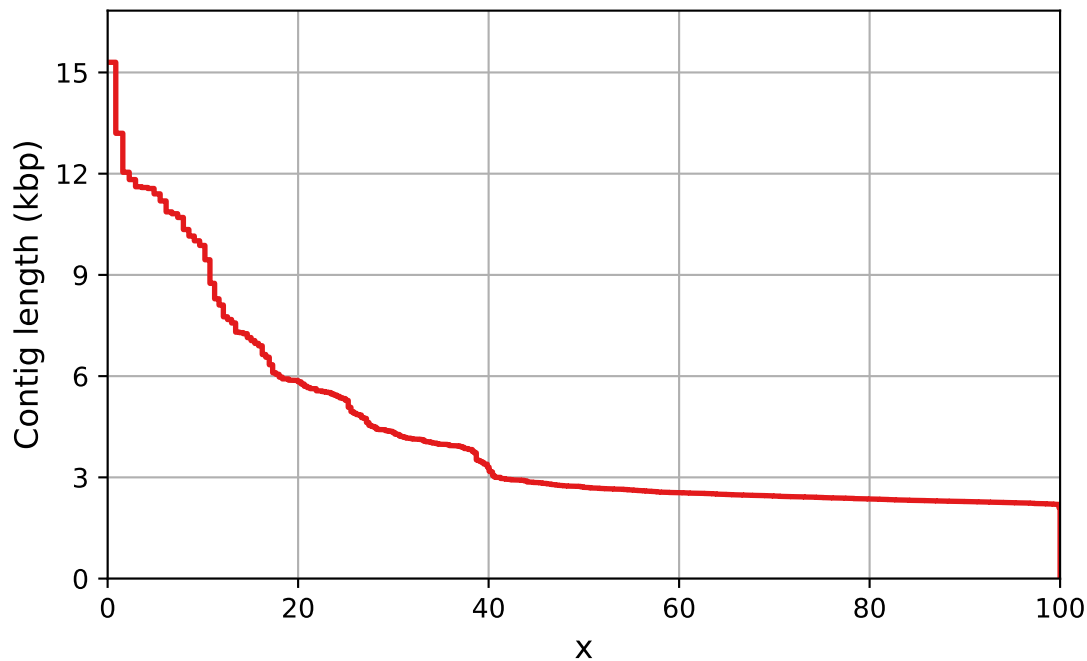


Report

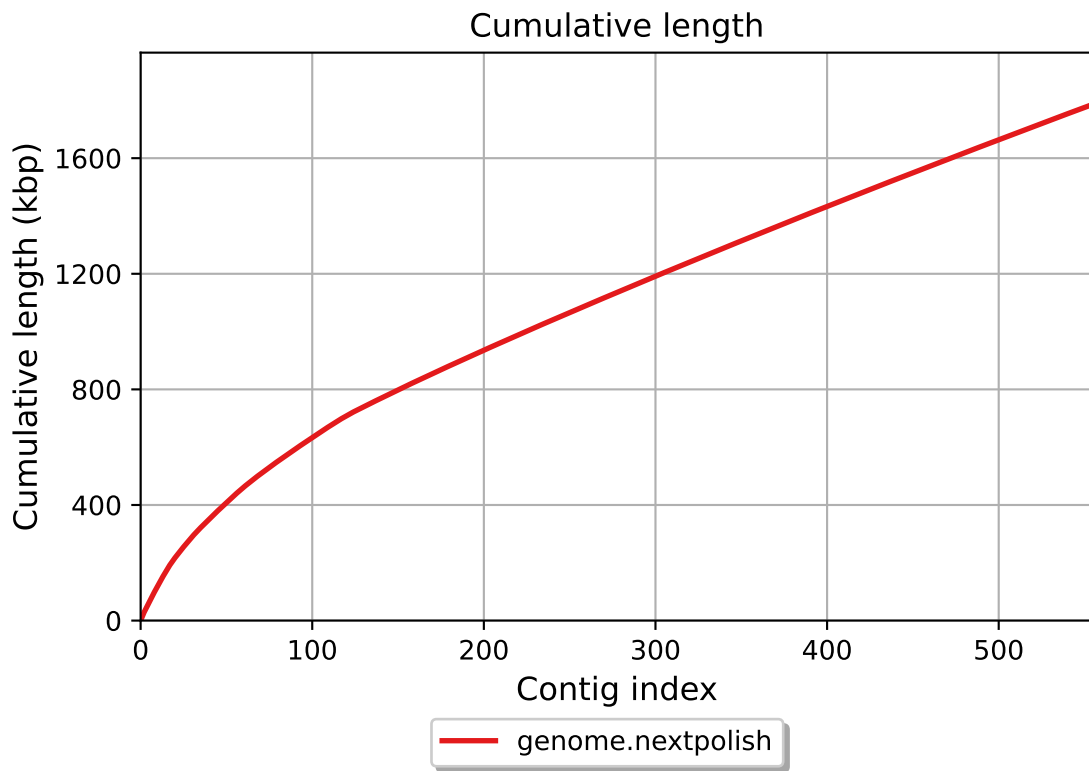
	genome.nextpolish
# contigs (>= 0 bp)	555
# contigs (>= 1000 bp)	555
# contigs (>= 5000 bp)	59
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1786549
Total length (>= 1000 bp)	1786549
Total length (>= 5000 bp)	456417
Total length (>= 10000 bp)	172613
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	555
Largest contig	15301
Total length	1786549
GC (%)	53.92
N50	2704
N90	2284
auN	4277.5
L50	185
L90	476
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

