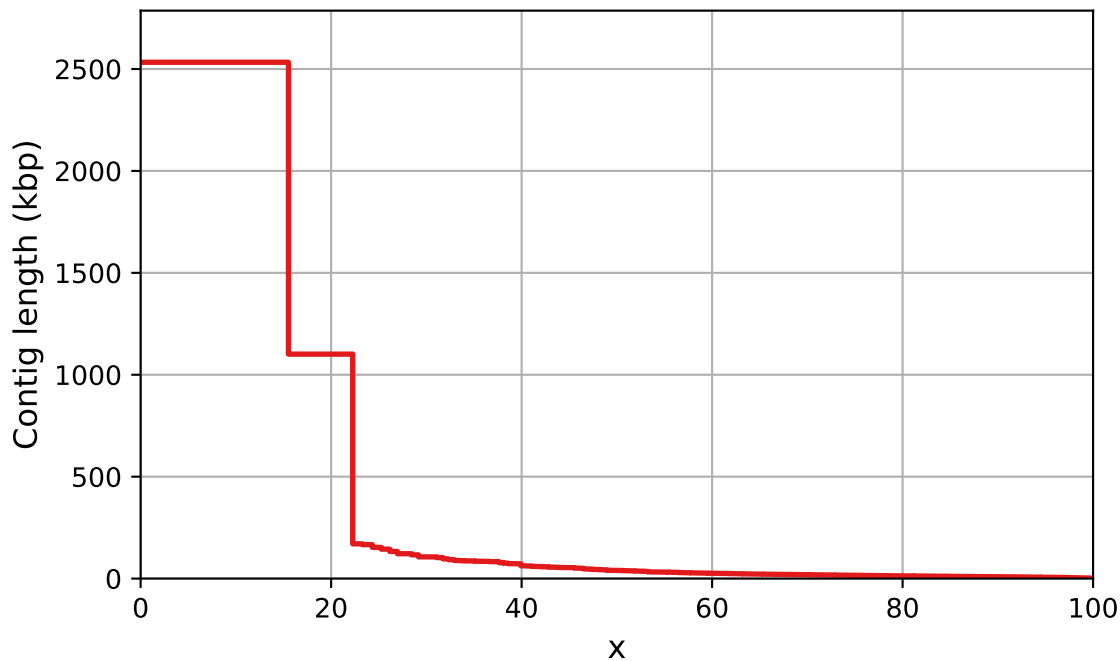


Report

	genome.nextpolish
# contigs (>= 0 bp)	745
# contigs (>= 1000 bp)	740
# contigs (>= 5000 bp)	607
# contigs (>= 10000 bp)	396
# contigs (>= 25000 bp)	123
# contigs (>= 50000 bp)	49
Total length (>= 0 bp)	16322232
Total length (>= 1000 bp)	16319022
Total length (>= 5000 bp)	15882896
Total length (>= 10000 bp)	14290397
Total length (>= 25000 bp)	10036093
Total length (>= 50000 bp)	7587542
# contigs	745
Largest contig	2533074
Total length	16322232
GC (%)	46.69
N50	39898
N90	9064
auN	500965.5
L50	63
L90	439
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 6 part

