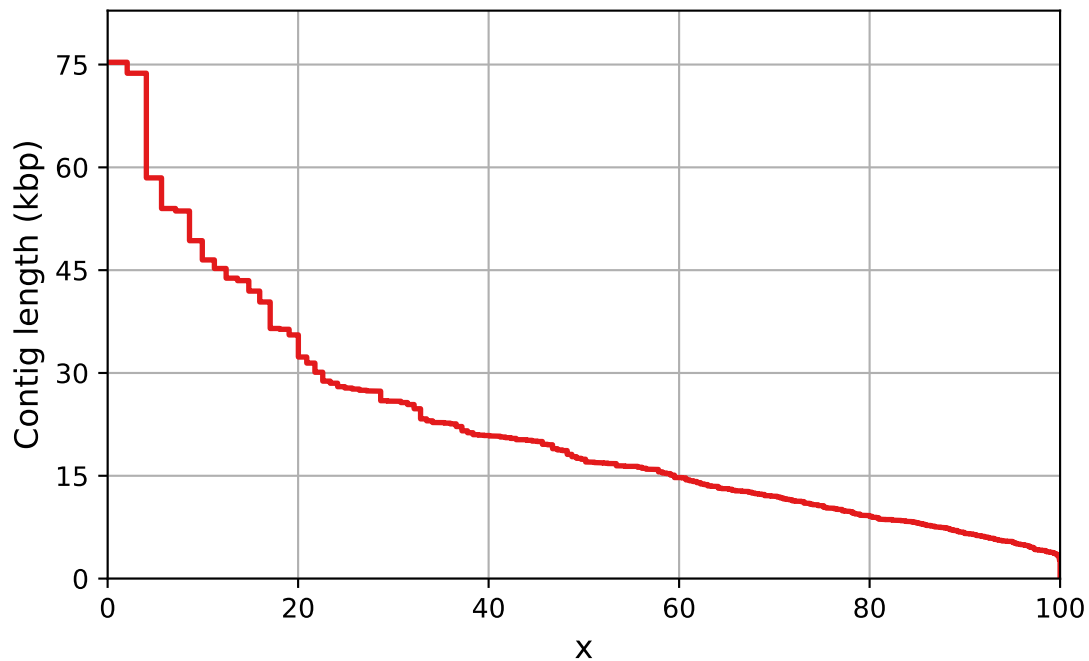


Report

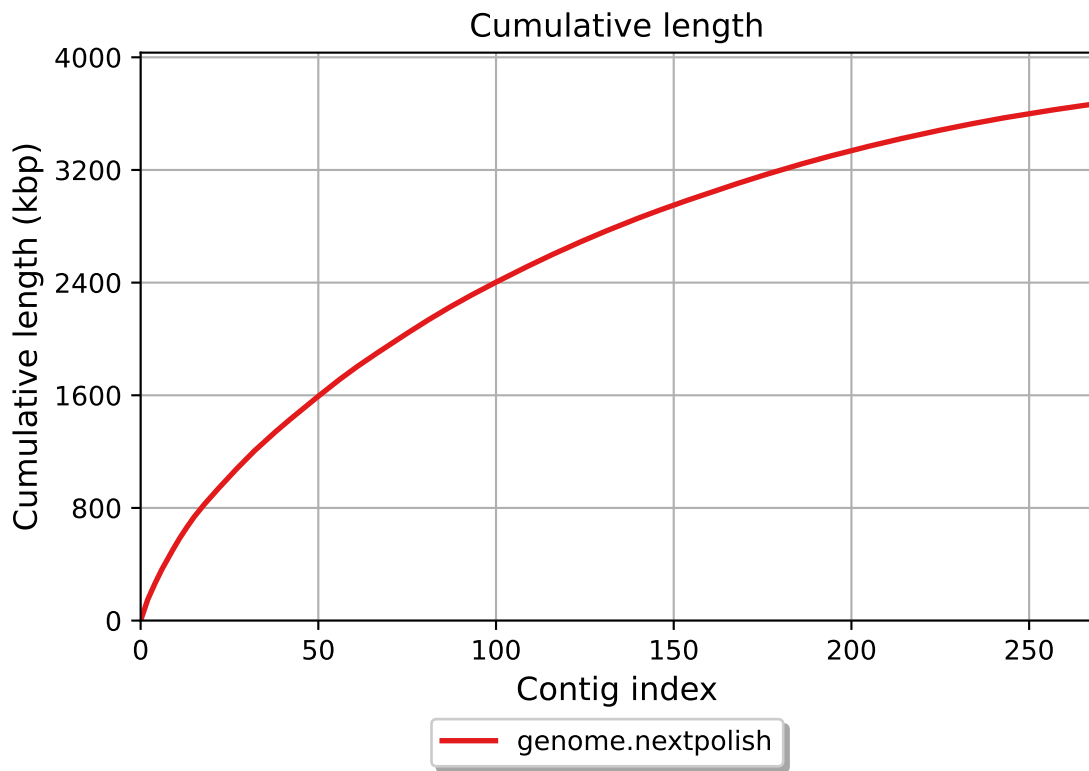
	genome.nextpolish
# contigs (>= 0 bp)	268
# contigs (>= 1000 bp)	268
# contigs (>= 5000 bp)	231
# contigs (>= 10000 bp)	137
# contigs (>= 25000 bp)	31
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	3666494
Total length (>= 1000 bp)	3666494
Total length (>= 5000 bp)	3513694
Total length (>= 10000 bp)	2828079
Total length (>= 25000 bp)	1179861
Total length (>= 50000 bp)	315158
# contigs	268
Largest contig	75332
Total length	3666494
GC (%)	40.64
N50	17412
N90	6584
auN	22812.5
L50	63
L90	195
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

