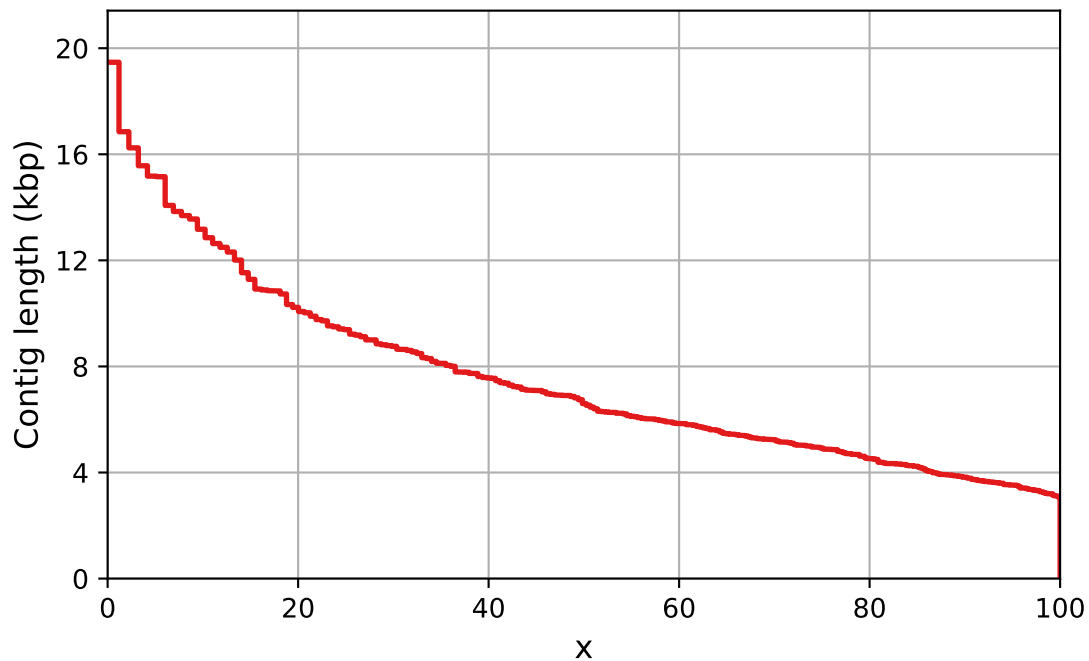


Report

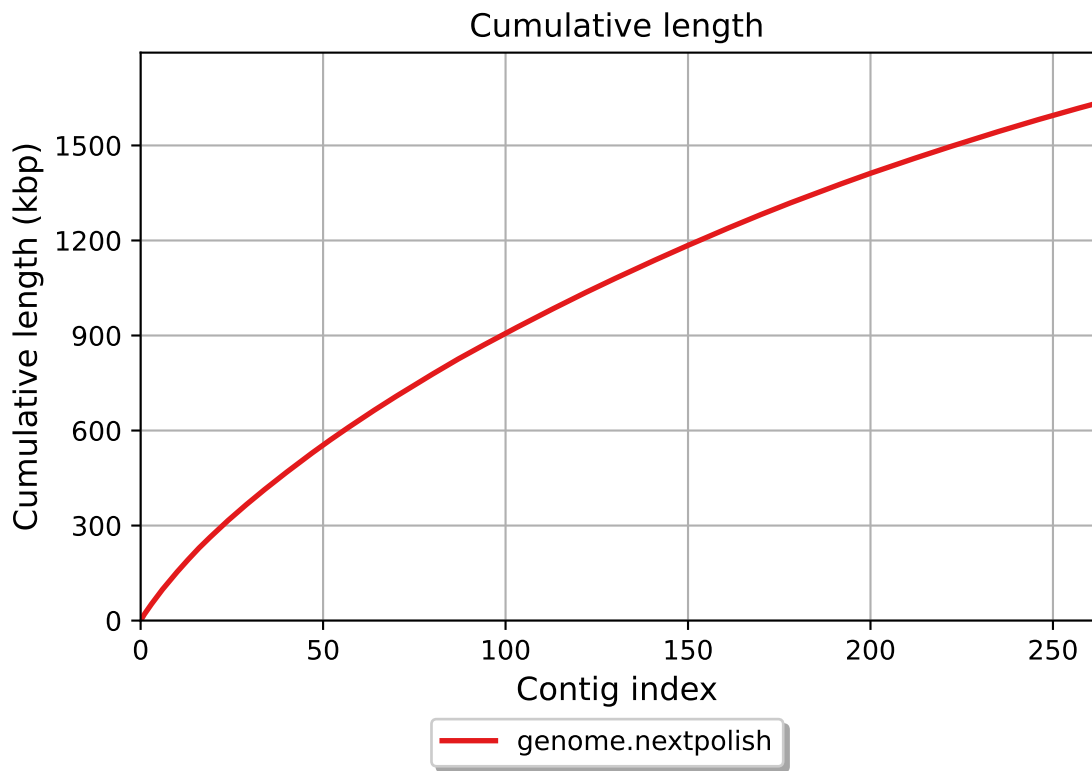
	genome.nextpolish
# contigs (>= 0 bp)	261
# contigs (>= 1000 bp)	261
# contigs (>= 5000 bp)	153
# contigs (>= 10000 bp)	27
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1630001
Total length (>= 1000 bp)	1630001
Total length (>= 5000 bp)	1199647
Total length (>= 10000 bp)	346817
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	261
Largest contig	19470
Total length	1630001
GC (%)	51.35
N50	6599
N90	3822
auN	7578.9
L50	86
L90	215
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

