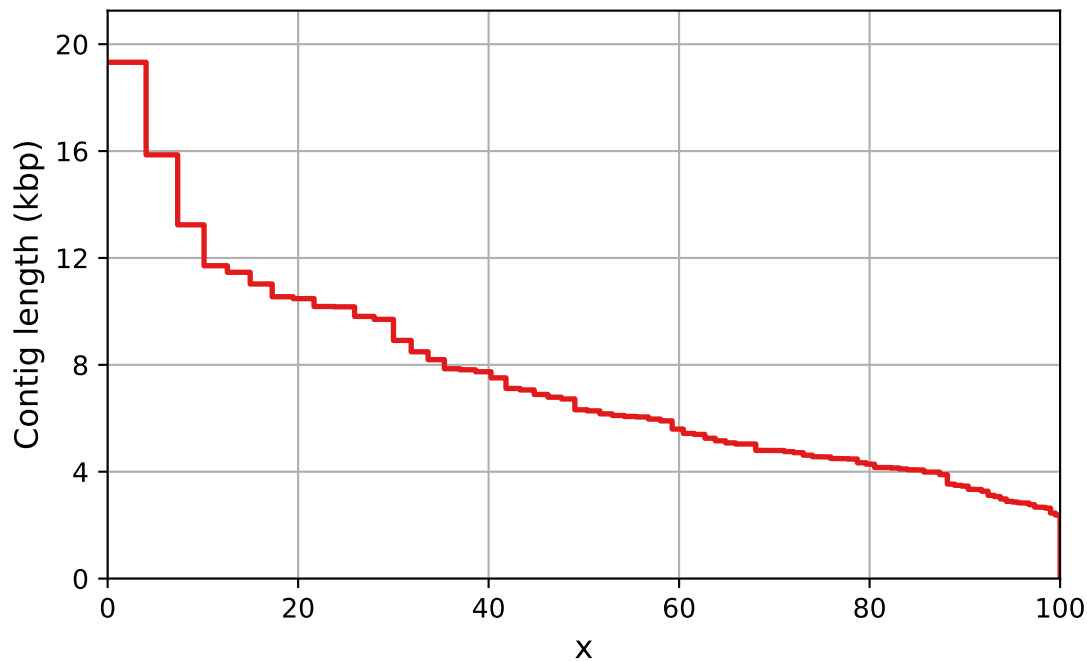


Report

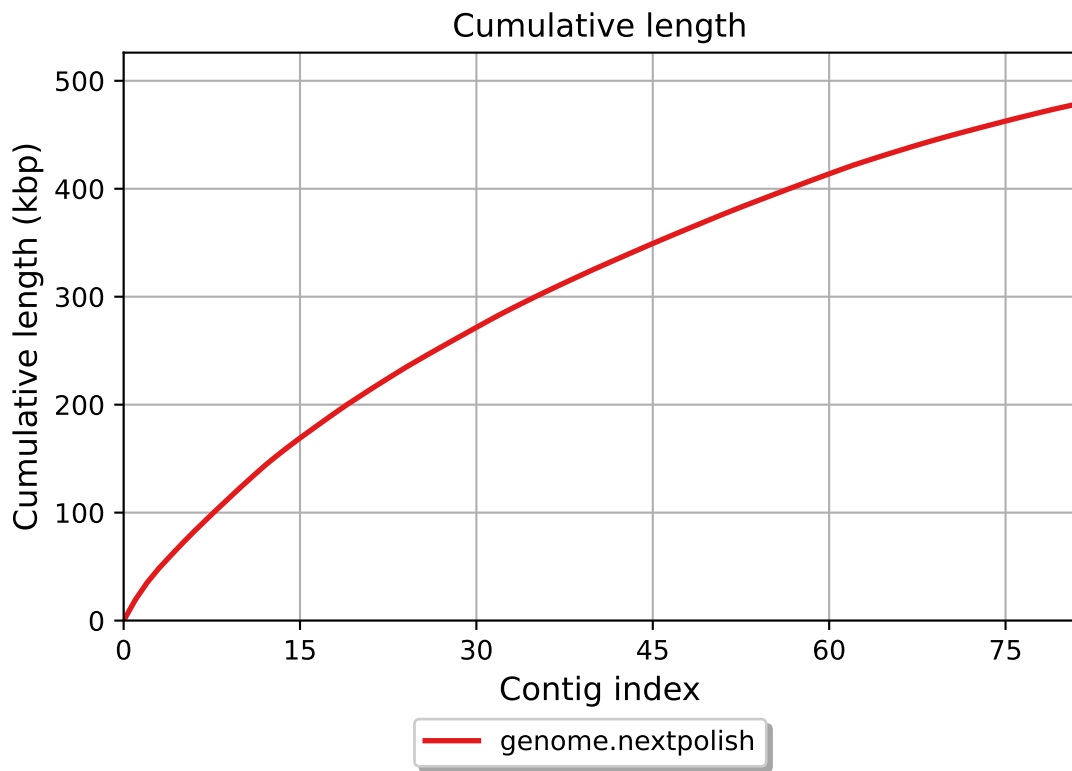
	genome.nextpolish
# contigs (>= 0 bp)	81
# contigs (>= 1000 bp)	81
# contigs (>= 5000 bp)	40
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	478255
Total length (>= 1000 bp)	478255
Total length (>= 5000 bp)	325444
Total length (>= 10000 bp)	123998
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	81
Largest contig	19323
Total length	478255
GC (%)	38.91
N50	6318
N90	3462
auN	7565.8
L50	25
L90	65
# 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).

