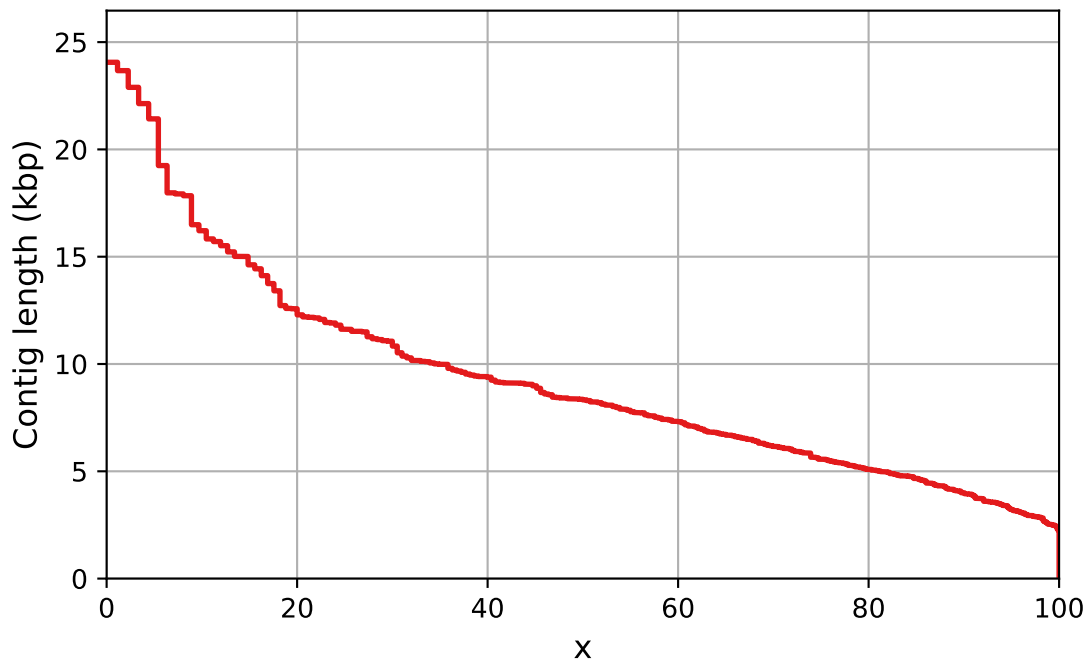


Report

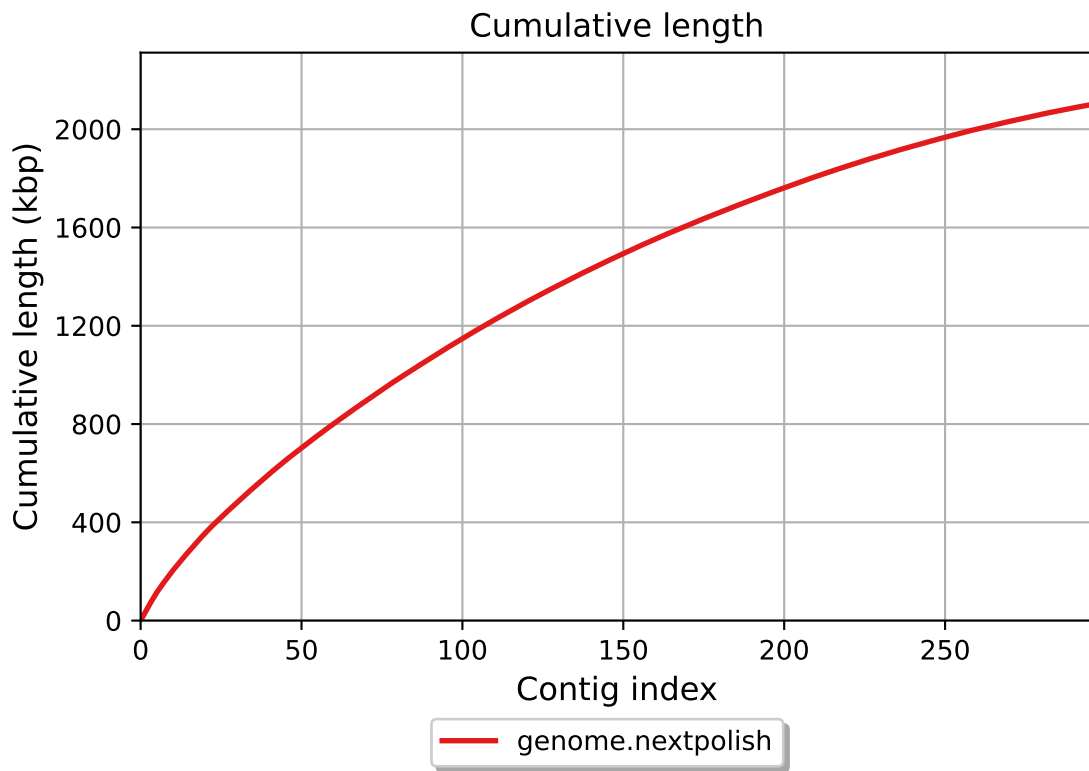
	genome.nextpolish
# contigs (>= 0 bp)	296
# contigs (>= 1000 bp)	296
# contigs (>= 5000 bp)	189
# contigs (>= 10000 bp)	53
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2101607
Total length (>= 1000 bp)	2101607
Total length (>= 5000 bp)	1707797
Total length (>= 10000 bp)	733069
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	296
Largest contig	24060
Total length	2101607
GC (%)	33.44
N50	8328
N90	3995
auN	9352.5
L50	89
L90	230
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 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).

