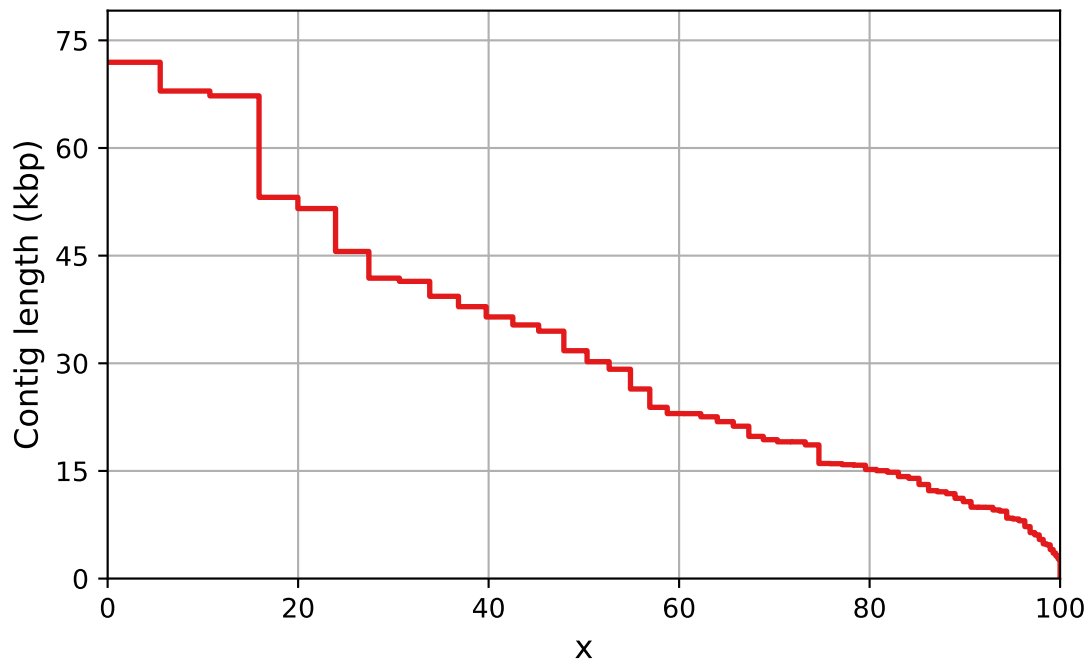


Report

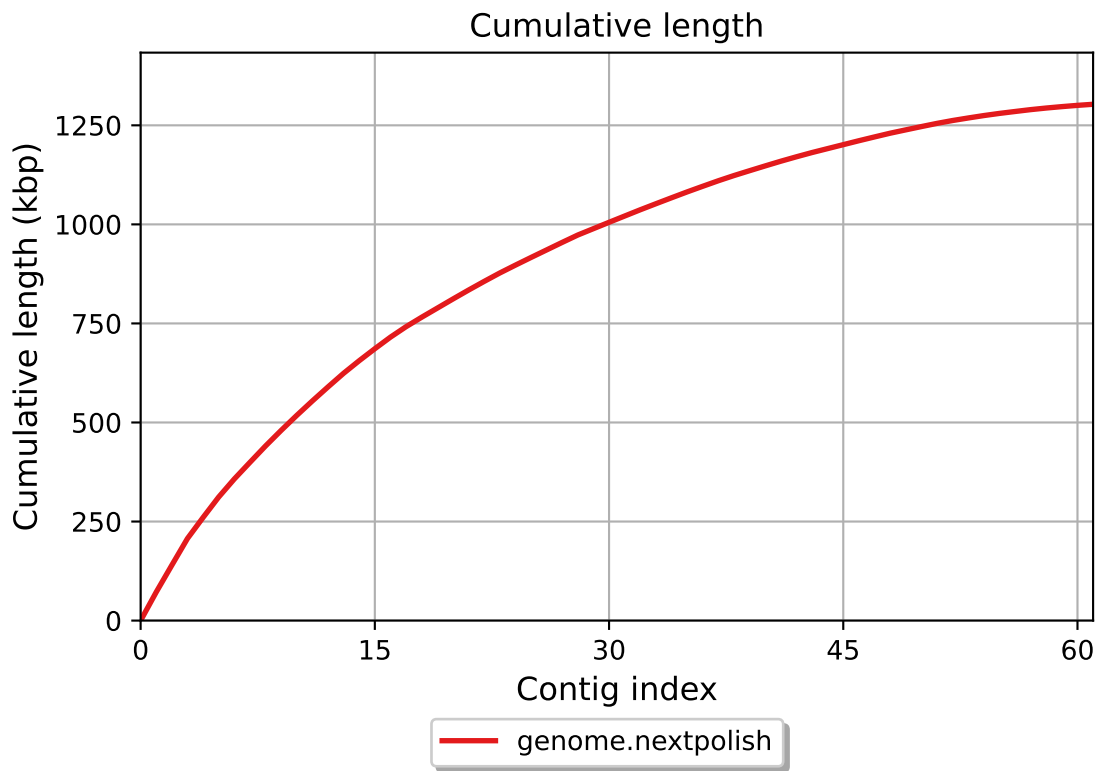
	genome.nextpolish
# contigs (>= 0 bp)	61
# contigs (>= 1000 bp)	61
# contigs (>= 5000 bp)	55
# contigs (>= 10000 bp)	43
# contigs (>= 25000 bp)	17
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	1303145
Total length (>= 1000 bp)	1303145
Total length (>= 5000 bp)	1280068
Total length (>= 10000 bp)	1181317
Total length (>= 25000 bp)	741772
Total length (>= 50000 bp)	311839
# contigs	61
Largest contig	71946
Total length	1303145
GC (%)	40.09
N50	31752
N90	10722
auN	34024.4
L50	14
L90	43
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

