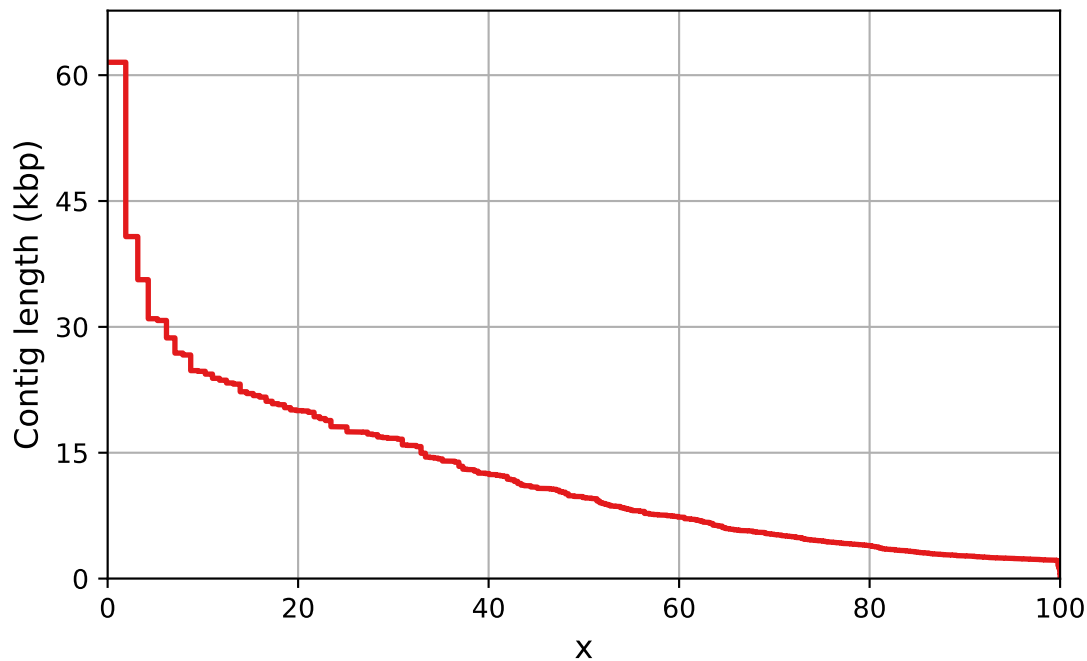


Report

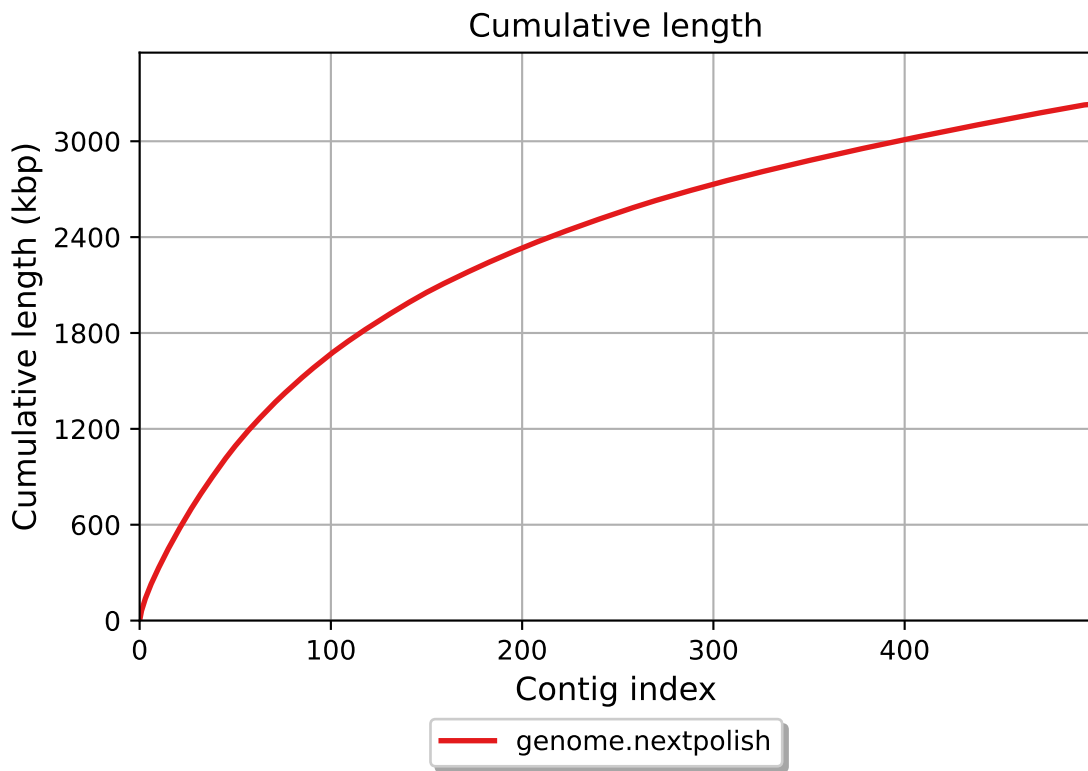
	genome.nextpolish
# contigs (>= 0 bp)	498
# contigs (>= 1000 bp)	496
# contigs (>= 5000 bp)	198
# contigs (>= 10000 bp)	89
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	3231462
Total length (>= 1000 bp)	3229867
Total length (>= 5000 bp)	2322096
Total length (>= 10000 bp)	1563238
Total length (>= 25000 bp)	281871
Total length (>= 50000 bp)	61537
# contigs	498
Largest contig	61537
Total length	3231462
GC (%)	58.86
N50	9657
N90	2707
auN	12714.5
L50	95
L90	361
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

