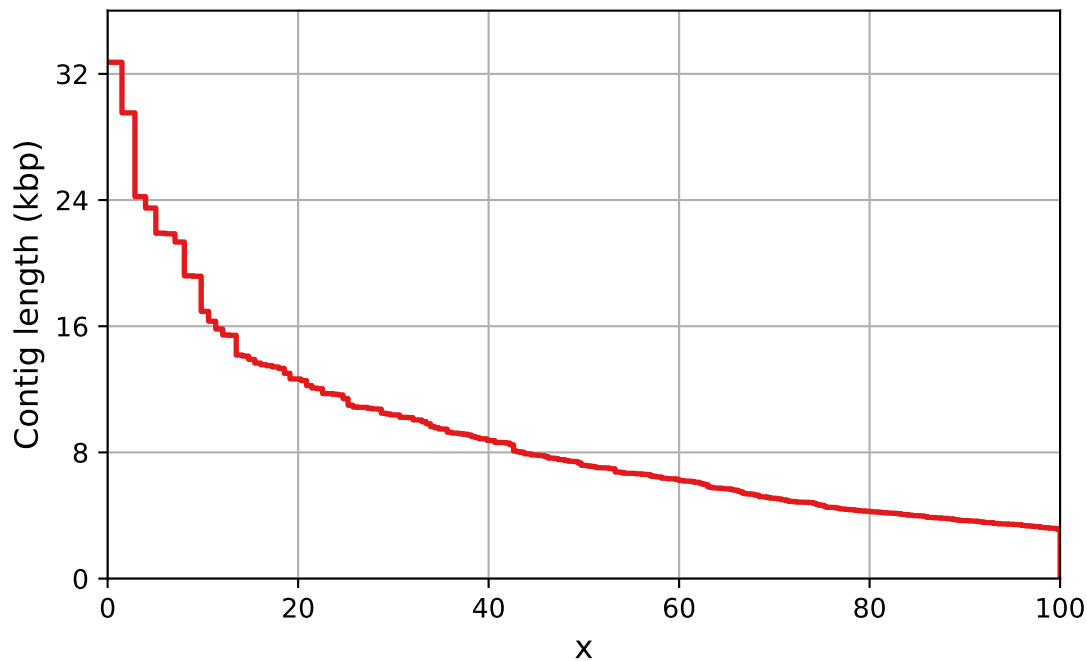


Report

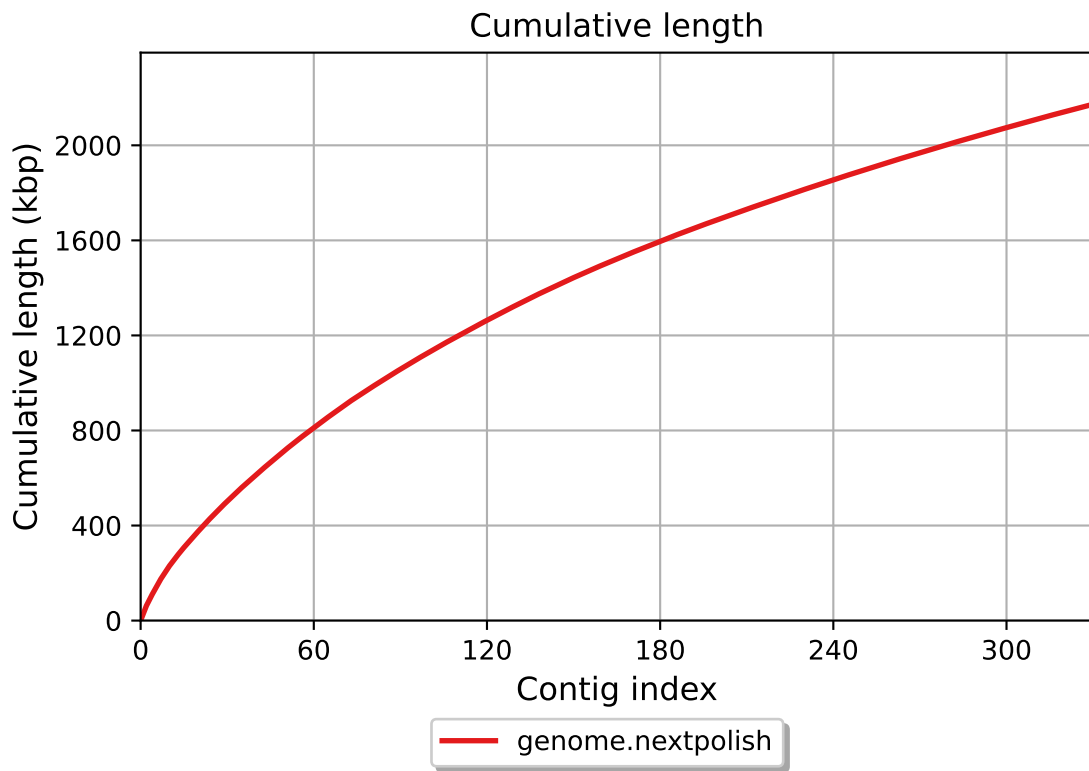
	genome.nextpolish
# contigs (>= 0 bp)	330
# contigs (>= 1000 bp)	330
# contigs (>= 5000 bp)	168
# contigs (>= 10000 bp)	50
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2172757
Total length (>= 1000 bp)	2172757
Total length (>= 5000 bp)	1537227
Total length (>= 10000 bp)	716791
Total length (>= 25000 bp)	62262
Total length (>= 50000 bp)	0
# contigs	330
Largest contig	32735
Total length	2172757
GC (%)	57.35
N50	7187
N90	3679
auN	9204.6
L50	94
L90	267
# 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).

