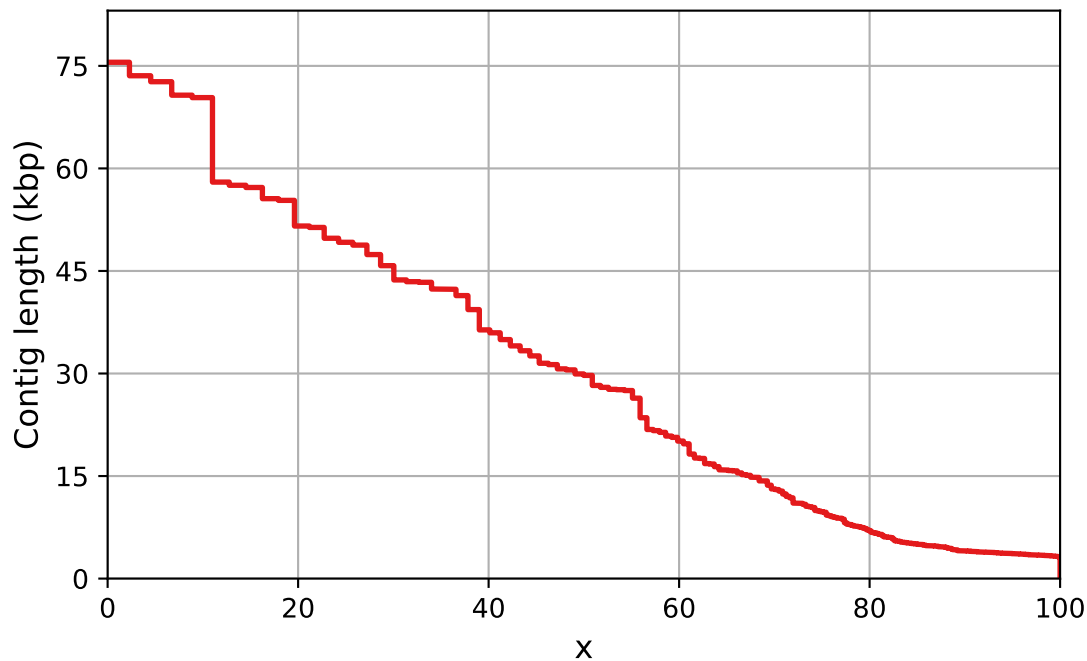


Report

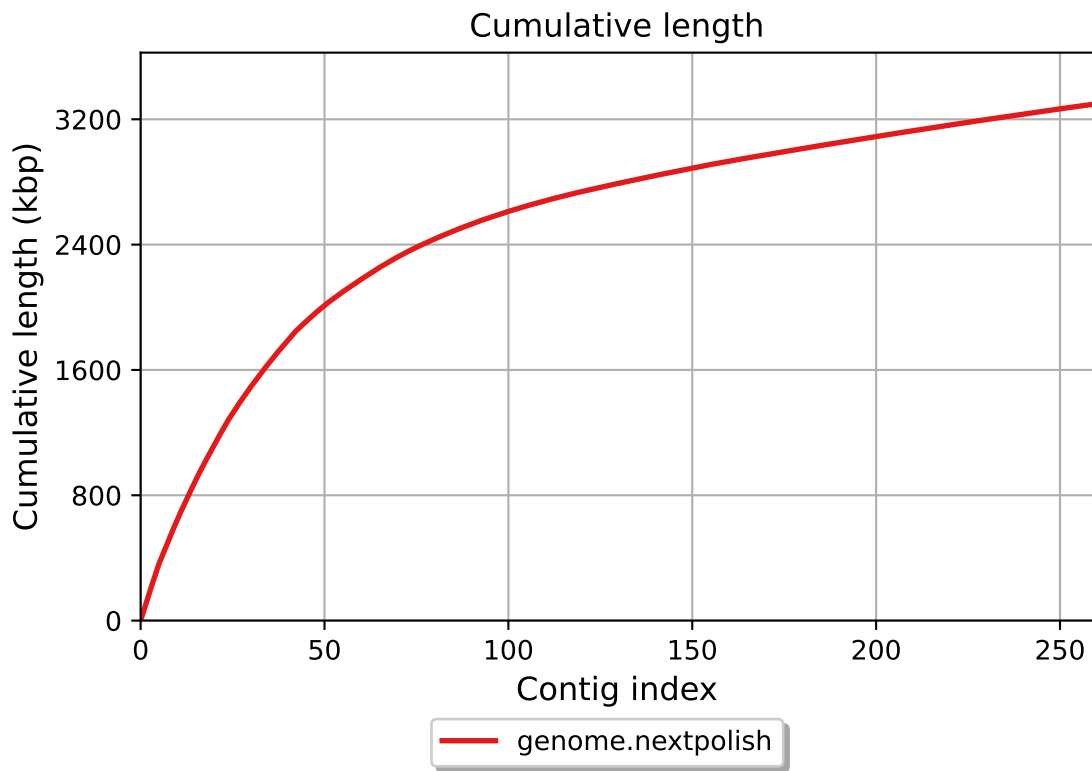
	genome.nextpolish
# contigs (>= 0 bp)	259
# contigs (>= 1000 bp)	259
# contigs (>= 5000 bp)	134
# contigs (>= 10000 bp)	81
# contigs (>= 25000 bp)	42
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	3295933
Total length (>= 1000 bp)	3295933
Total length (>= 5000 bp)	2811622
Total length (>= 10000 bp)	2447193
Total length (>= 25000 bp)	1842589
Total length (>= 50000 bp)	749427
# contigs	259
Largest contig	75528
Total length	3295933
GC (%)	55.83
N50	29724
N90	4048
auN	31451.2
L50	36
L90	169
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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).

