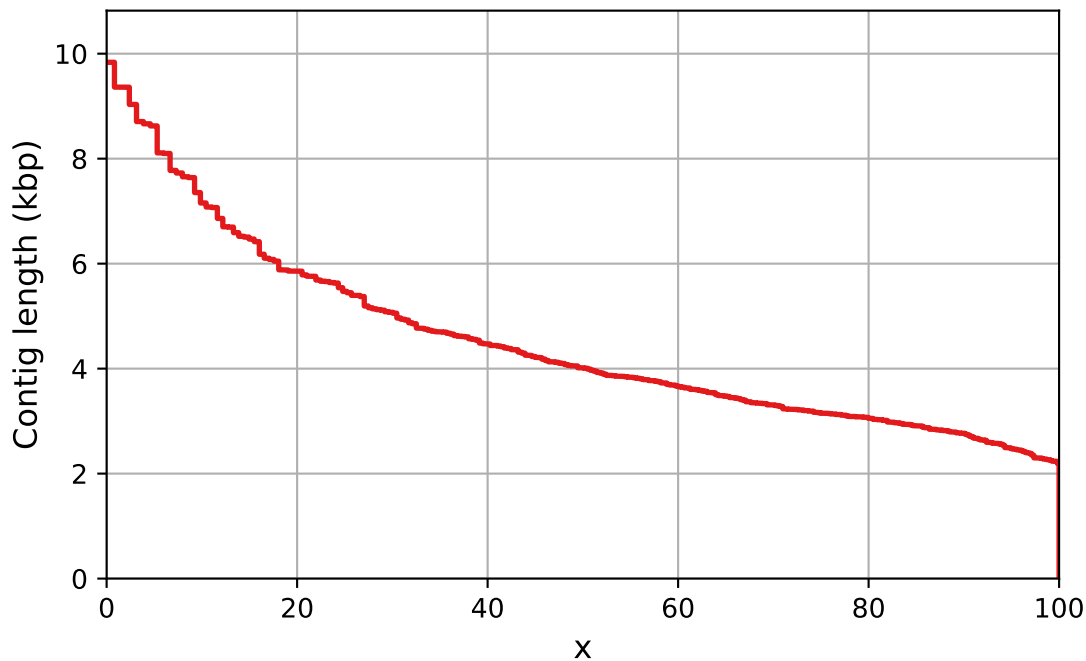


Report

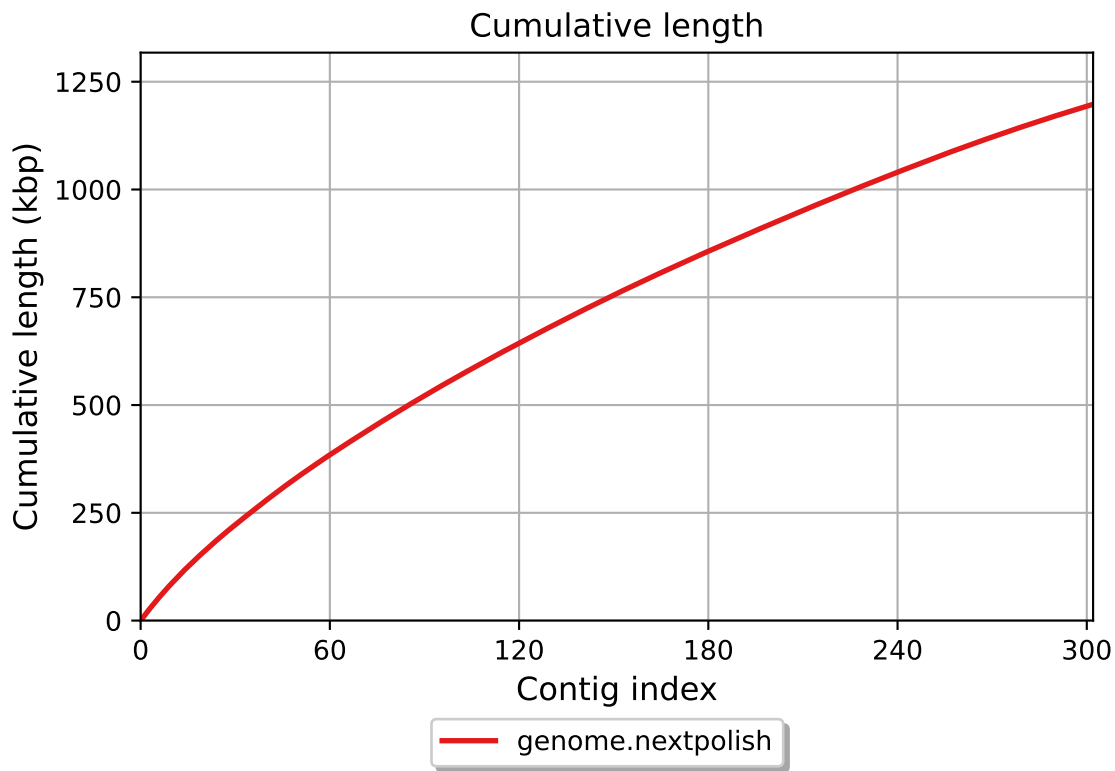
	genome.nextpolish
# contigs (>= 0 bp)	302
# contigs (>= 1000 bp)	302
# contigs (>= 5000 bp)	56
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1197704
Total length (>= 1000 bp)	1197704
Total length (>= 5000 bp)	364909
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	302
Largest contig	9836
Total length	1197704
GC (%)	56.18
N50	4014
N90	2767
auN	4518.8
L50	109
L90	254
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

