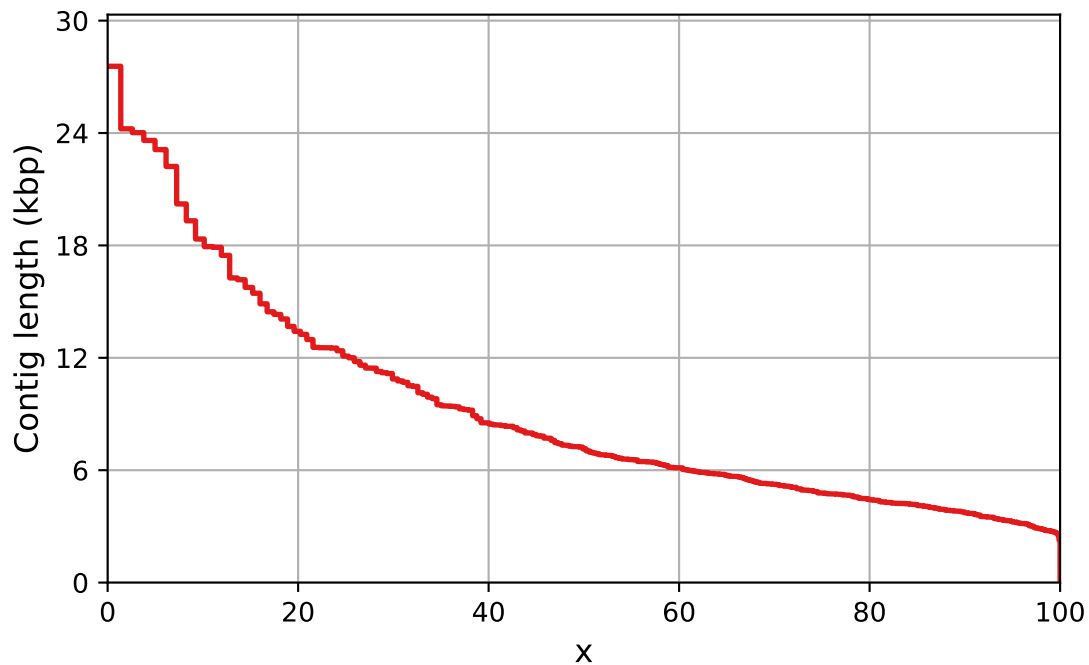


Report

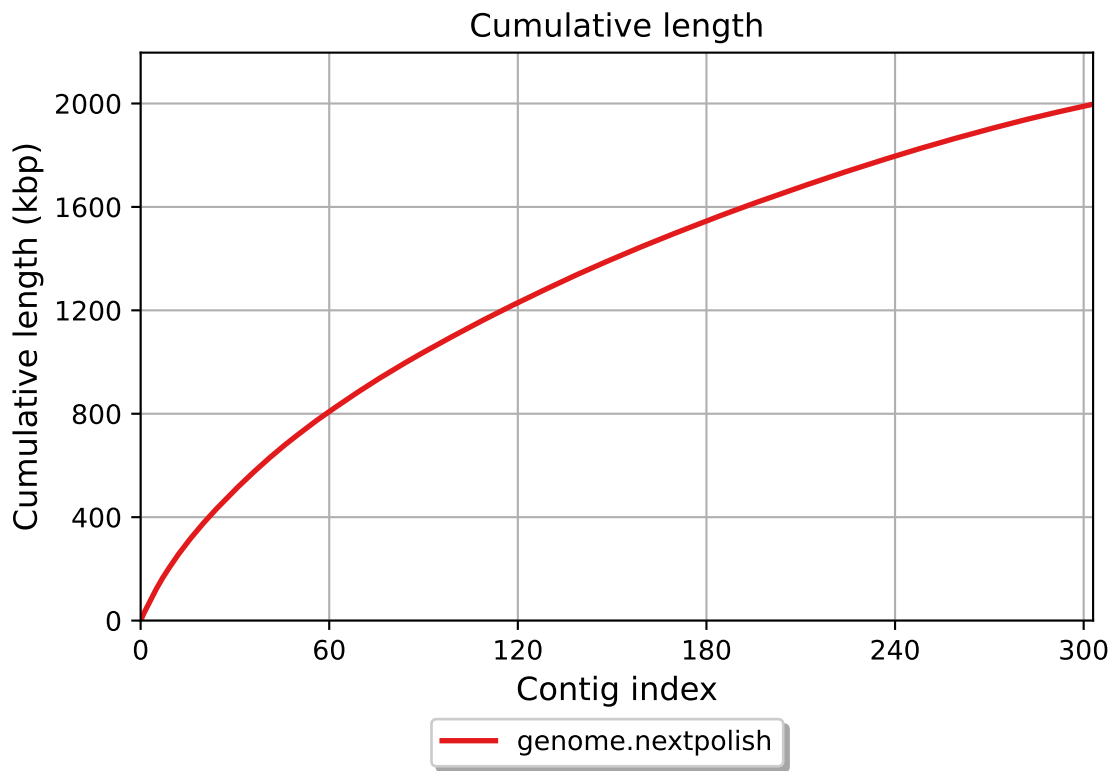
	genome.nextpolish
# contigs (>= 0 bp)	303
# contigs (>= 1000 bp)	303
# contigs (>= 5000 bp)	161
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1997023
Total length (>= 1000 bp)	1997023
Total length (>= 5000 bp)	1454375
Total length (>= 10000 bp)	670642
Total length (>= 25000 bp)	27563
Total length (>= 50000 bp)	0
# contigs	303
Largest contig	27563
Total length	1997023
GC (%)	56.05
N50	7143
N90	3730
auN	9273.4
L50	85
L90	241
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

