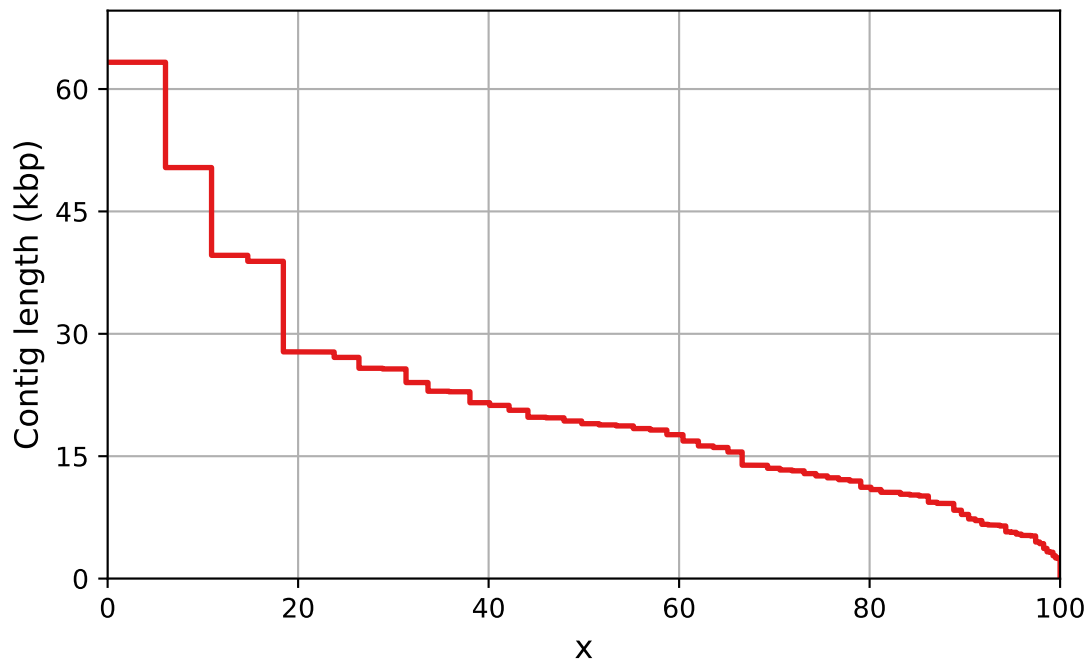


Report

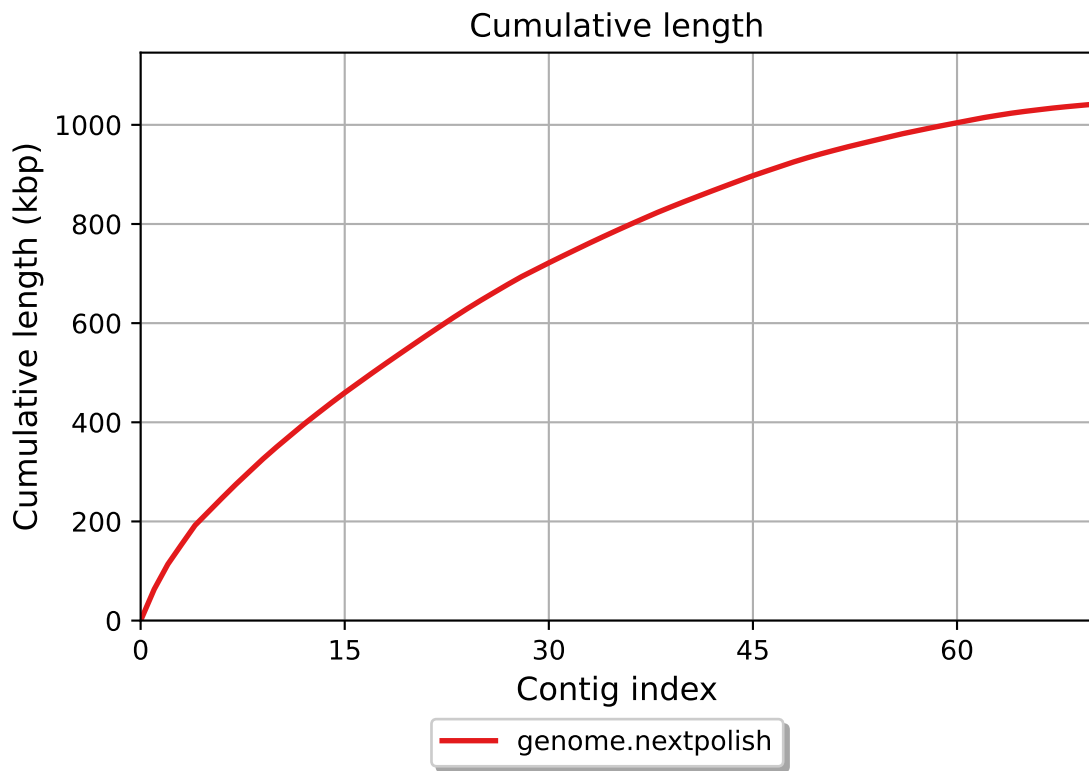
	genome.nextpolish
# contigs (>= 0 bp)	70
# contigs (>= 1000 bp)	70
# contigs (>= 5000 bp)	62
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	9
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	1041517
Total length (>= 1000 bp)	1041517
Total length (>= 5000 bp)	1014758
Total length (>= 10000 bp)	897464
Total length (>= 25000 bp)	326297
Total length (>= 50000 bp)	113654
# contigs	70
Largest contig	63280
Total length	1041517
GC (%)	54.29
N50	18983
N90	7859
auN	22895.4
L50	19
L90	50
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

