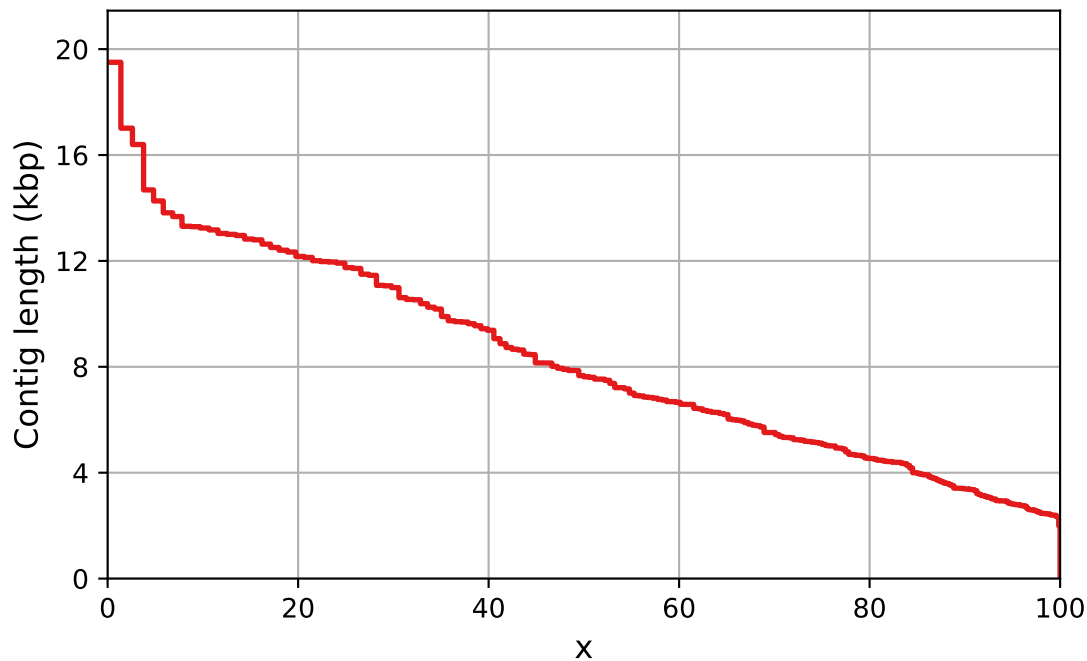


Report

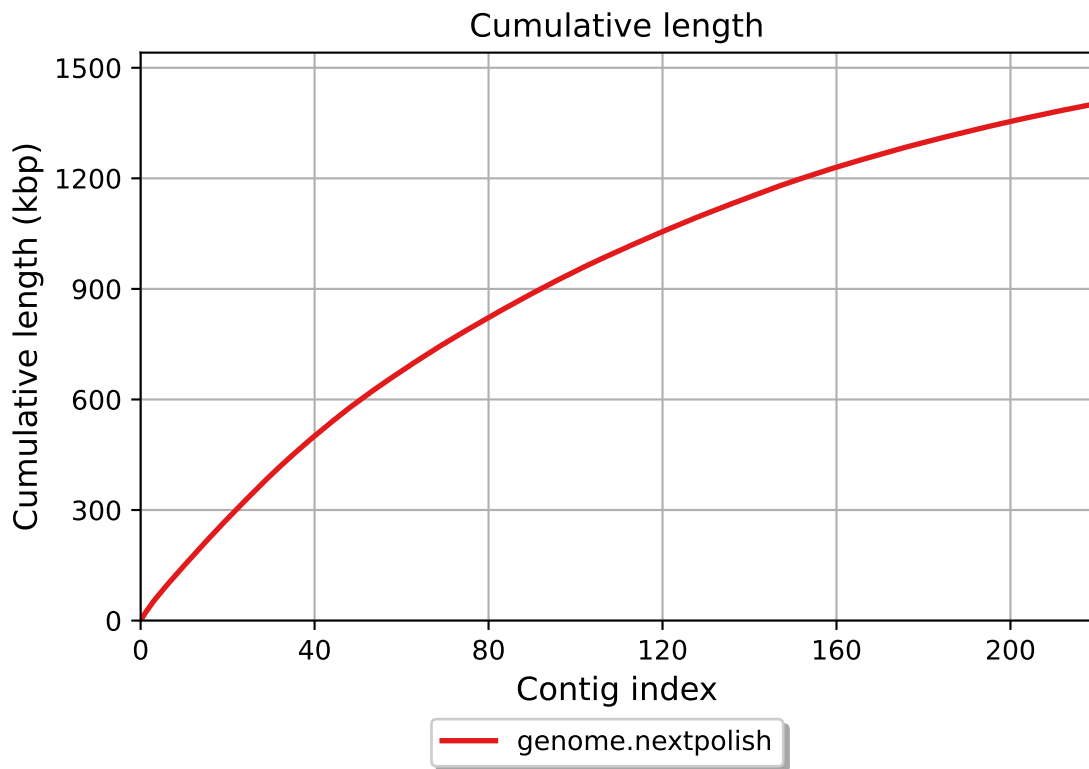
	genome.nextpolish
# contigs (>= 0 bp)	219
# contigs (>= 1000 bp)	219
# contigs (>= 5000 bp)	123
# contigs (>= 10000 bp)	39
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1400990
Total length (>= 1000 bp)	1400990
Total length (>= 5000 bp)	1070518
Total length (>= 10000 bp)	491075
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	219
Largest contig	19503
Total length	1400990
GC (%)	40.73
N50	7666
N90	3395
auN	8335.3
L50	63
L90	169
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

