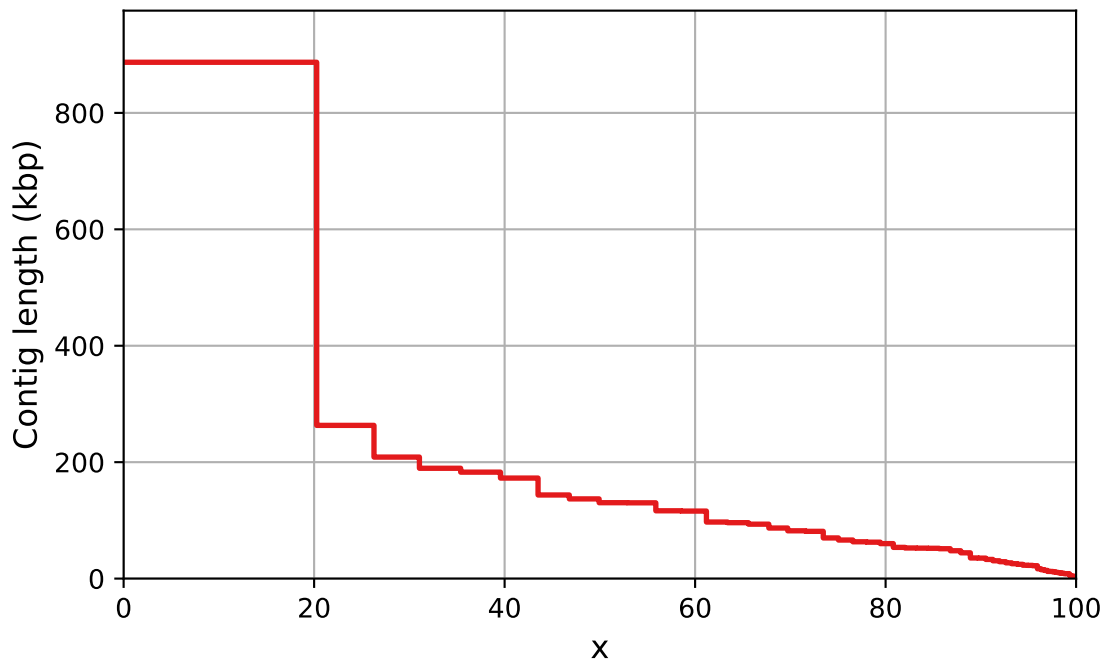


Report

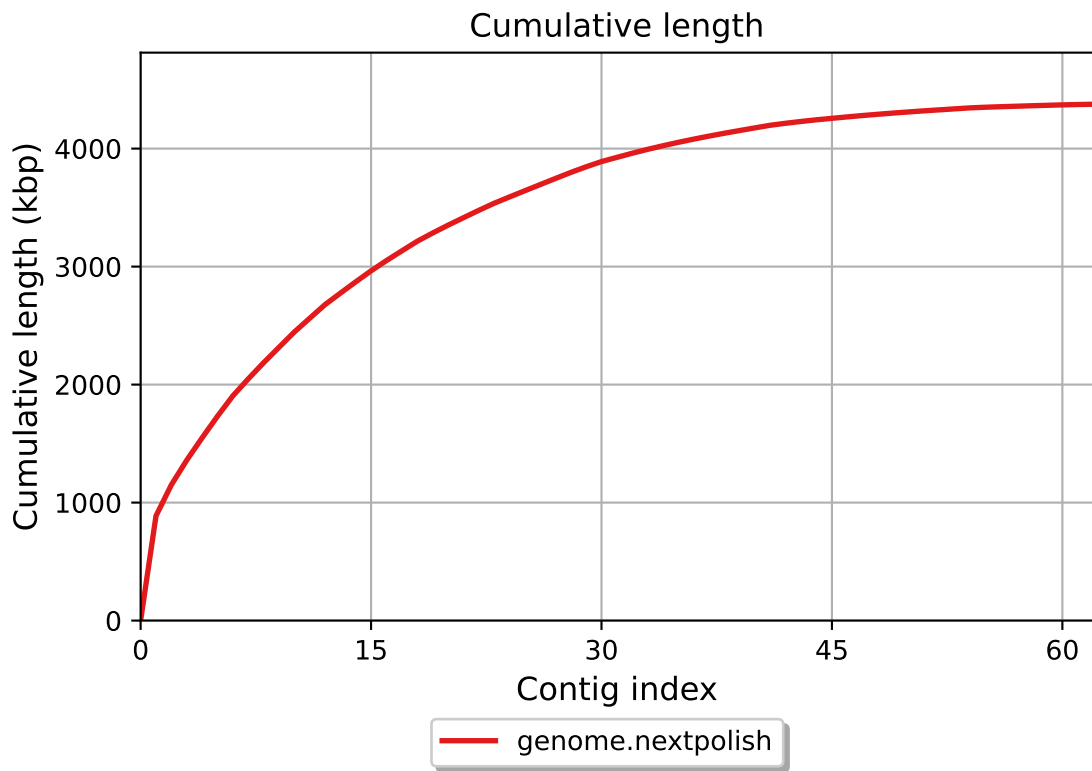
	genome.nextpolish
# contigs (>= 0 bp)	62
# contigs (>= 1000 bp)	62
# contigs (>= 5000 bp)	55
# contigs (>= 10000 bp)	49
# contigs (>= 25000 bp)	37
# contigs (>= 50000 bp)	28
Total length (>= 0 bp)	4375712
Total length (>= 1000 bp)	4375712
Total length (>= 5000 bp)	4350717
Total length (>= 10000 bp)	4301321
Total length (>= 25000 bp)	4105776
Total length (>= 50000 bp)	3797956
# contigs	62
Largest contig	886978
Total length	4375712
GC (%)	45.77
N50	130368
N90	35156
auN	273513.1
L50	9
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 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).