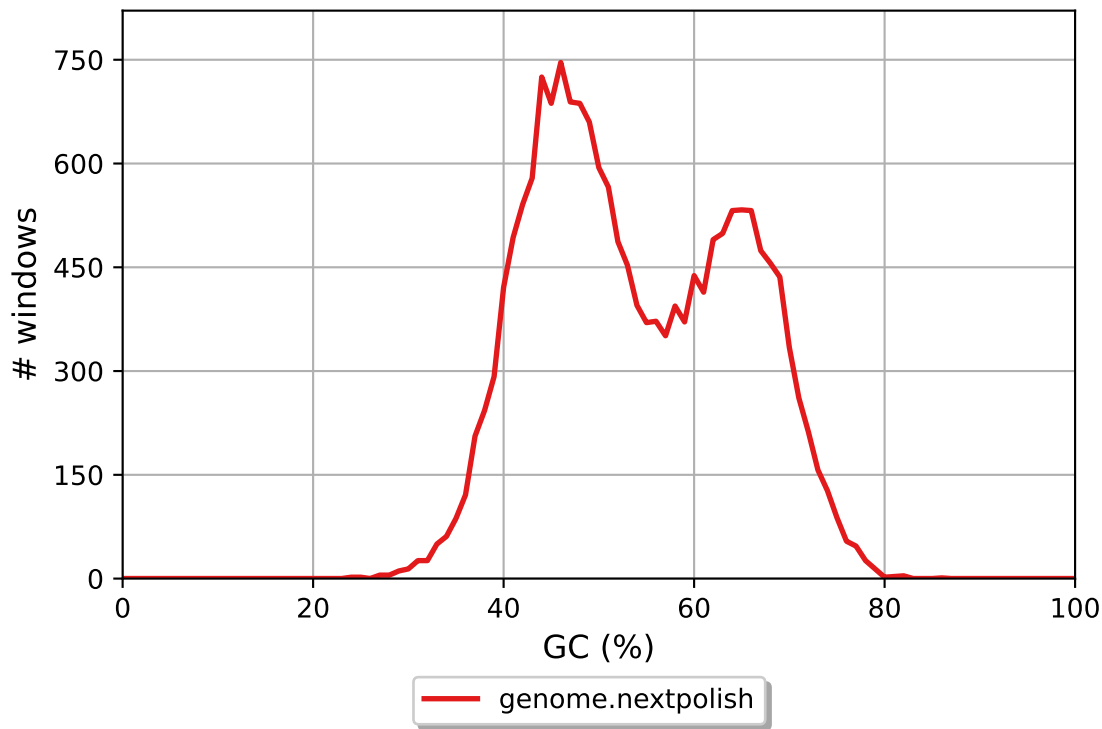
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