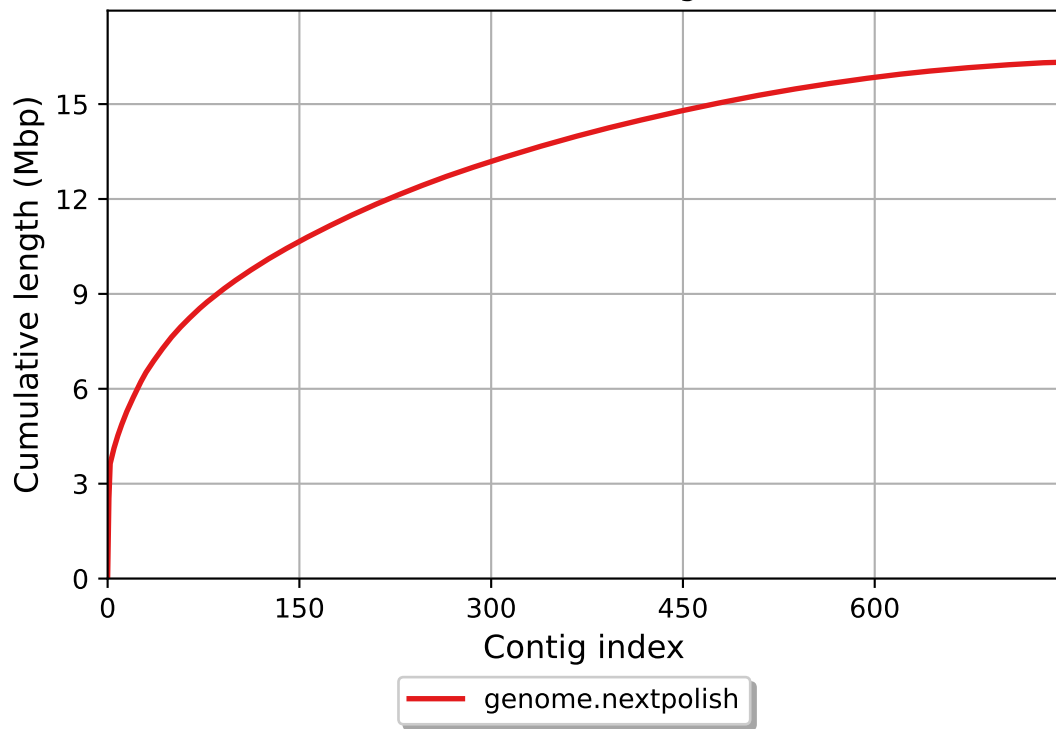
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

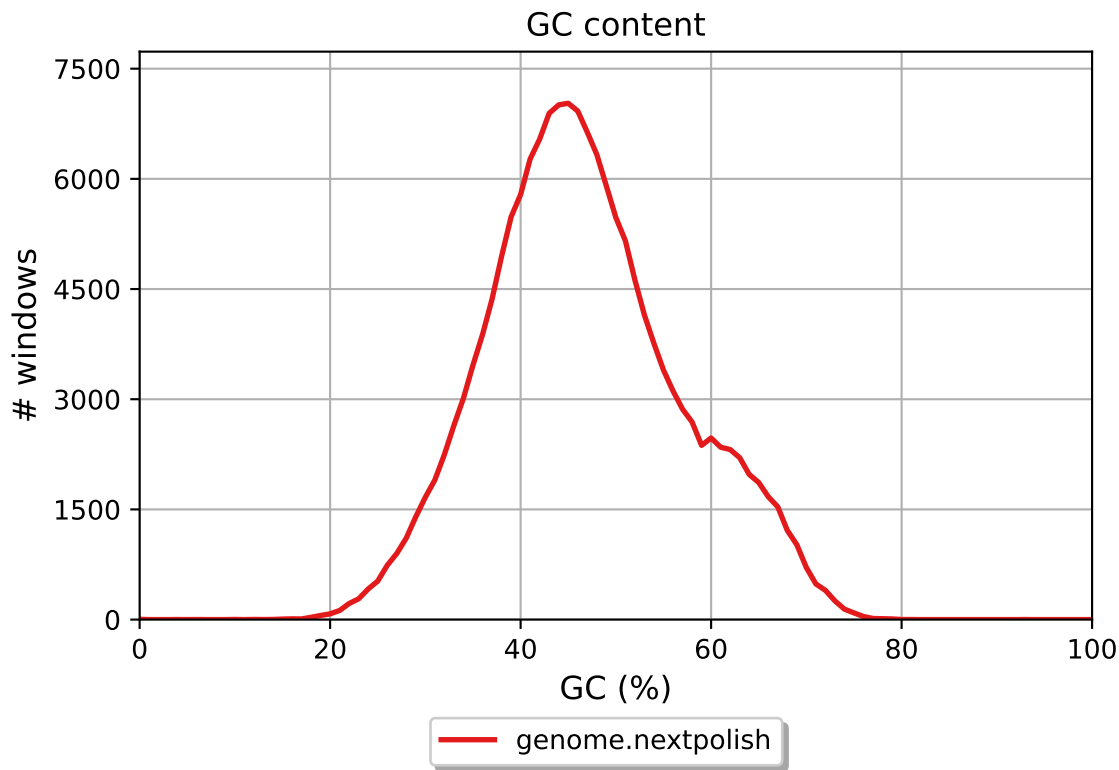
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

