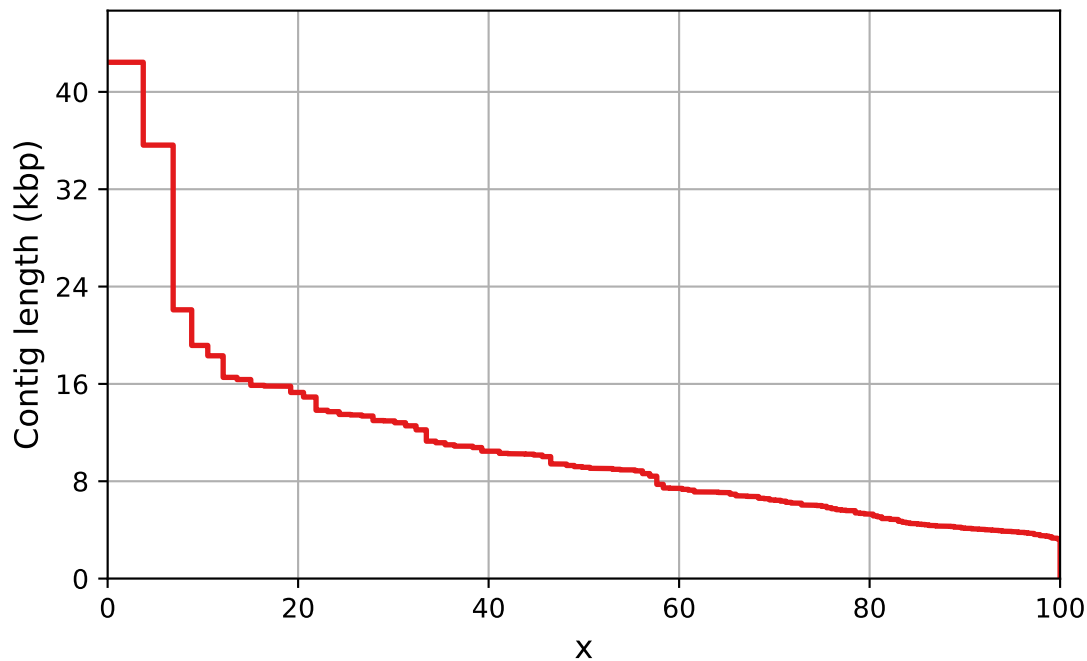


Report

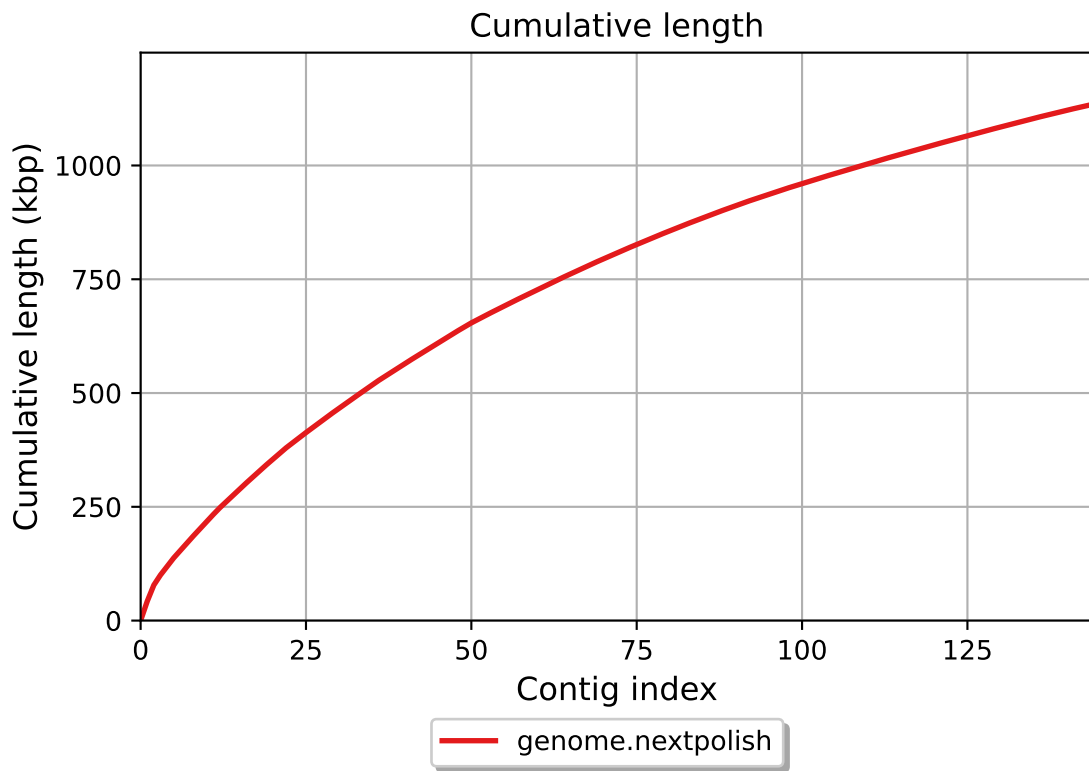
	genome.nextpolish
# contigs (>= 0 bp)	144
# contigs (>= 1000 bp)	144
# contigs (>= 5000 bp)	92
# contigs (>= 10000 bp)	36
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1134508
Total length (>= 1000 bp)	1134508
Total length (>= 5000 bp)	922117
Total length (>= 10000 bp)	527755
Total length (>= 25000 bp)	78058
Total length (>= 50000 bp)	0
# contigs	144
Largest contig	42433
Total length	1134508
GC (%)	56.92
N50	9144
N90	4131
auN	11510.9
L50	41
L90	115
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

