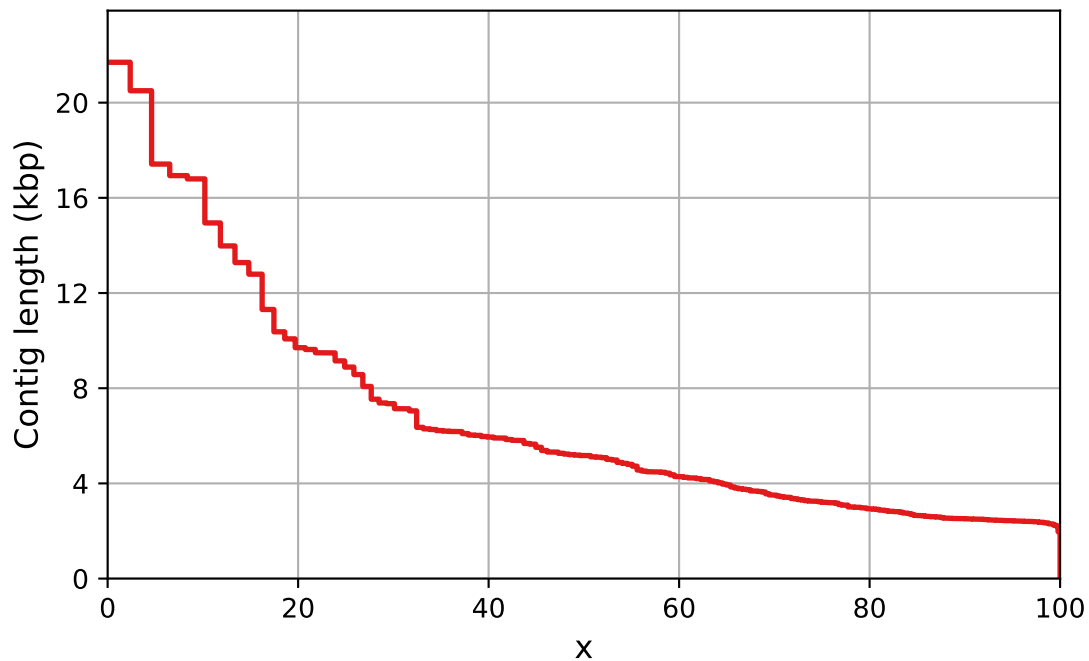


Report

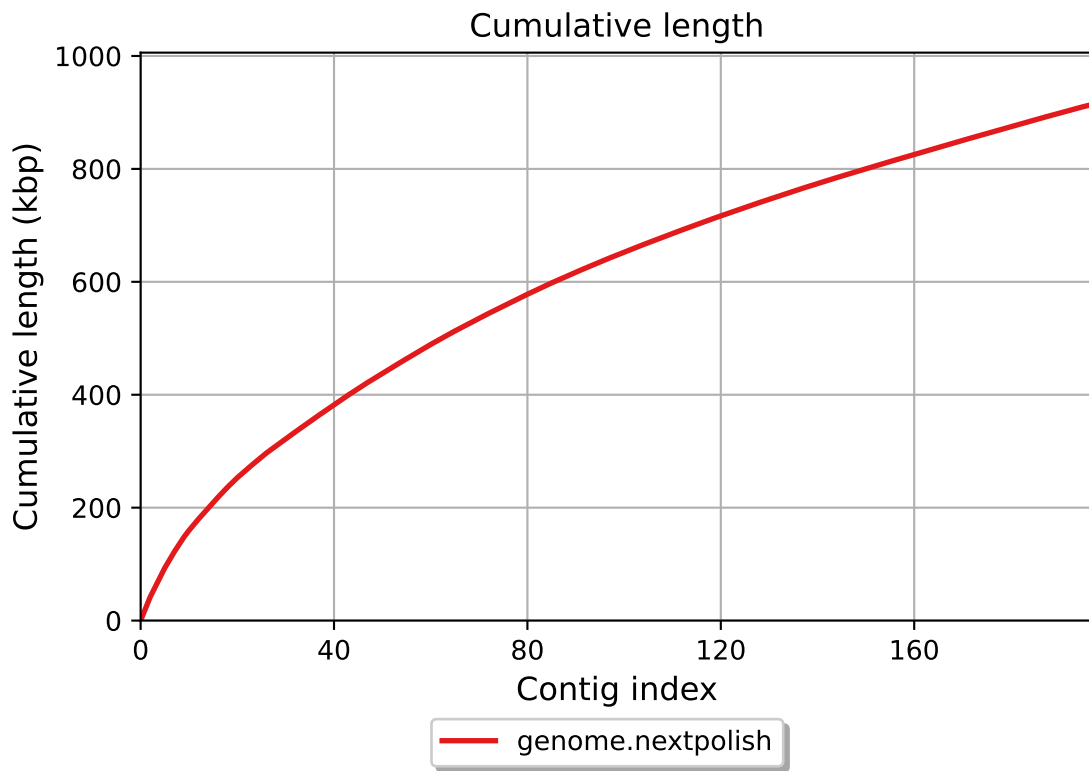
	genome.nextpolish
# contigs (>= 0 bp)	197
# contigs (>= 1000 bp)	197
# contigs (>= 5000 bp)	59
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	914467
Total length (>= 1000 bp)	914467
Total length (>= 5000 bp)	484103
Total length (>= 10000 bp)	180096
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	197
Largest contig	21697
Total length	914467
GC (%)	41.78
N50	5172
N90	2518
auN	6924.2
L50	54
L90	159
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

