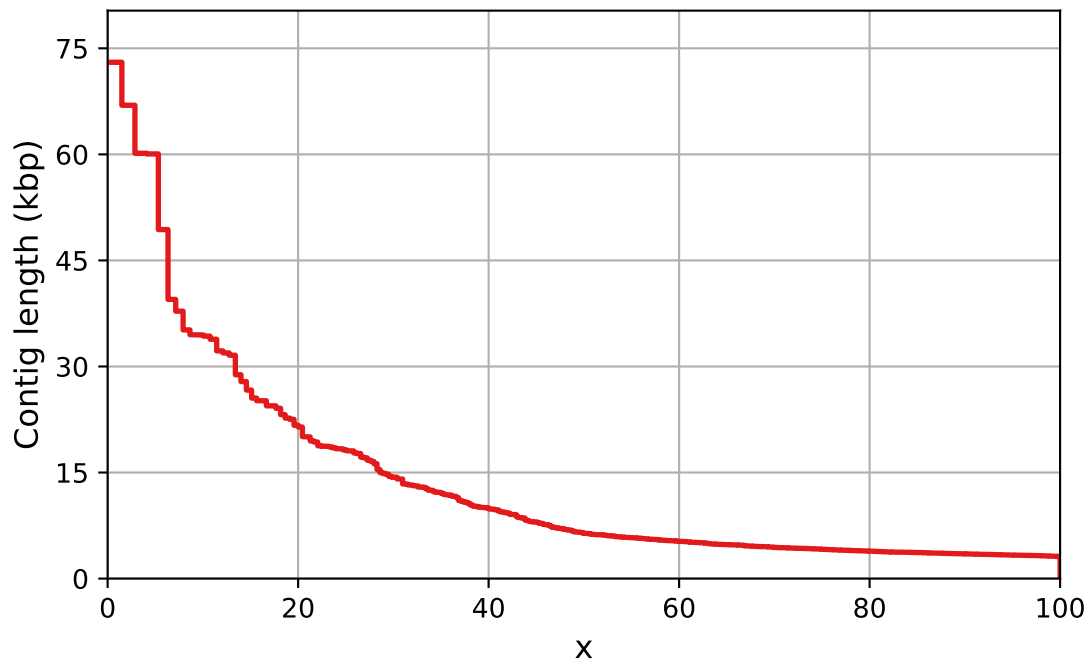


Report

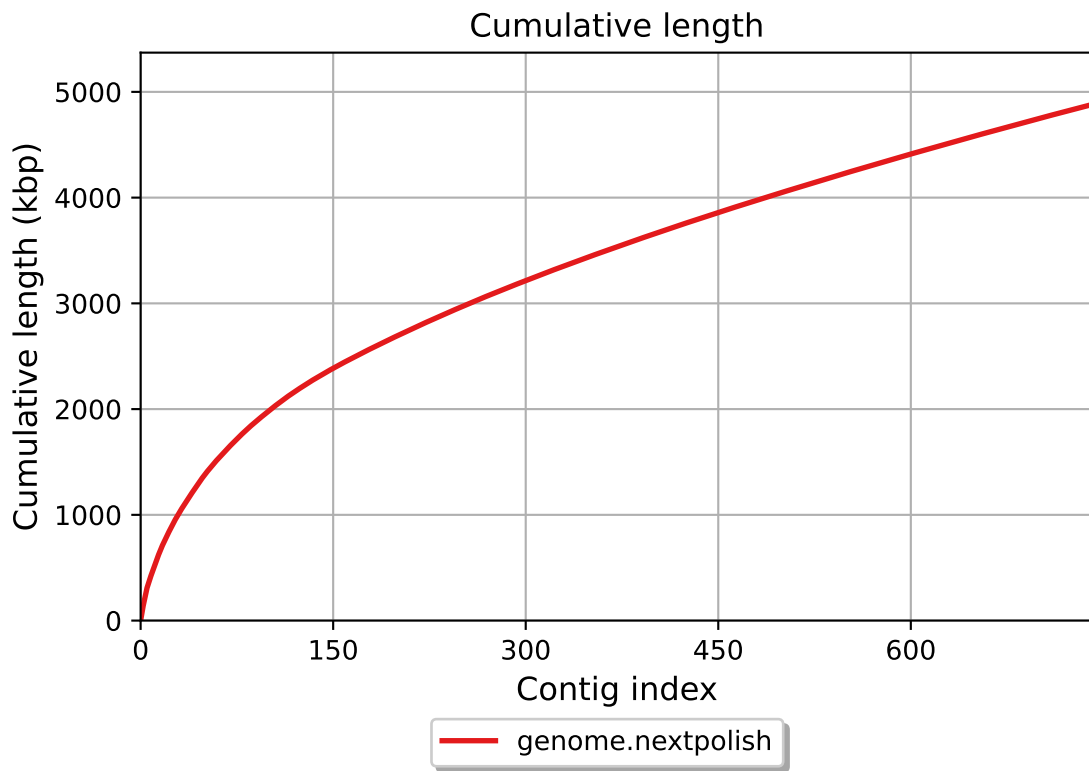
	genome.nextpolish
# contigs (>= 0 bp)	742
# contigs (>= 1000 bp)	742
# contigs (>= 5000 bp)	269
# contigs (>= 10000 bp)	96
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	4882814
Total length (>= 1000 bp)	4882814
Total length (>= 5000 bp)	3064851
Total length (>= 10000 bp)	1945259
Total length (>= 25000 bp)	814030
Total length (>= 50000 bp)	260188
# contigs	742
Largest contig	73026
Total length	4882814
GC (%)	53.55
N50	6408
N90	3497
auN	14055.0
L50	159
L90	595
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

