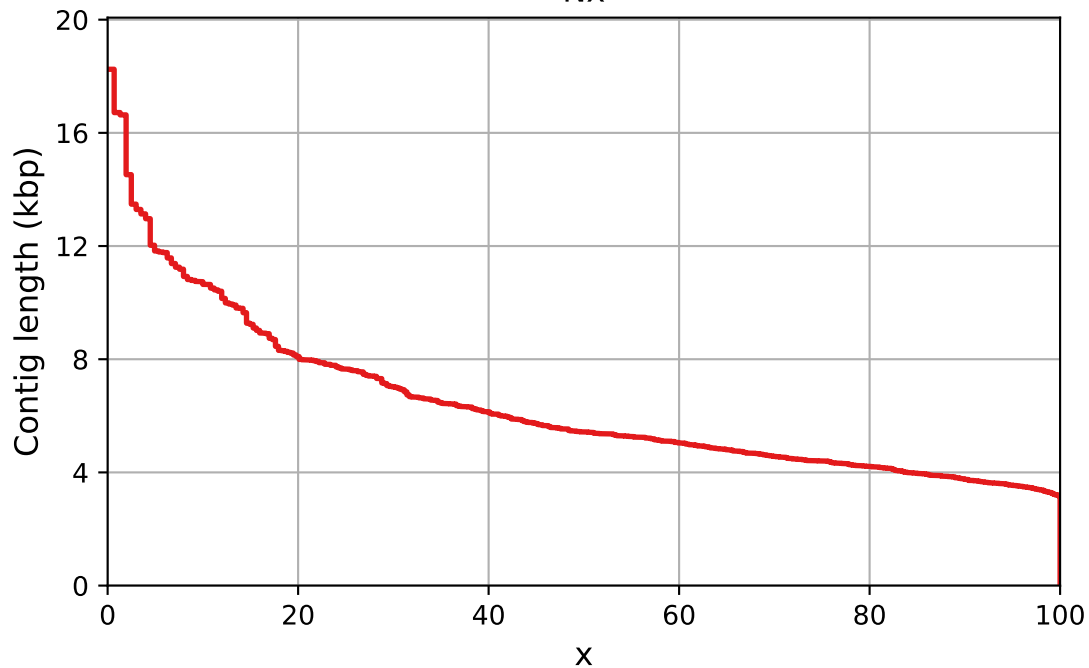


Report

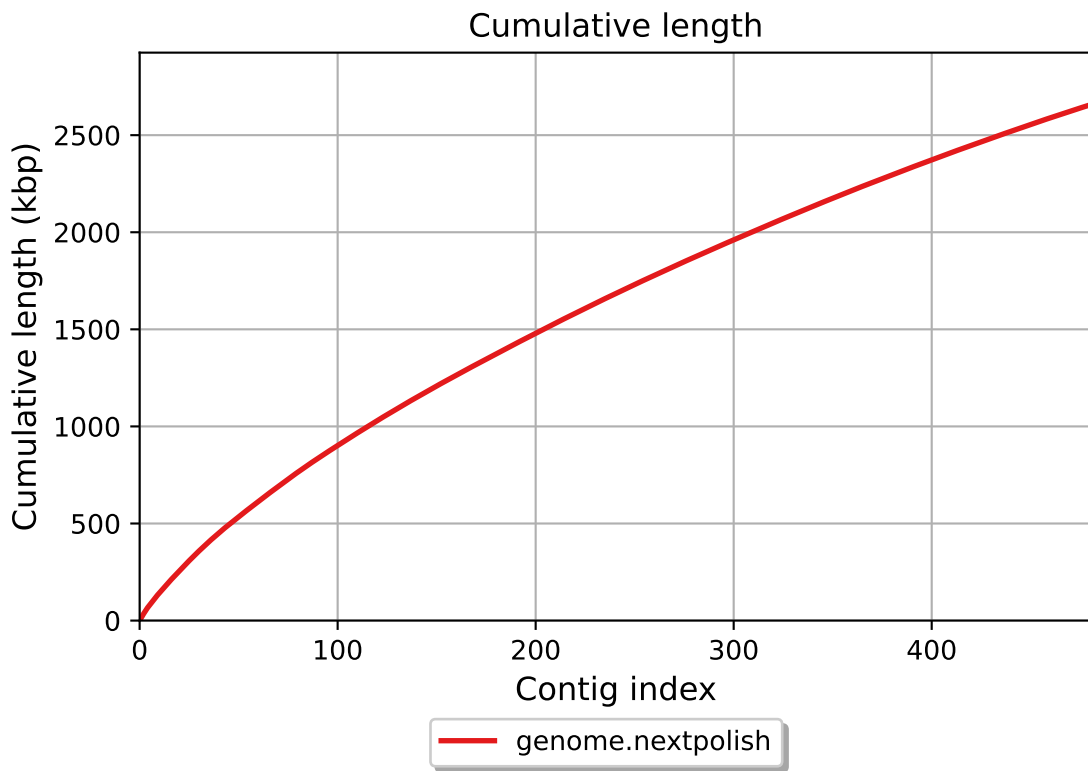
	genome.nextpolish
# contigs (>= 0 bp)	481
# contigs (>= 1000 bp)	481
# contigs (>= 5000 bp)	227
# contigs (>= 10000 bp)	27
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2659080
Total length (>= 1000 bp)	2659080
Total length (>= 5000 bp)	1618149
Total length (>= 10000 bp)	328625
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	481
Largest contig	18247
Total length	2659080
GC (%)	54.77
N50	5430
N90	3755
auN	6419.9
L50	172
L90	406
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