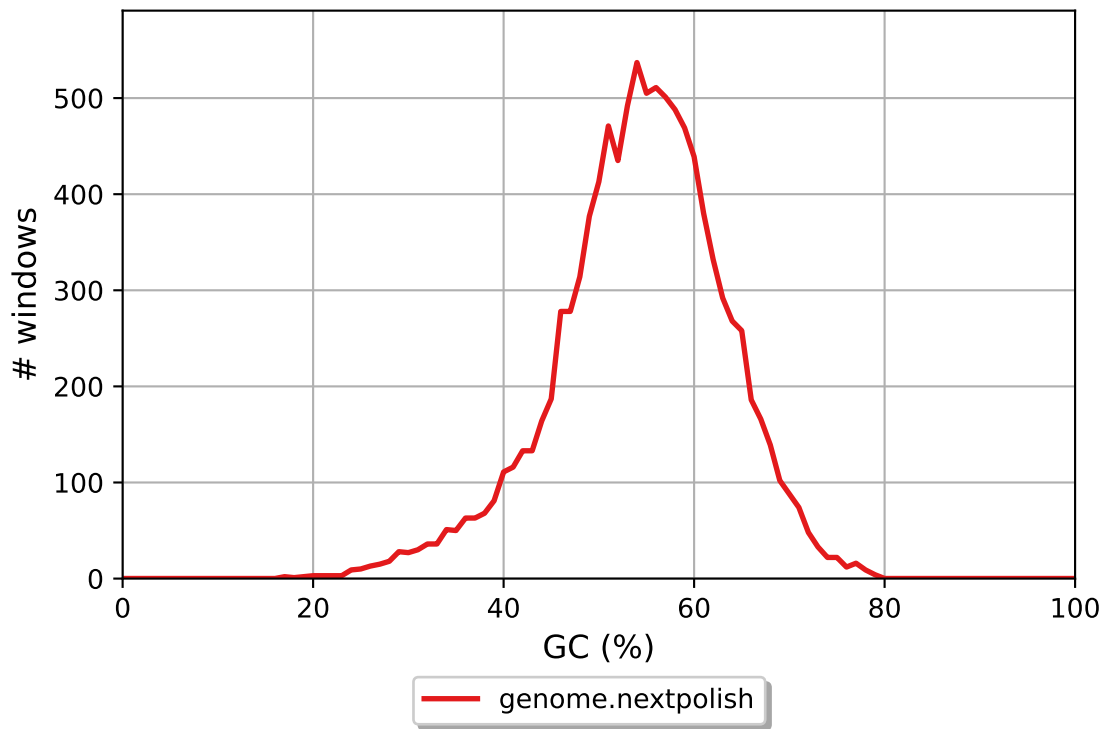
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