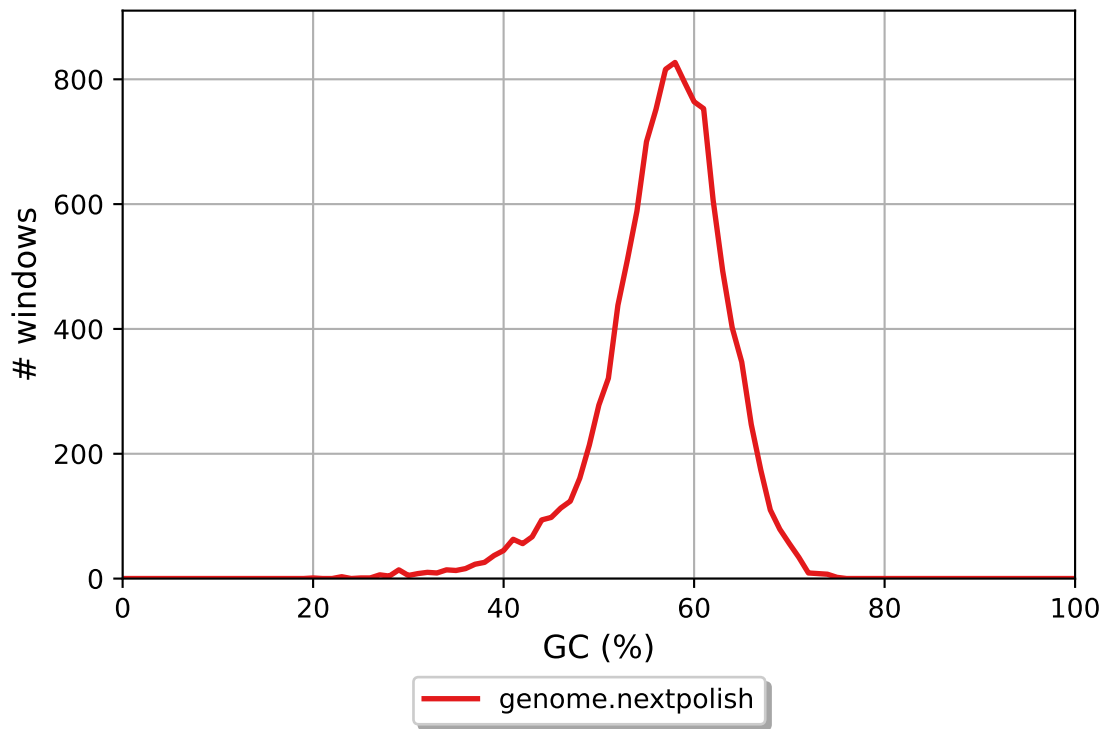
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