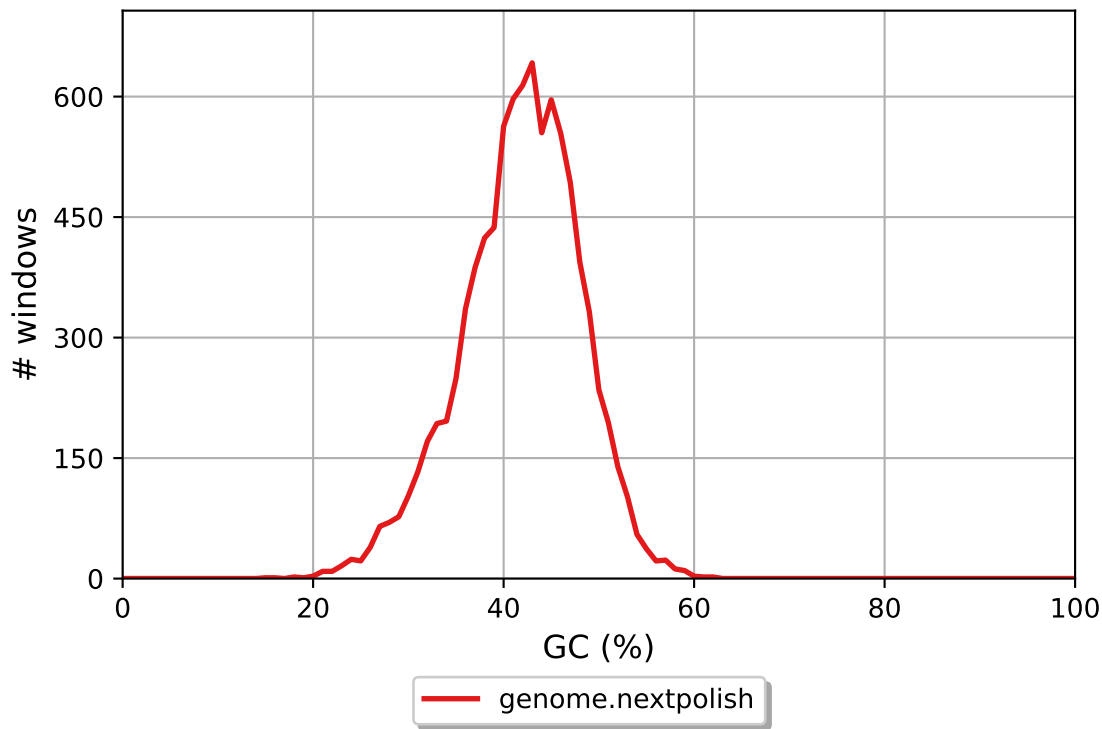
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