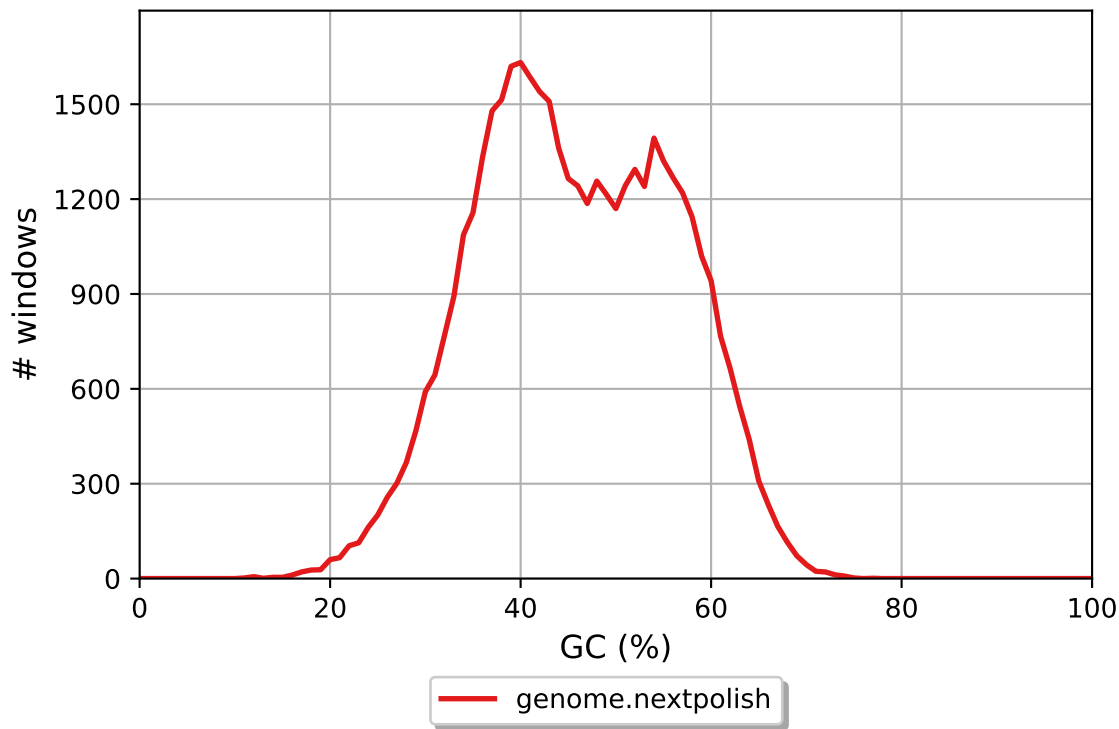
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

