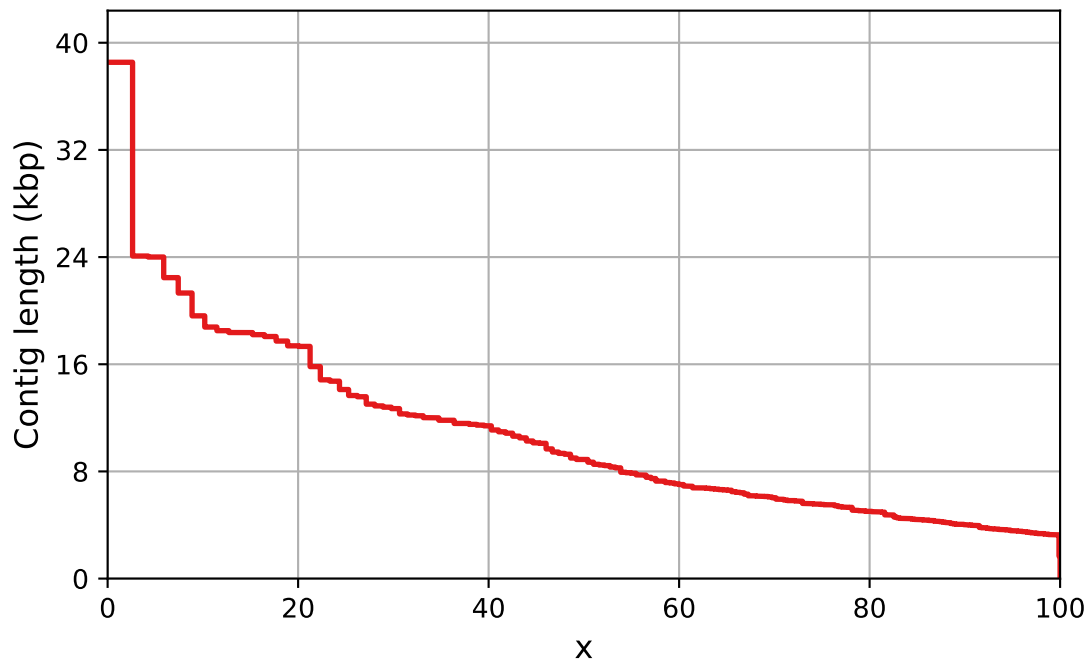


# Report

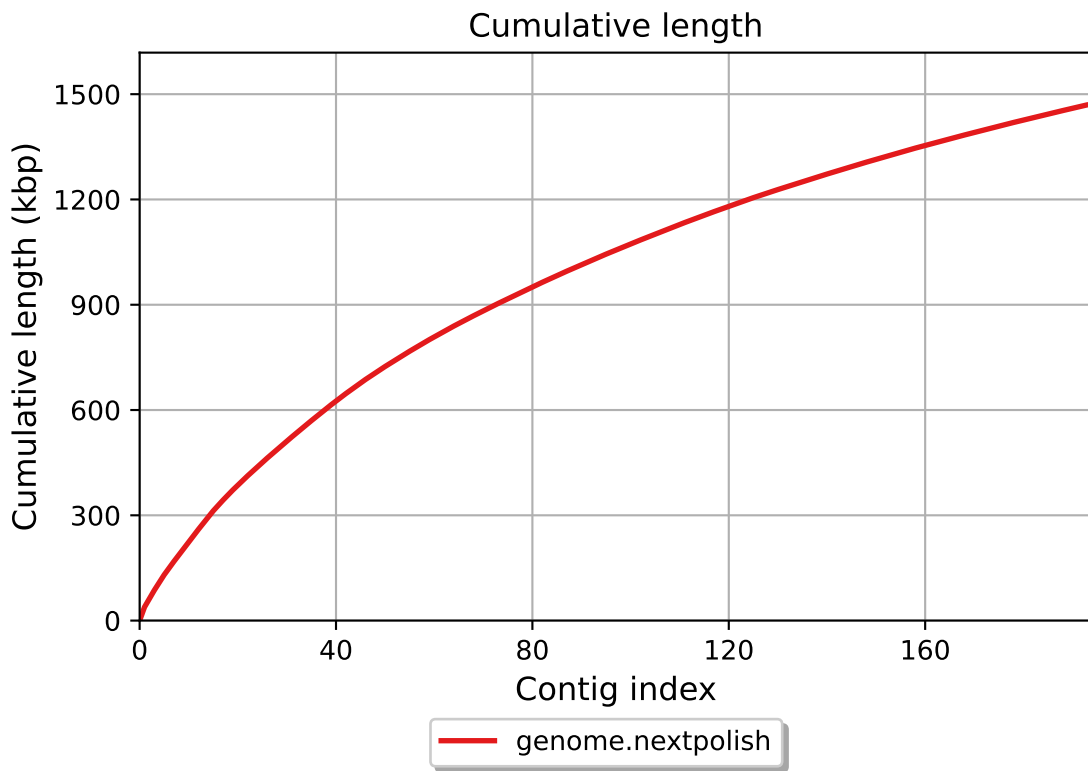
	genome.nextpolish
# contigs (>= 0 bp)	194
# contigs (>= 1000 bp)	194
# contigs (>= 5000 bp)	120
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1471234
Total length (>= 1000 bp)	1471234
Total length (>= 5000 bp)	1180361
Total length (>= 10000 bp)	677273
Total length (>= 25000 bp)	38542
Total length (>= 50000 bp)	0
# contigs	194
Largest contig	38542
Total length	1471234
GC (%)	56.89
N50	8890
N90	4028
auN	10928.4
L50	52
L90	153
# 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).