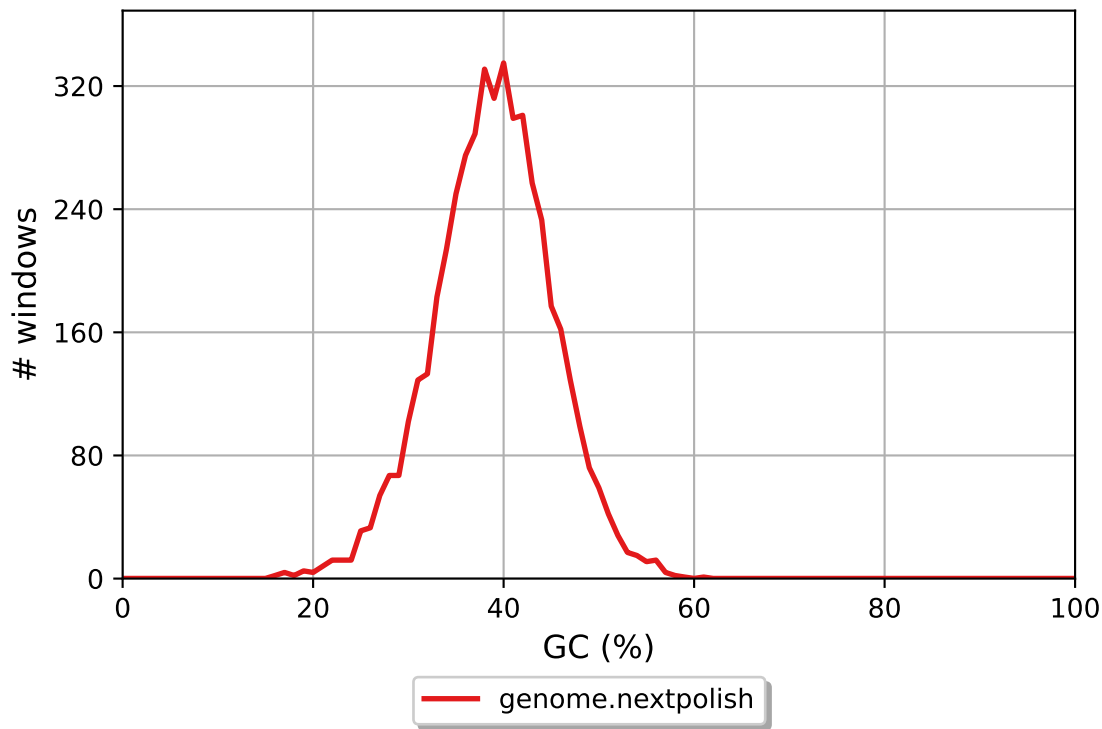
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