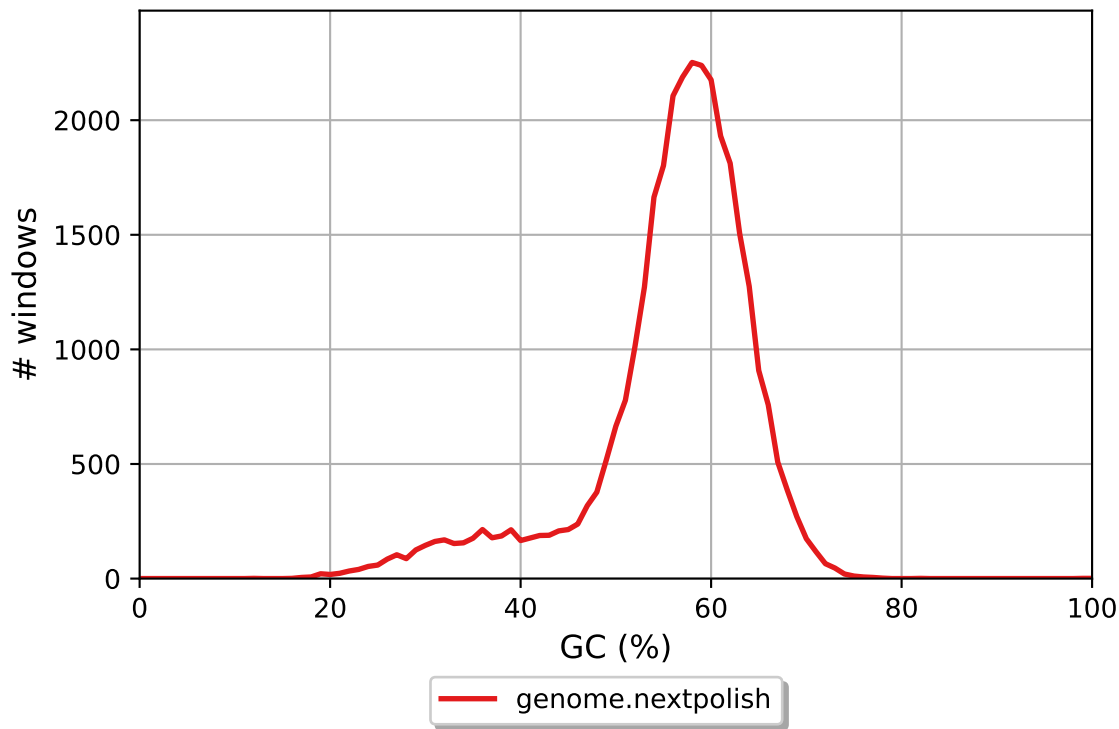
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