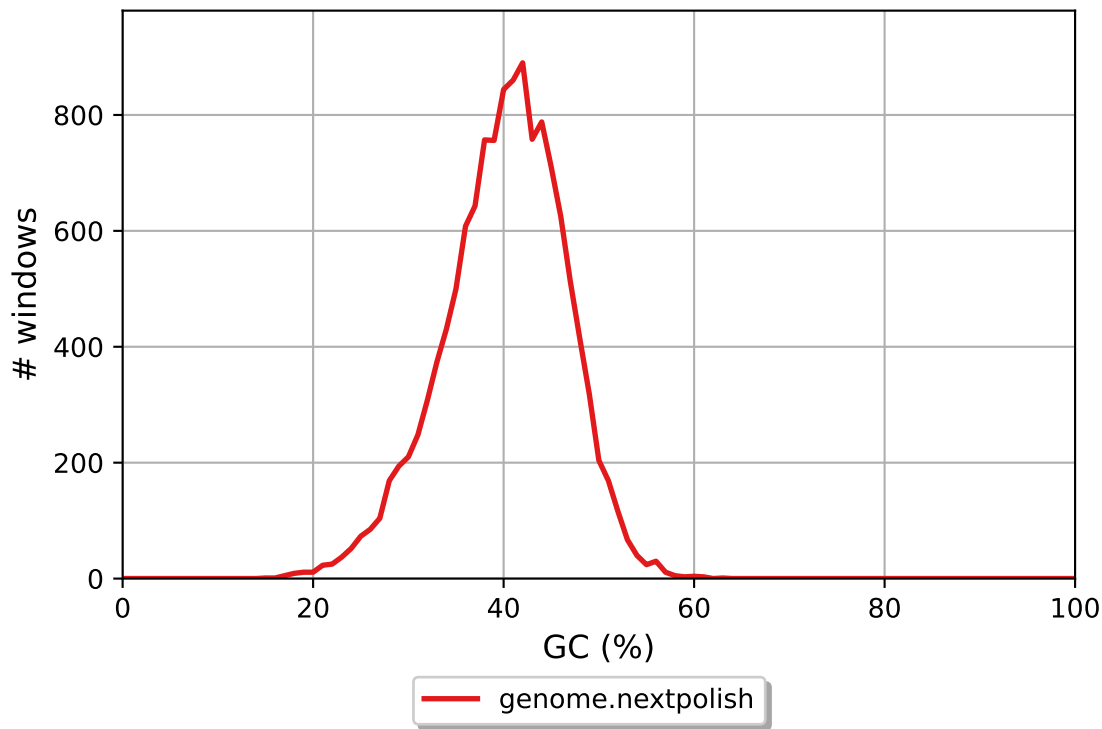
Nx



genome.nextpolish



GC content



genome.nextpolish GC content