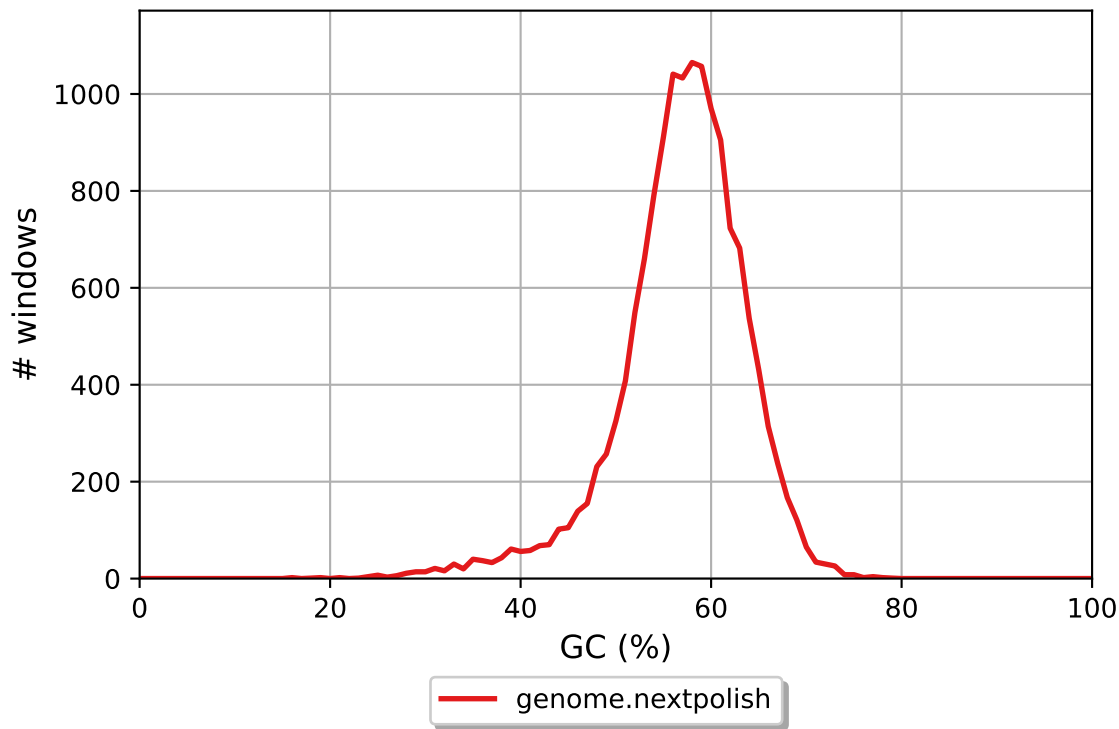
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

