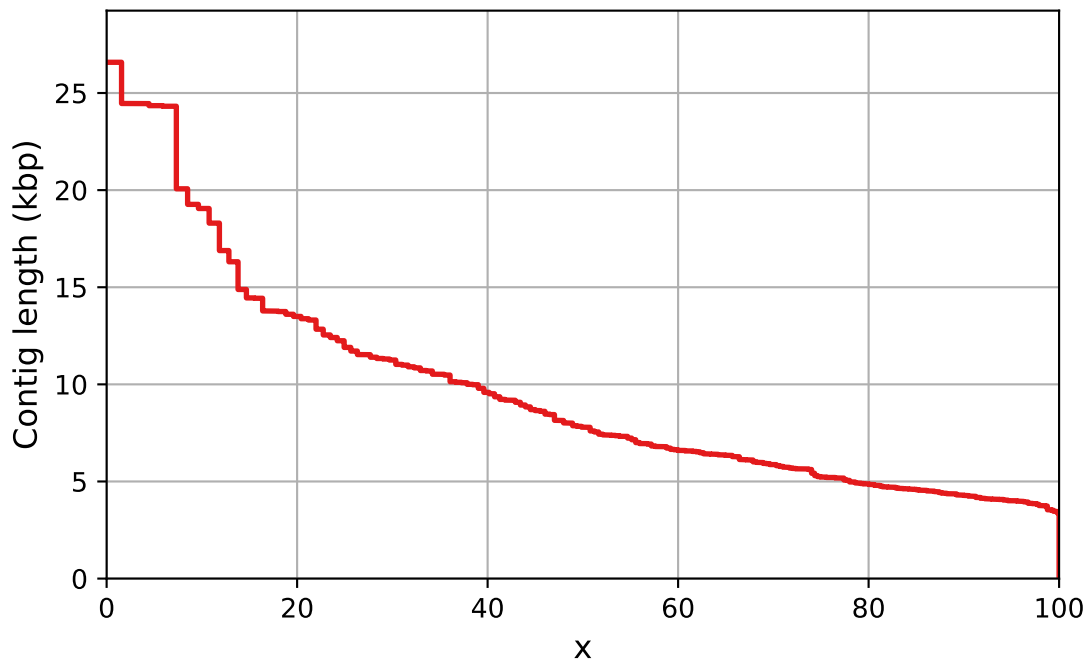


# Report

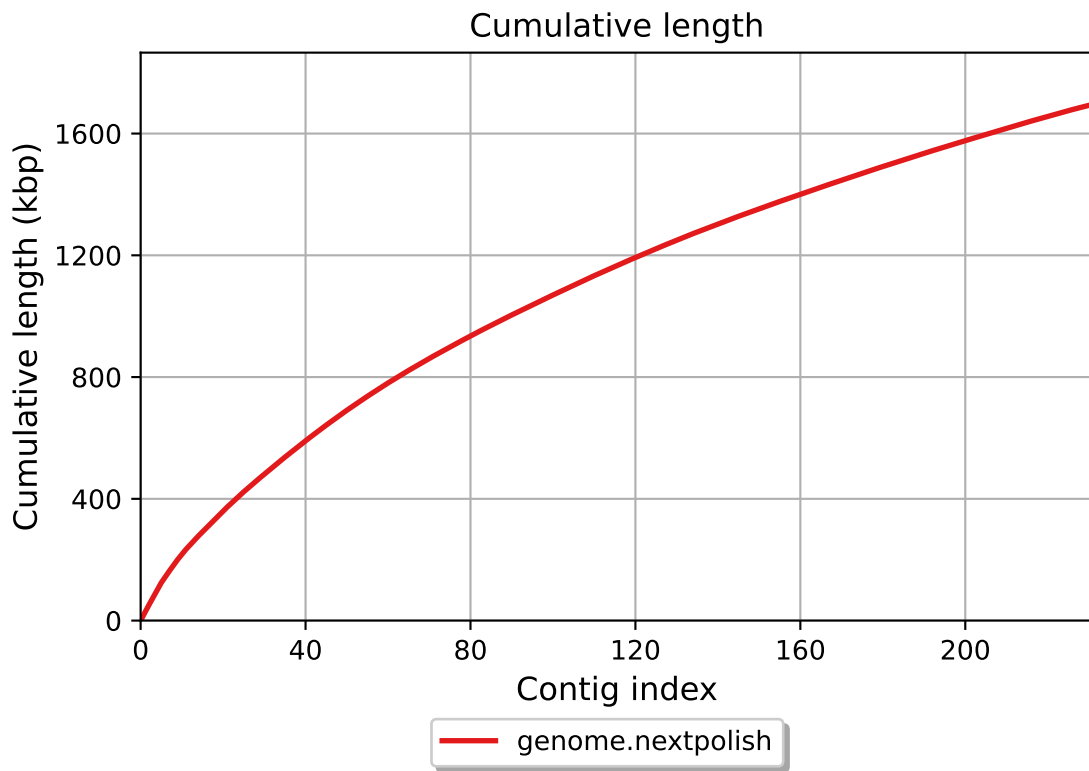
	genome.nextpolish
# contigs (>= 0 bp)	231
# contigs (>= 1000 bp)	231
# contigs (>= 5000 bp)	144
# contigs (>= 10000 bp)	46
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1696207
Total length (>= 1000 bp)	1696207
Total length (>= 5000 bp)	1323128
Total length (>= 10000 bp)	651937
Total length (>= 25000 bp)	26582
Total length (>= 50000 bp)	0
# contigs	231
Largest contig	26582
Total length	1696207
GC (%)	41.12
N50	7796
N90	4283
auN	9750.8
L50	69
L90	189
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 2 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).

