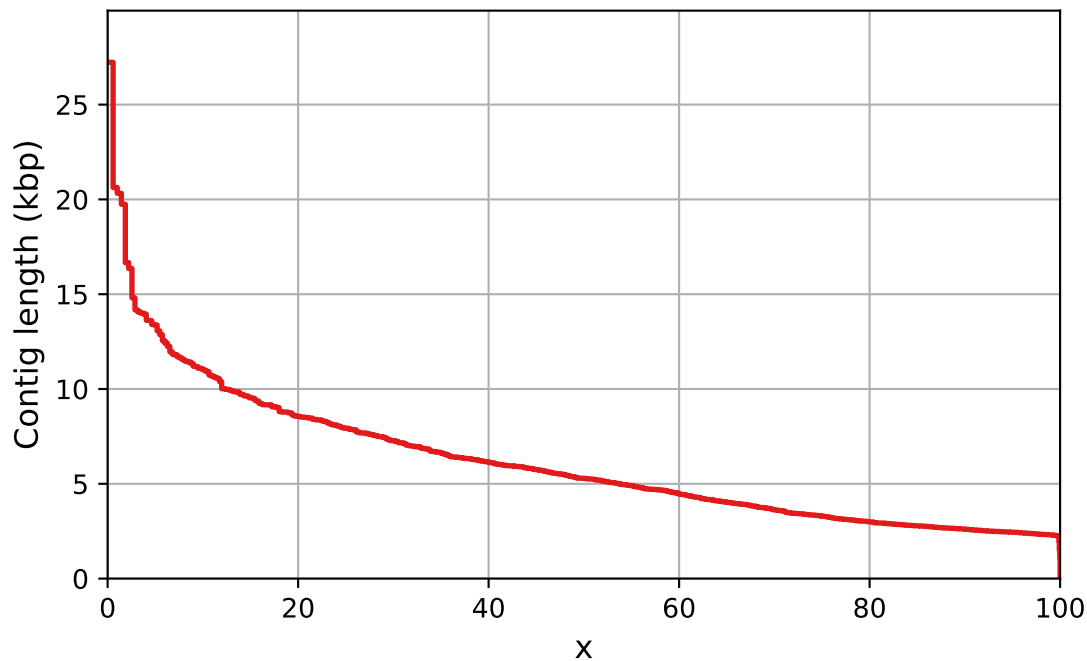


Report

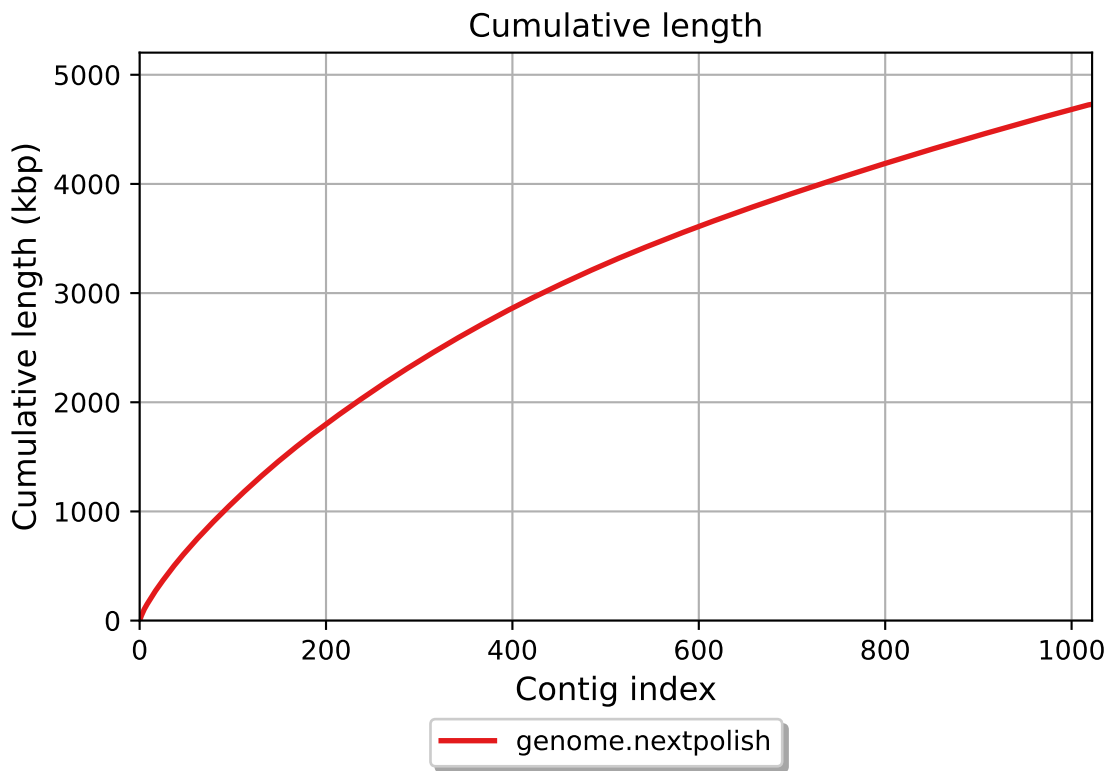
	genome.nextpolish
# contigs (>= 0 bp)	1022
# contigs (>= 1000 bp)	1022
# contigs (>= 5000 bp)	332
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4729776
Total length (>= 1000 bp)	4729776
Total length (>= 5000 bp)	2542500
Total length (>= 10000 bp)	586392
Total length (>= 25000 bp)	27233
Total length (>= 50000 bp)	0
# contigs	1022
Largest contig	27233
Total length	4729776
GC (%)	65.00
N50	5286
N90	2607
auN	6201.8
L50	298
L90	827
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

