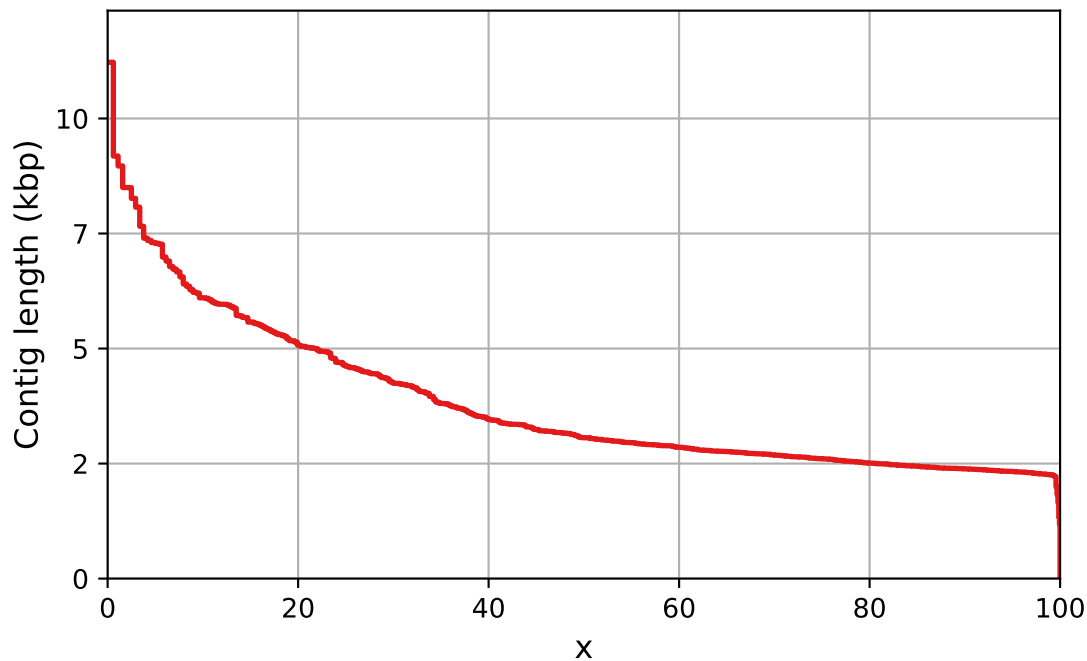


Report

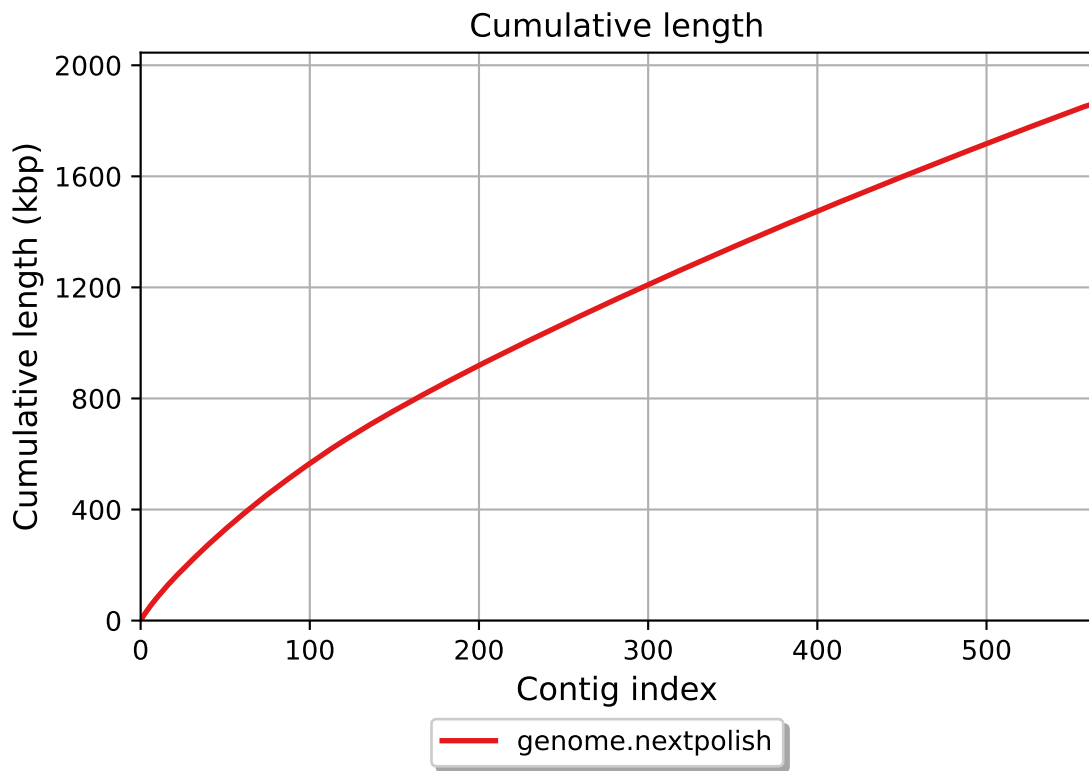
	genome.nextpolish
# contigs (>= 0 bp)	563
# contigs (>= 1000 bp)	563
# contigs (>= 5000 bp)	65
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1859548
Total length (>= 1000 bp)	1859548
Total length (>= 5000 bp)	405689
Total length (>= 10000 bp)	11221
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	563
Largest contig	11221
Total length	1859548
GC (%)	56.68
N50	3062
N90	2386
auN	3808.2
L50	204
L90	482
# 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).