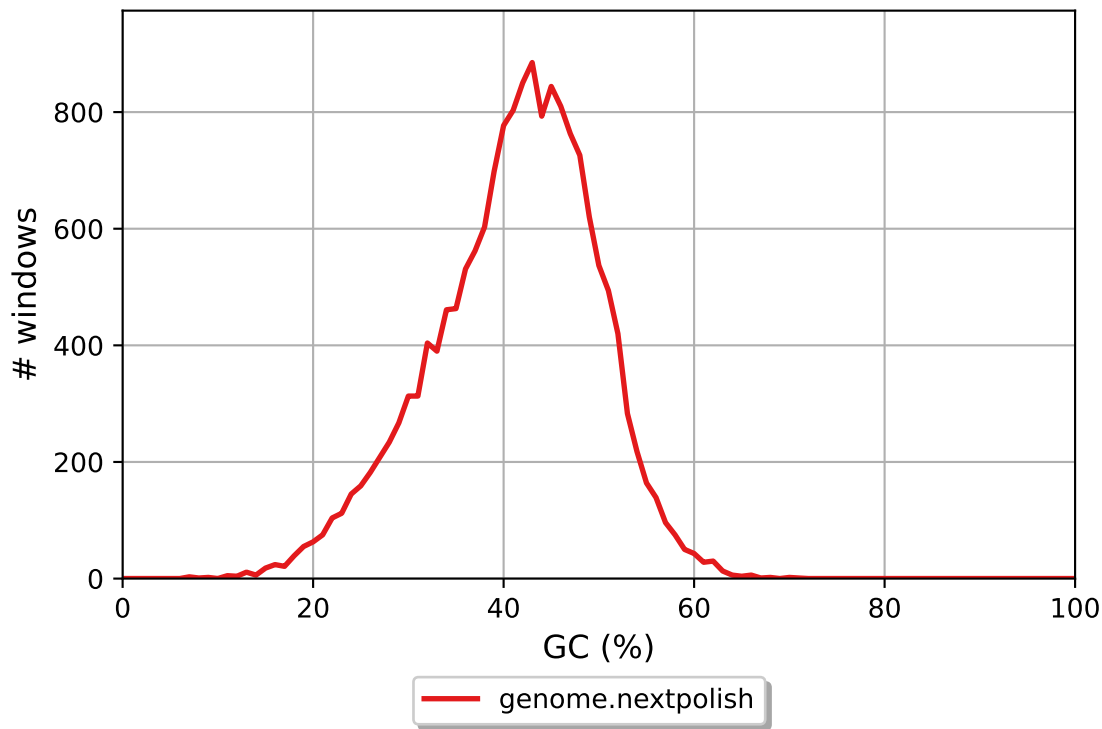
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