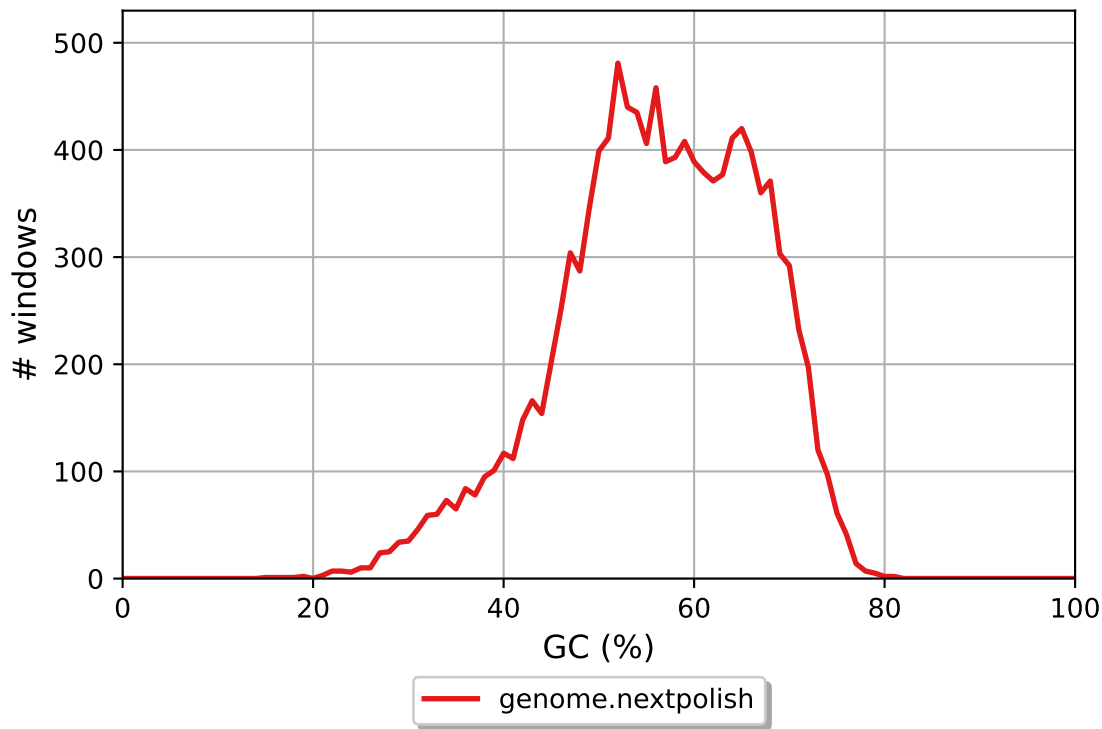
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