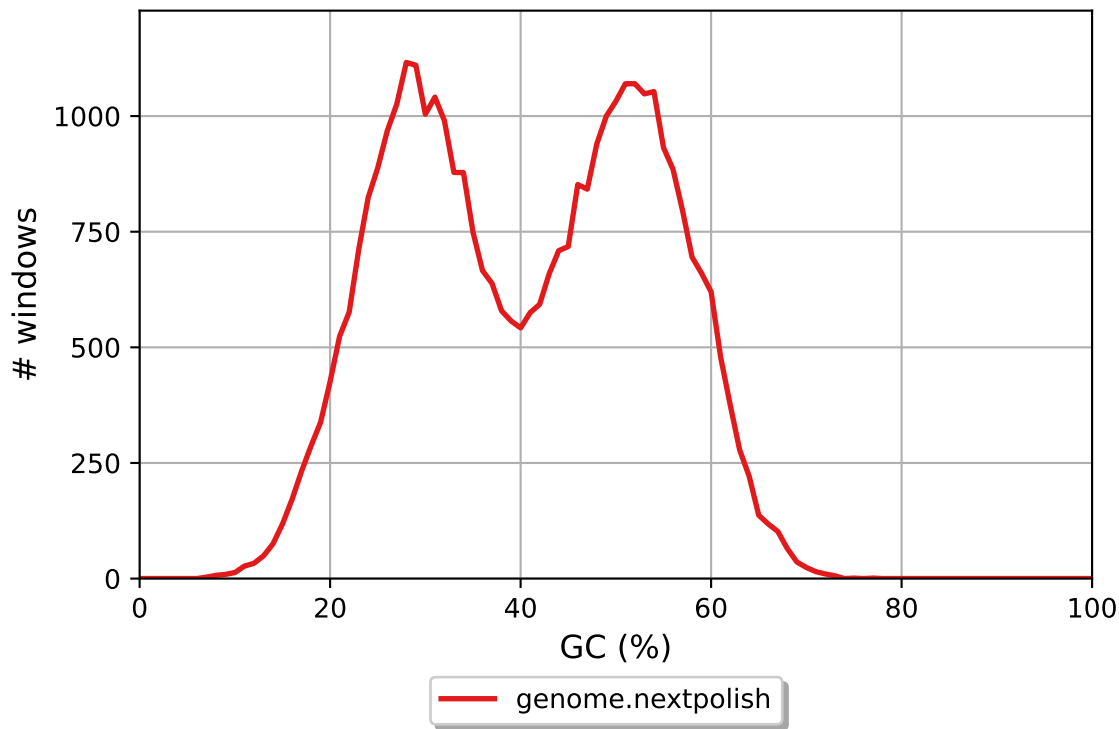
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

