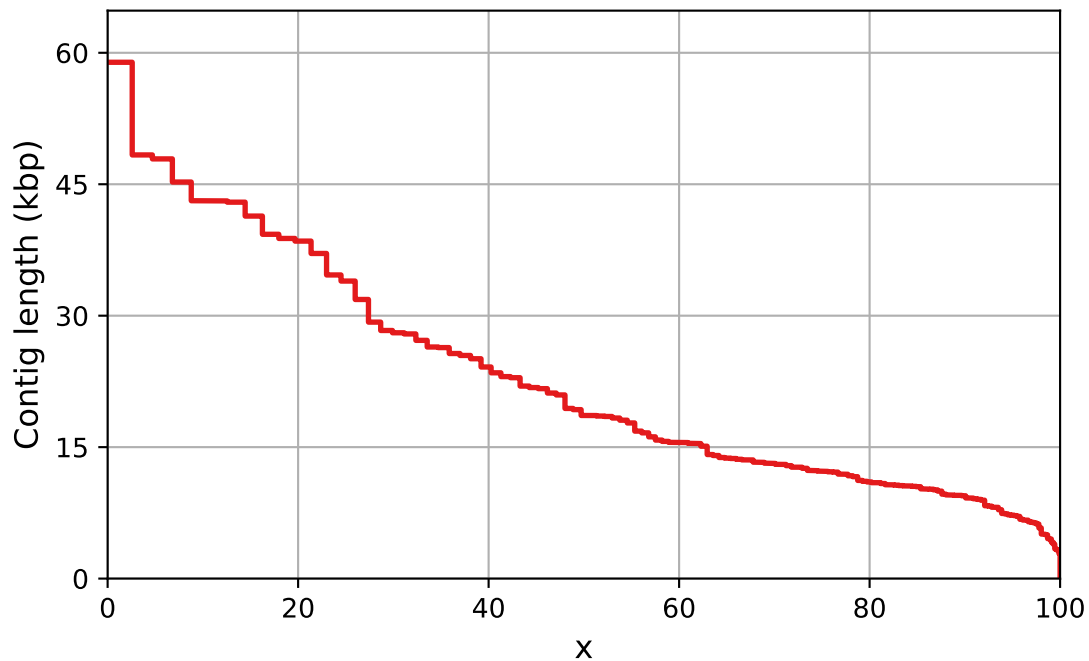


Report

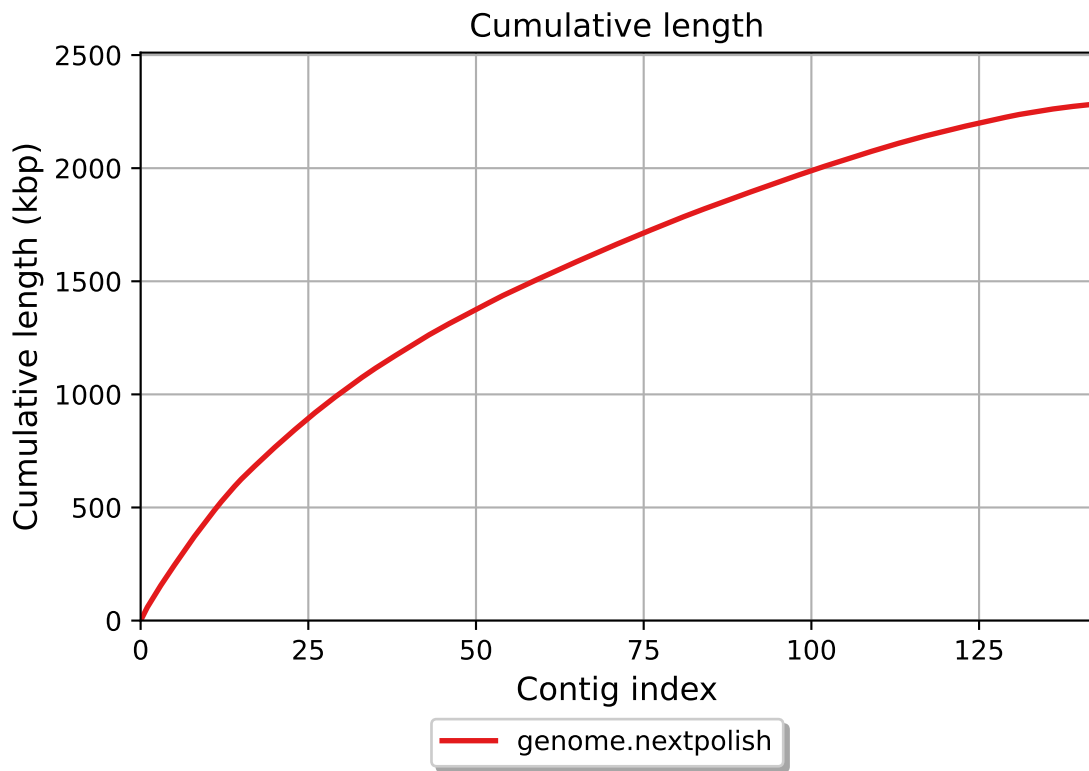
	genome.nextpolish
# contigs (>= 0 bp)	142
# contigs (>= 1000 bp)	142
# contigs (>= 5000 bp)	133
# contigs (>= 10000 bp)	100
# contigs (>= 25000 bp)	25
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2282513
Total length (>= 1000 bp)	2282513
Total length (>= 5000 bp)	2247550
Total length (>= 10000 bp)	1989109
Total length (>= 25000 bp)	894848
Total length (>= 50000 bp)	58918
# contigs	142
Largest contig	58918
Total length	2282513
GC (%)	56.32
N50	18613
N90	9450
auN	23342.6
L50	37
L90	107
# 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).

