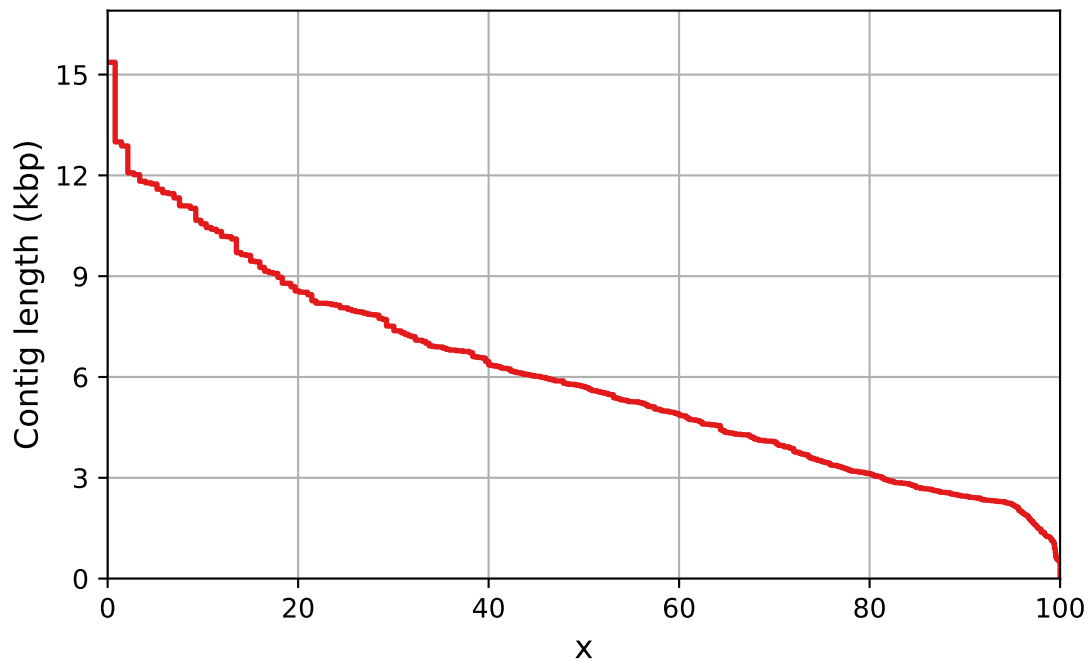


Report

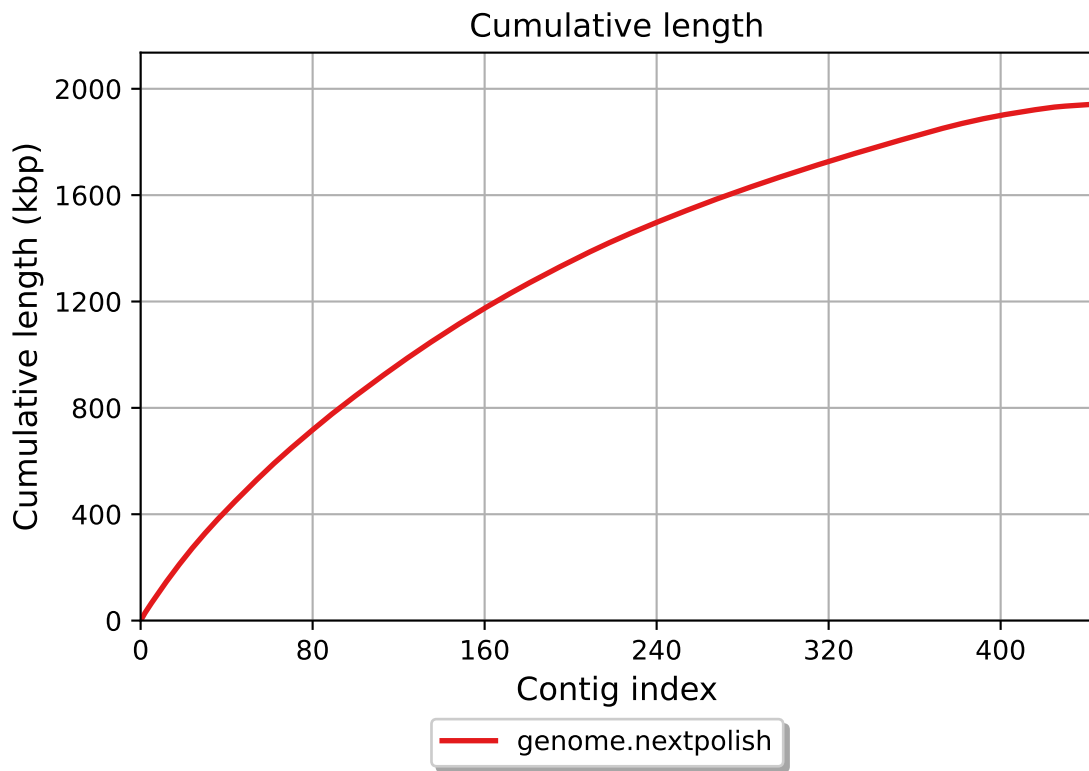
	genome.nextpolish
# contigs (>= 0 bp)	444
# contigs (>= 1000 bp)	424
# contigs (>= 5000 bp)	151
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1942063
Total length (>= 1000 bp)	1930257
Total length (>= 5000 bp)	1130298
Total length (>= 10000 bp)	262609
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	443
Largest contig	15366
Total length	1941795
GC (%)	57.81
N50	5708
N90	2452
auN	6047.0
L50	122
L90	329
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 3 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).

