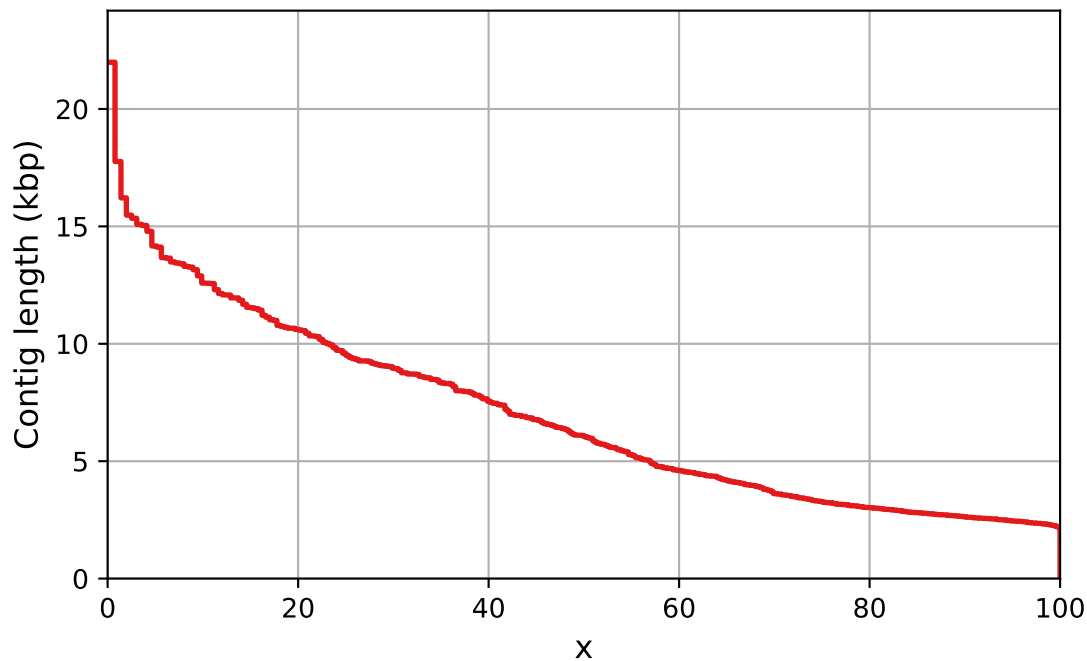


Report

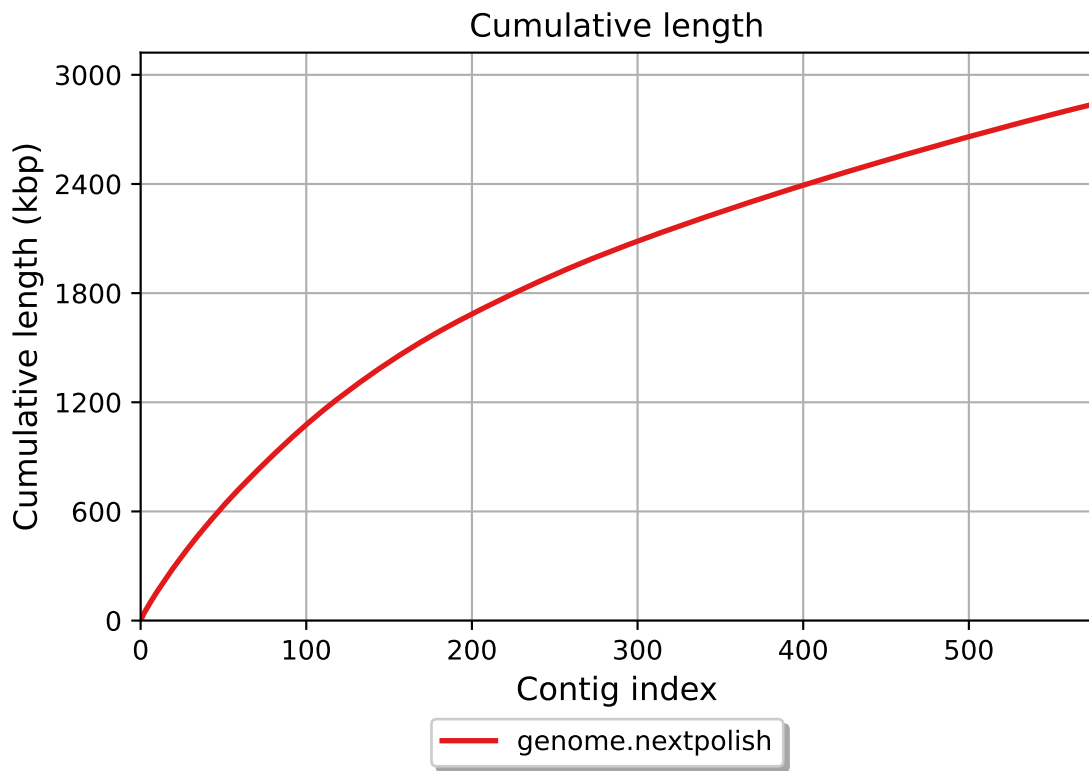
	genome.nextpolish
# contigs (>= 0 bp)	575
# contigs (>= 1000 bp)	575
# contigs (>= 5000 bp)	186
# contigs (>= 10000 bp)	53
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2838169
Total length (>= 1000 bp)	2838169
Total length (>= 5000 bp)	1618581
Total length (>= 10000 bp)	661378
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	575
Largest contig	21989
Total length	2838169
GC (%)	56.01
N50	6082
N90	2633
auN	6937.2
L50	150
L90	459
# 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).