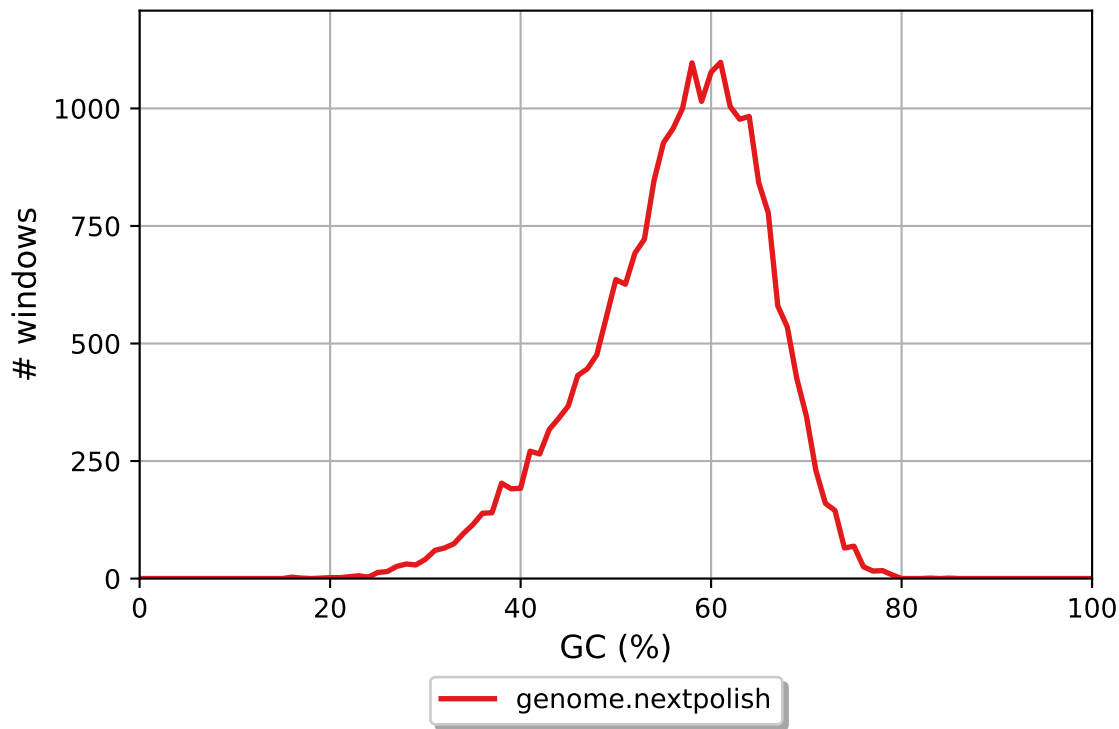
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