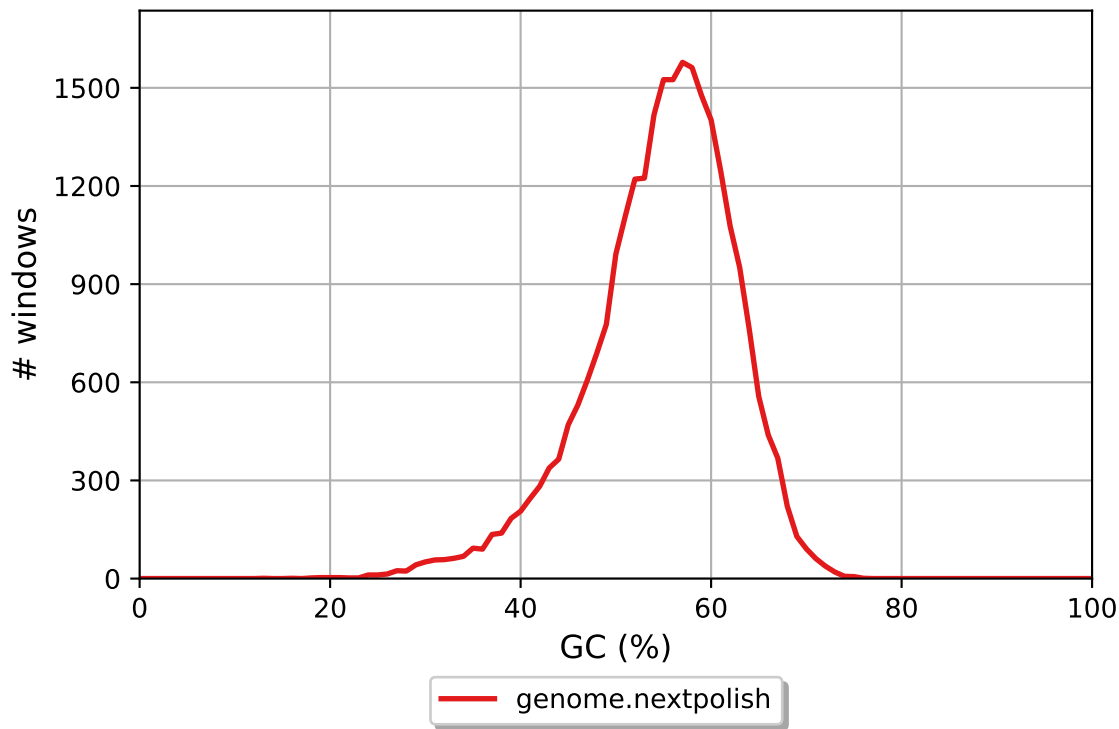
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