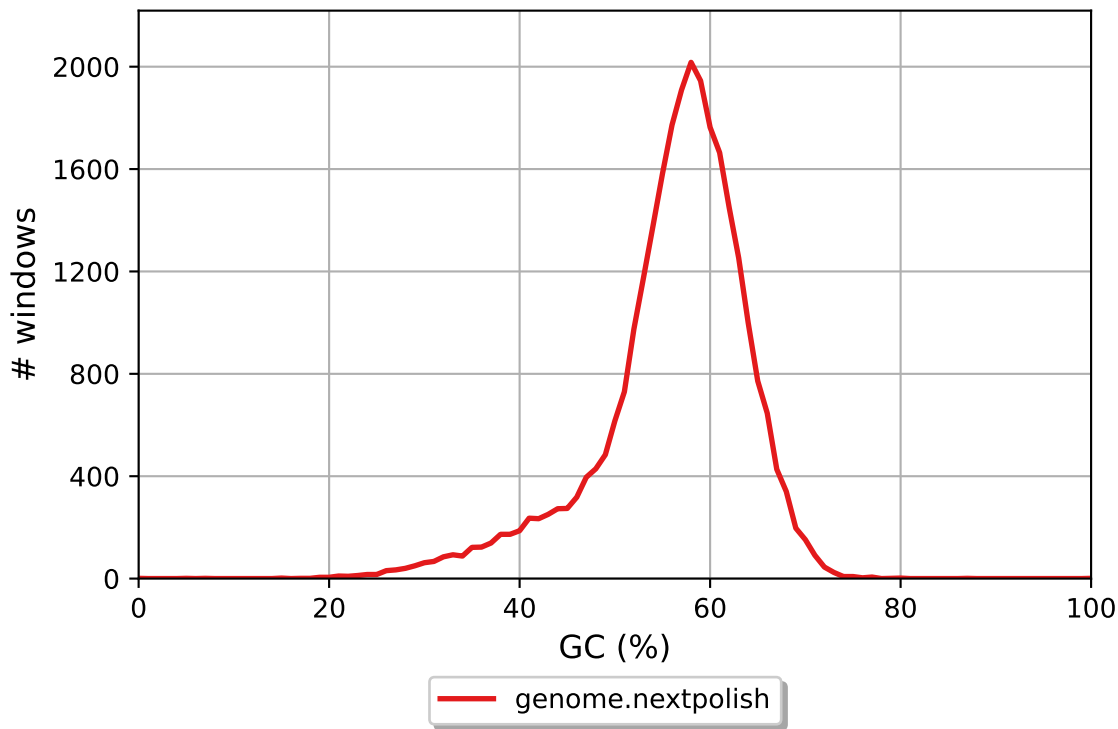
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

