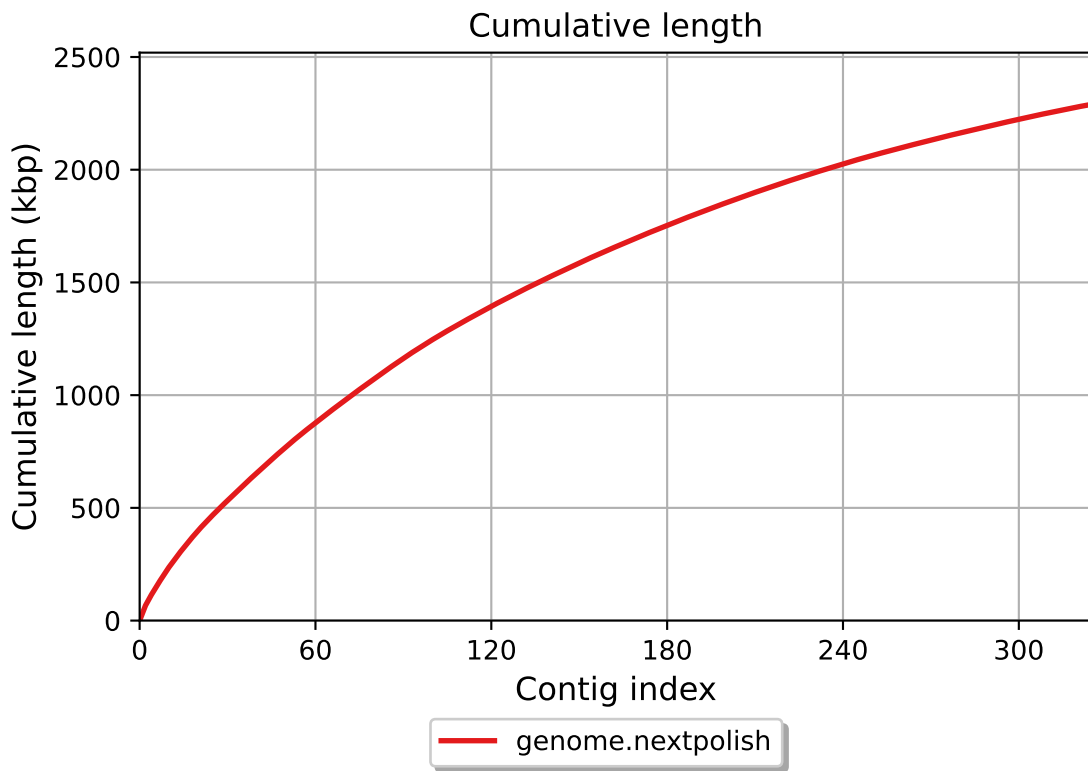
	genome.nextpolish
# contigs (>= 0 bp)	325
# contigs (>= 1000 bp)	325
# contigs (>= 5000 bp)	189
# contigs (>= 10000 bp)	65
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2290309
Total length (>= 1000 bp)	2290309
Total length (>= 5000 bp)	1799152
Total length (>= 10000 bp)	927582
Total length (>= 25000 bp)	66802
Total length (>= 50000 bp)	0
# contigs	325
Largest contig	33881
Total length	2290309
GC (%)	56.65
N50	8991
N90	3566
auN	10057.2
L50	88
L90	250
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

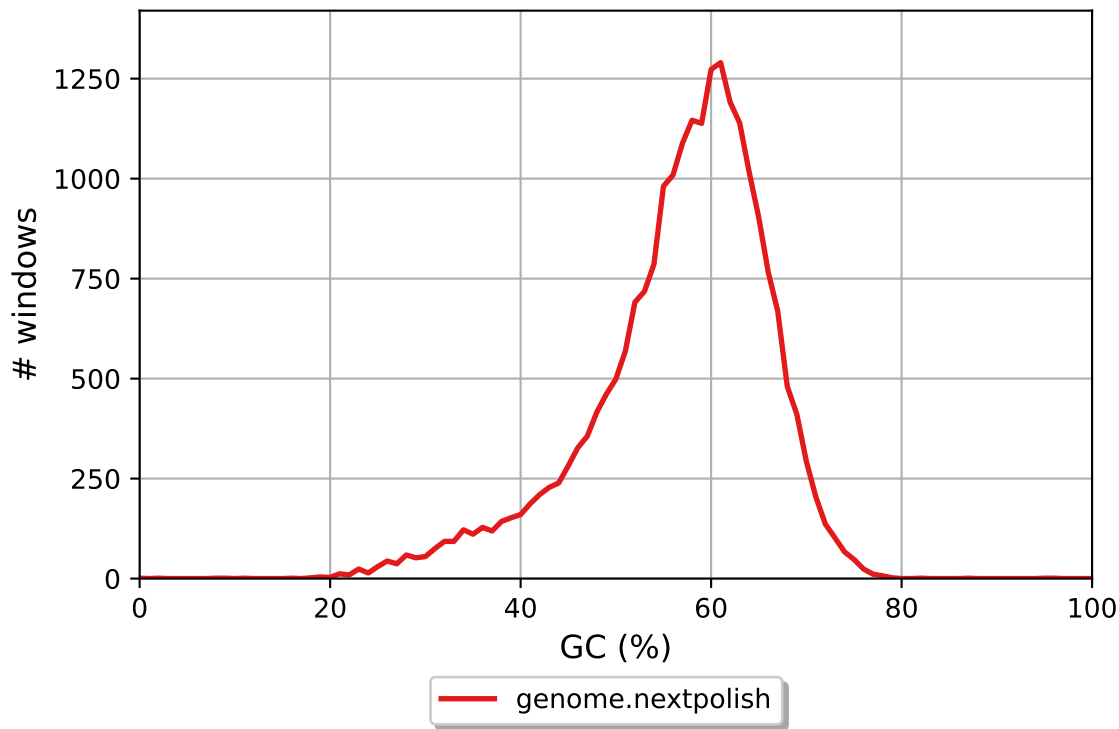
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