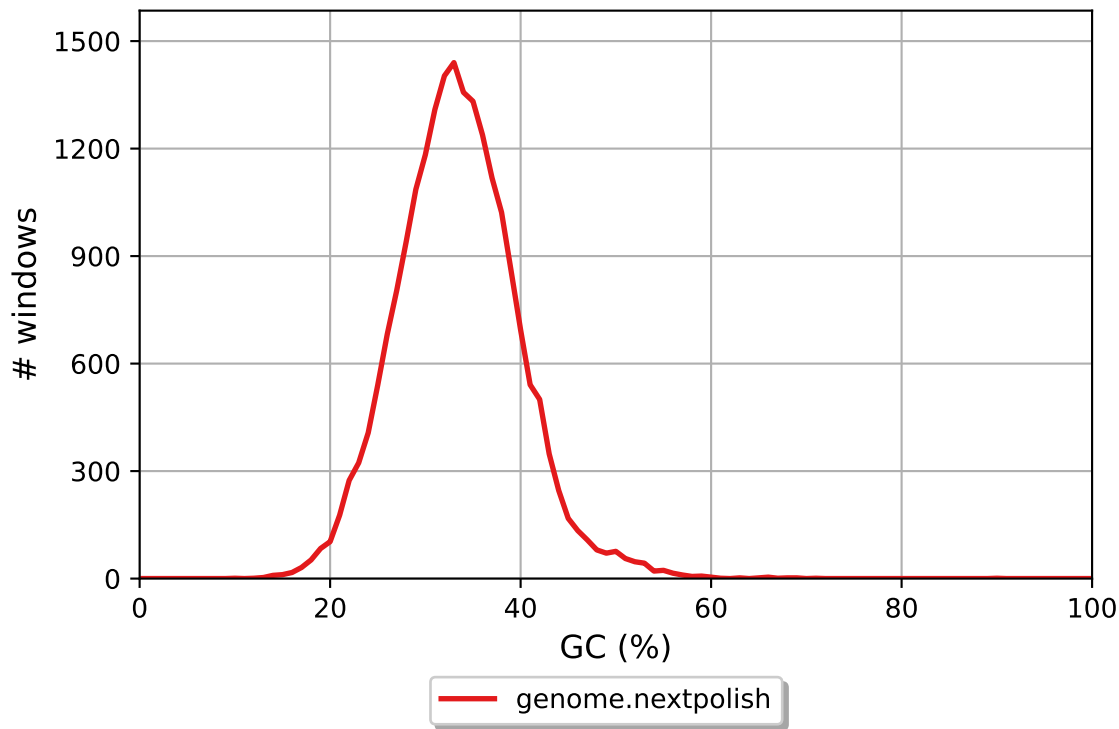
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

