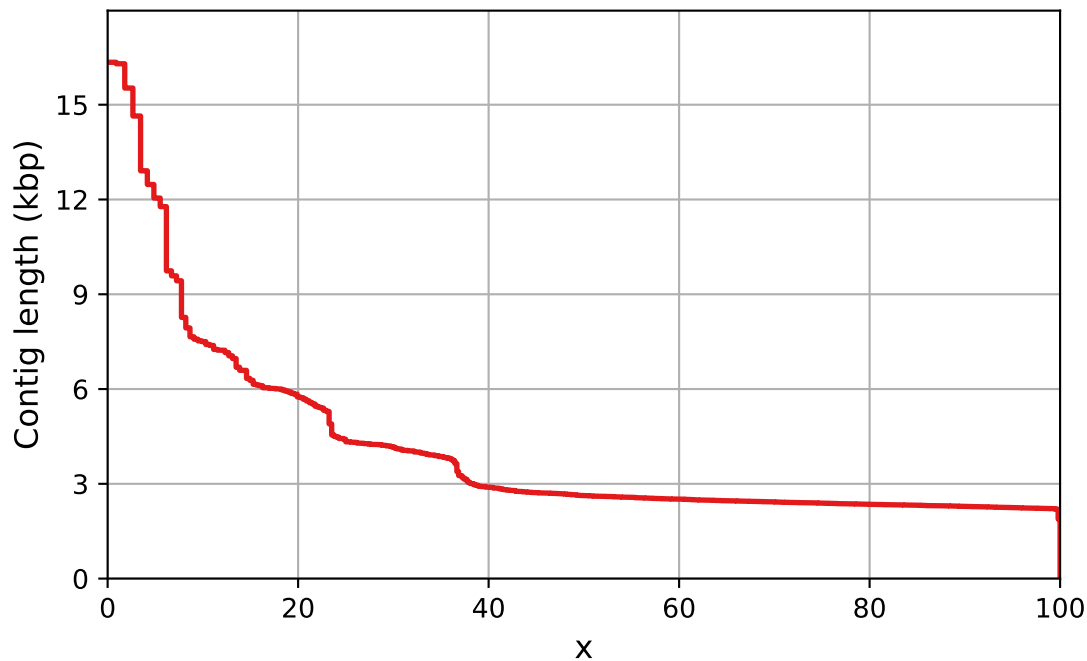


Report

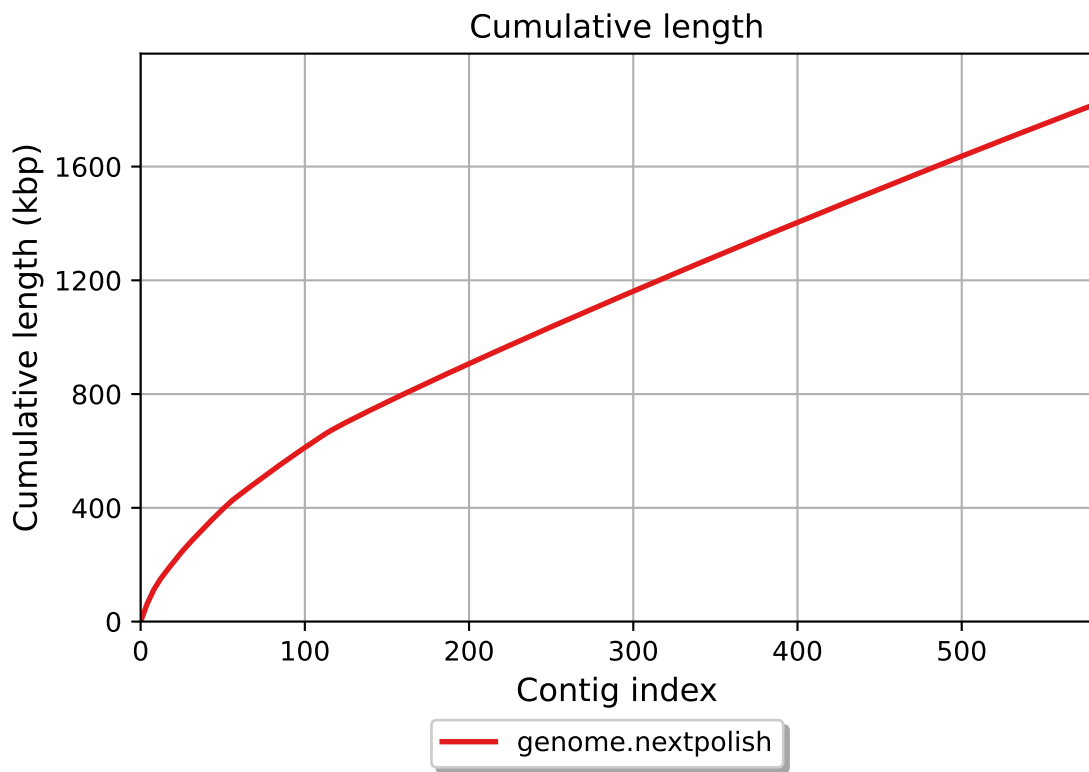
	genome.nextpolish
# contigs (>= 0 bp)	580
# contigs (>= 1000 bp)	580
# contigs (>= 5000 bp)	55
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1815544
Total length (>= 1000 bp)	1815544
Total length (>= 5000 bp)	422178
Total length (>= 10000 bp)	112012
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	580
Largest contig	16343
Total length	1815544
GC (%)	54.15
N50	2624
N90	2283
auN	4165.5
L50	201
L90	499
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 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).