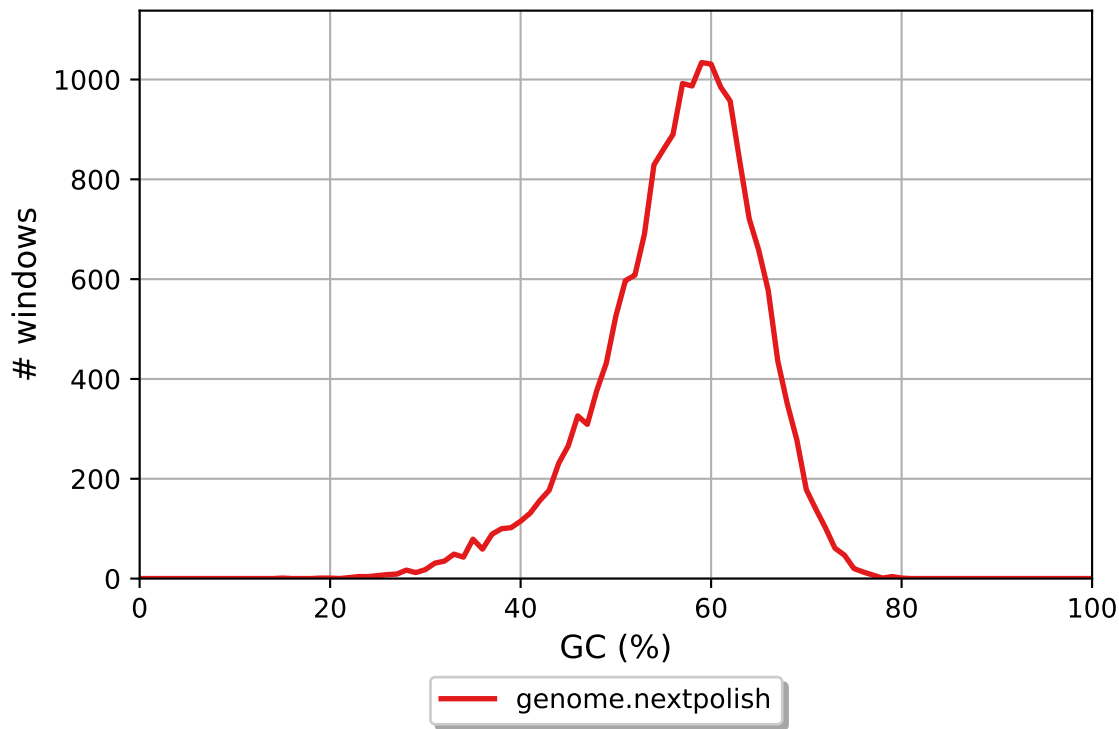
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