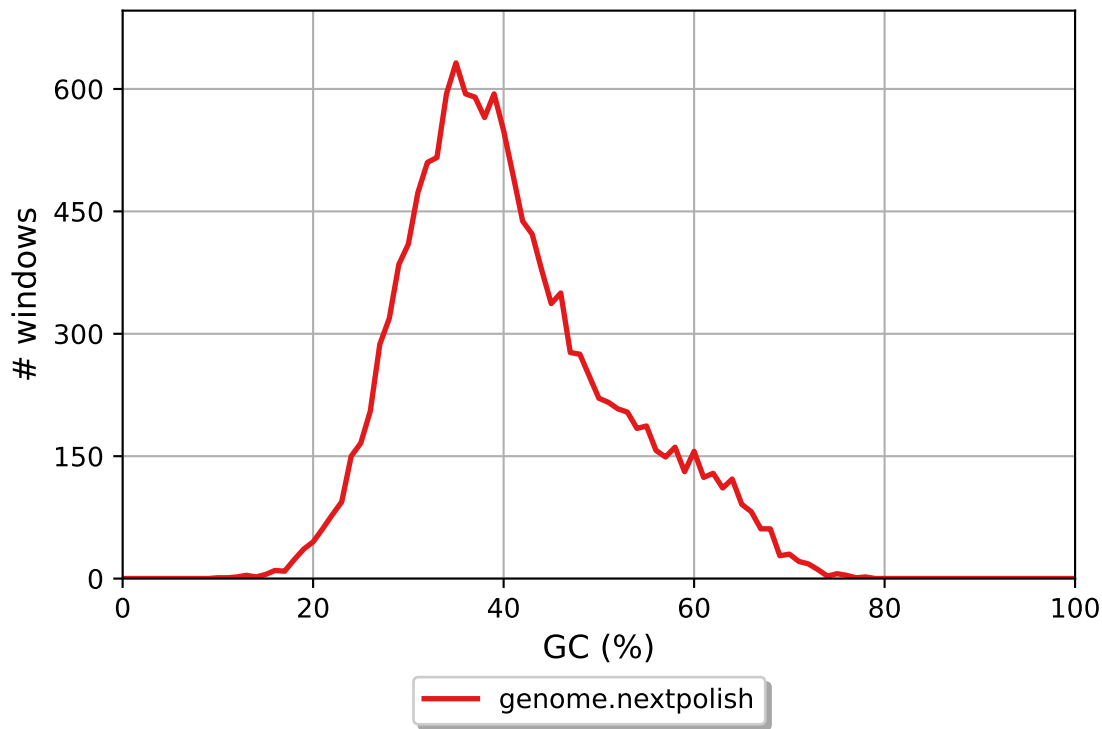
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