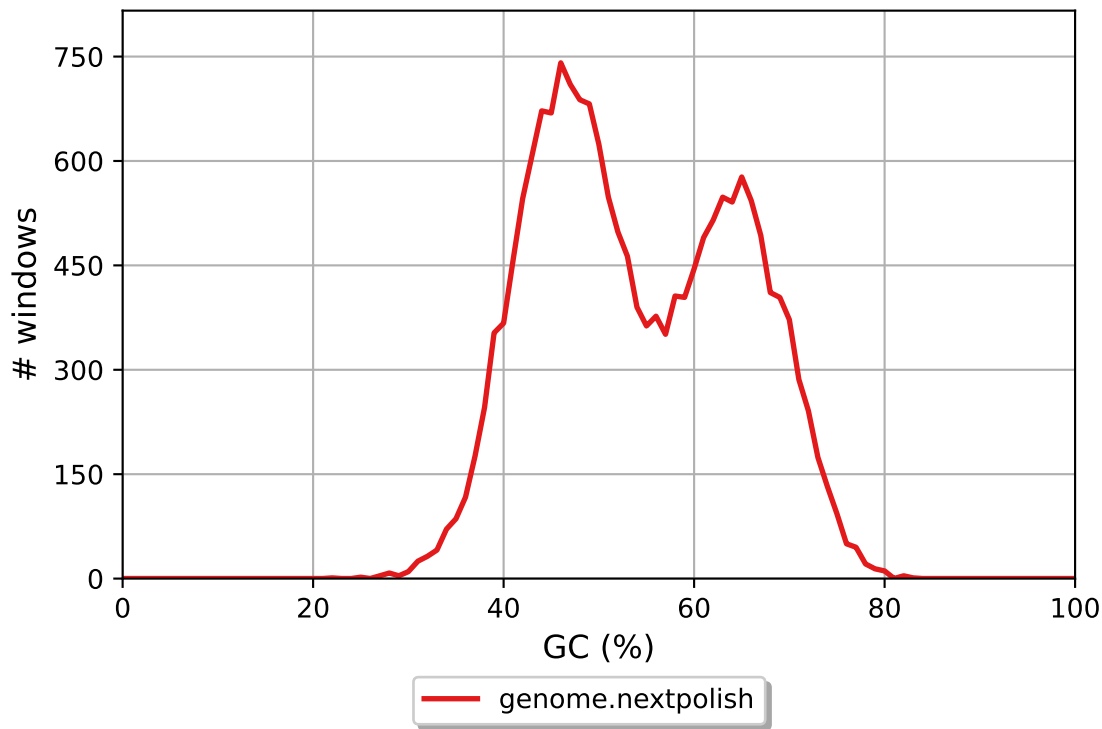
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