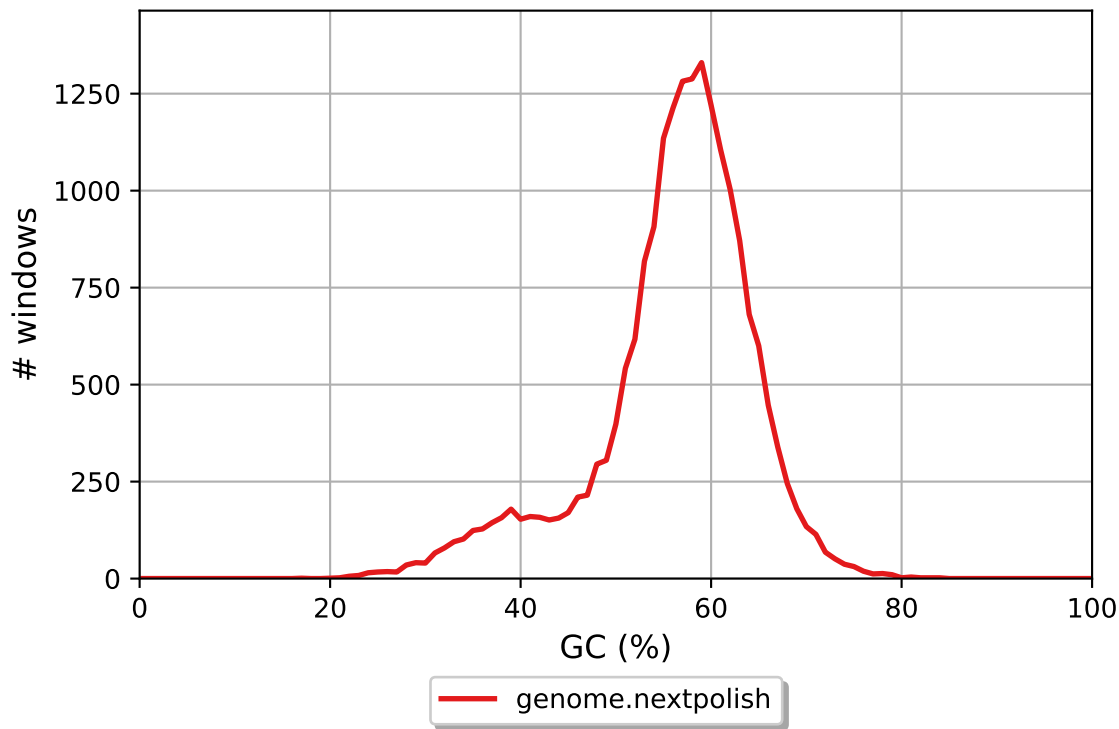
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