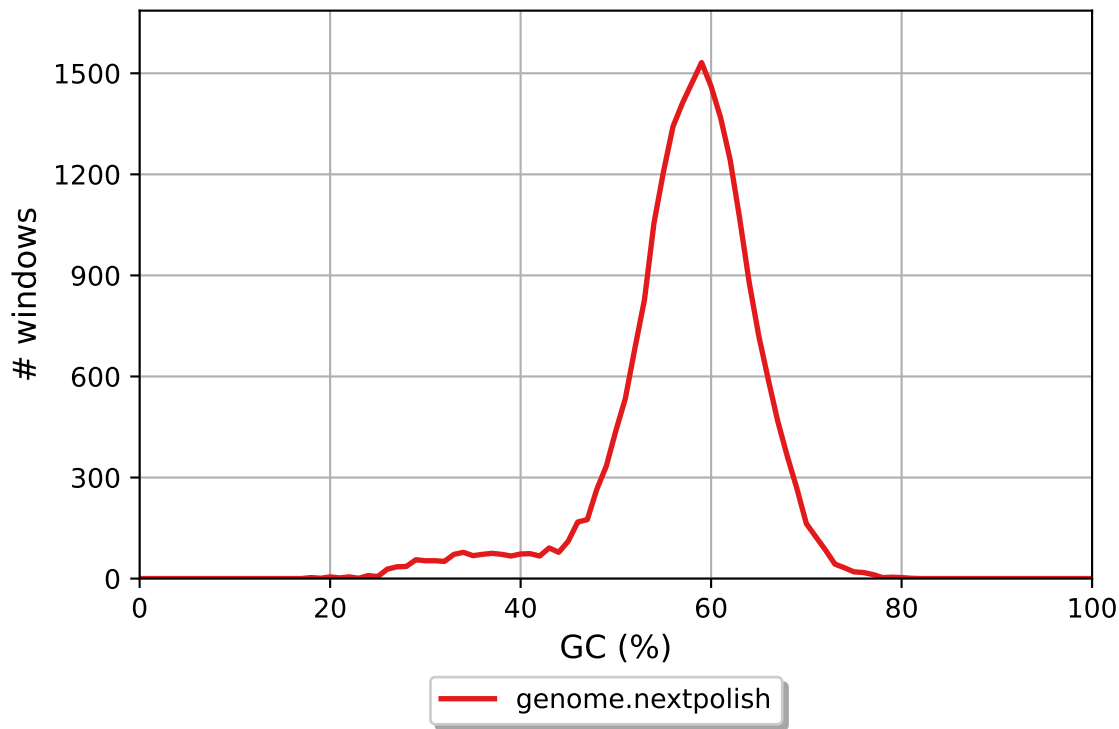
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