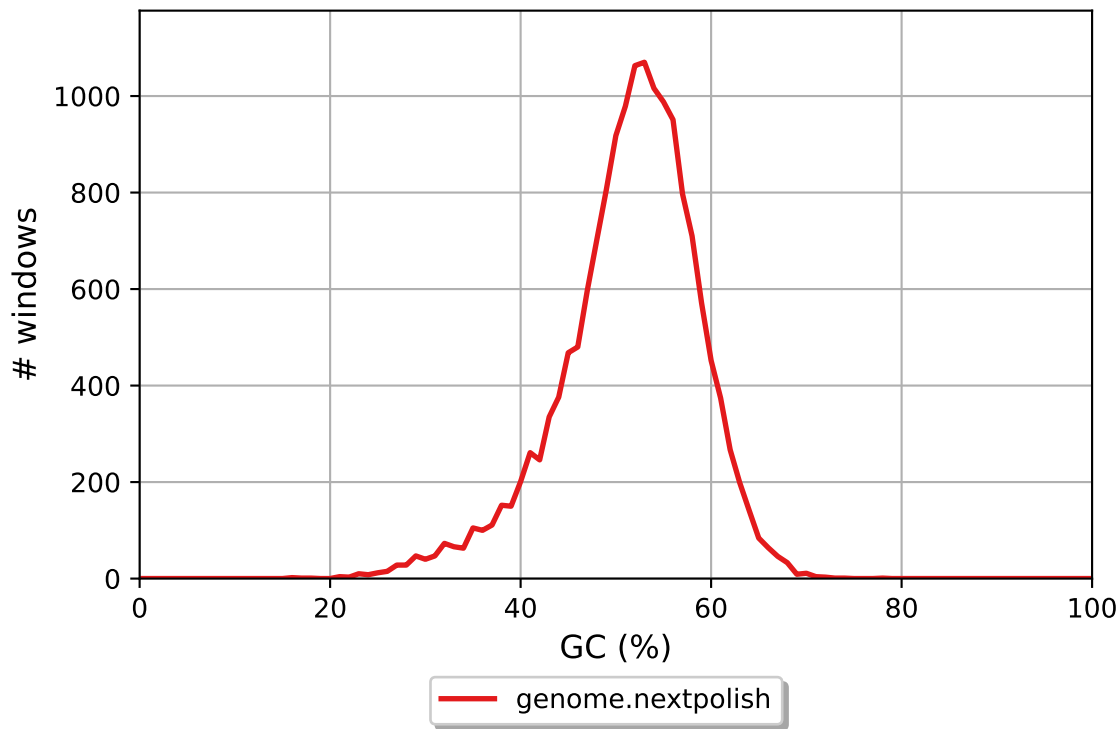
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