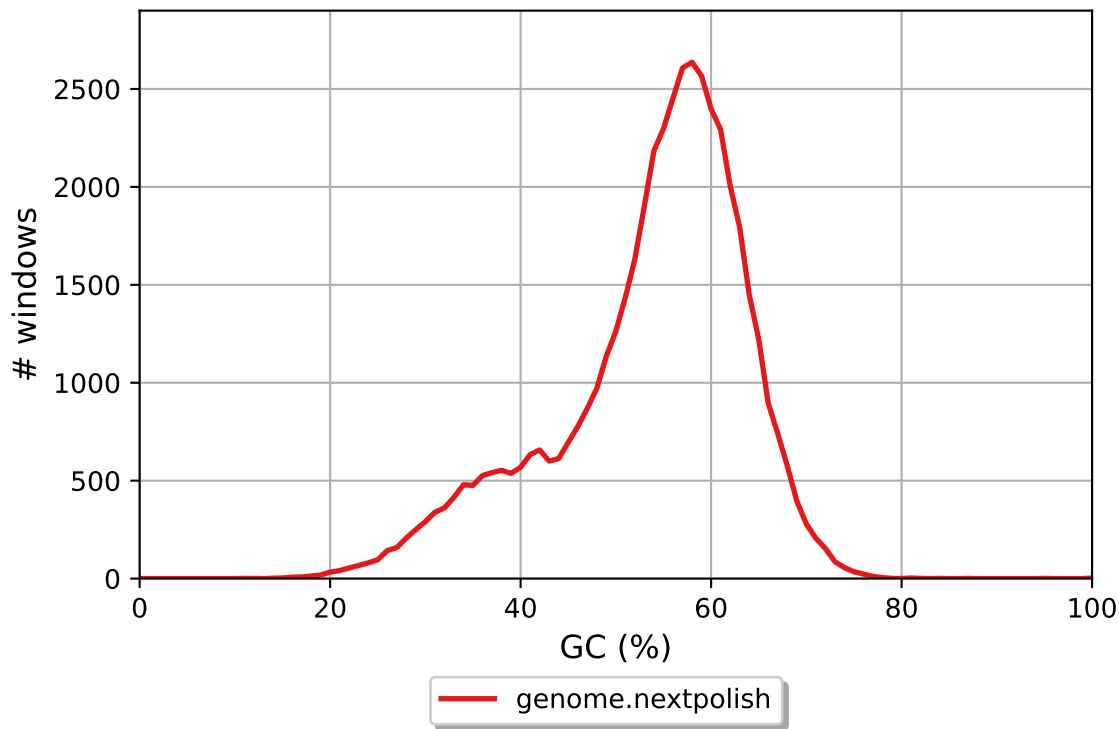
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