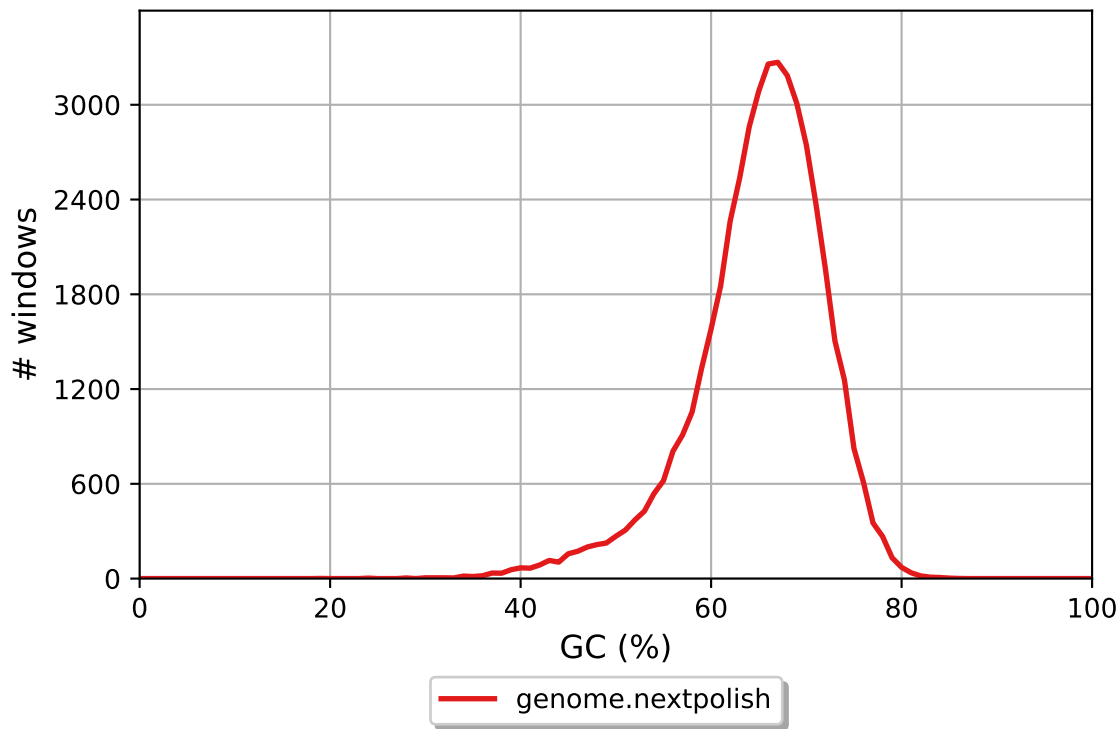
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