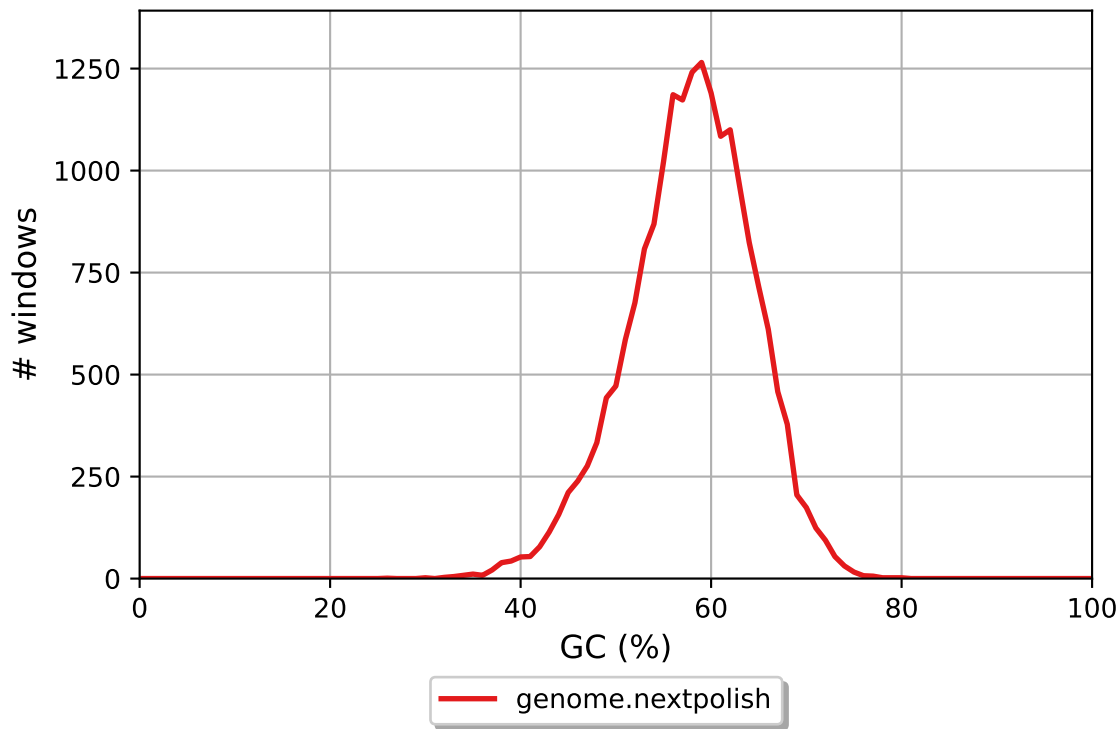
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