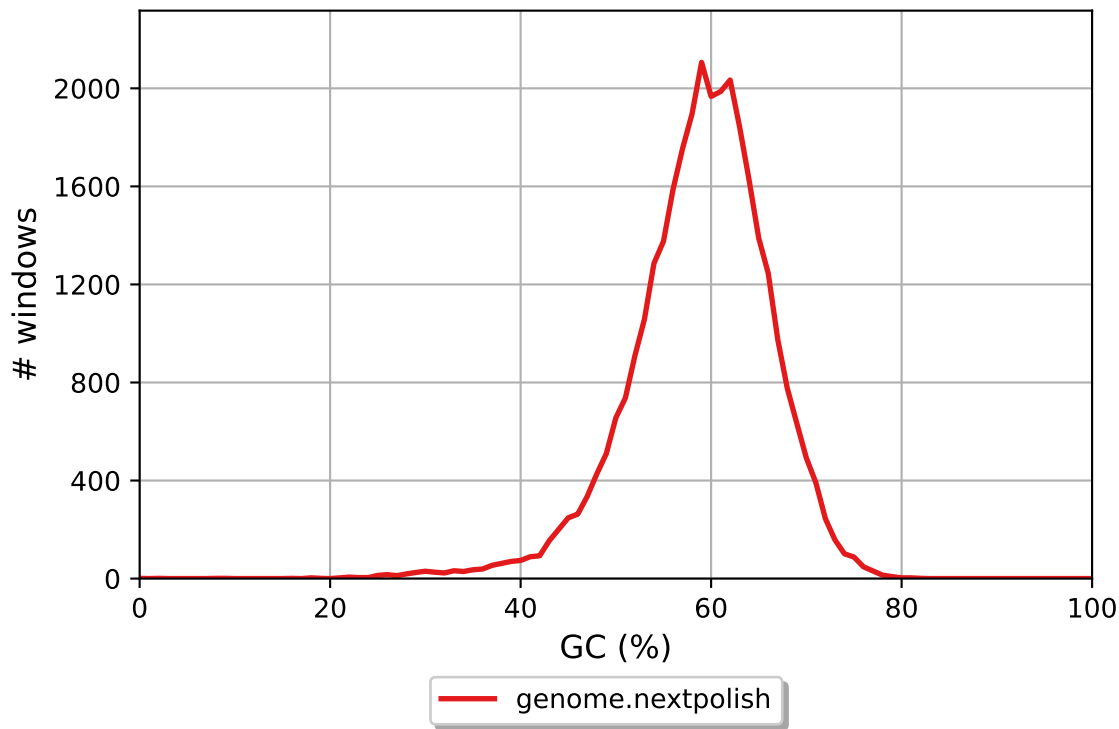
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

