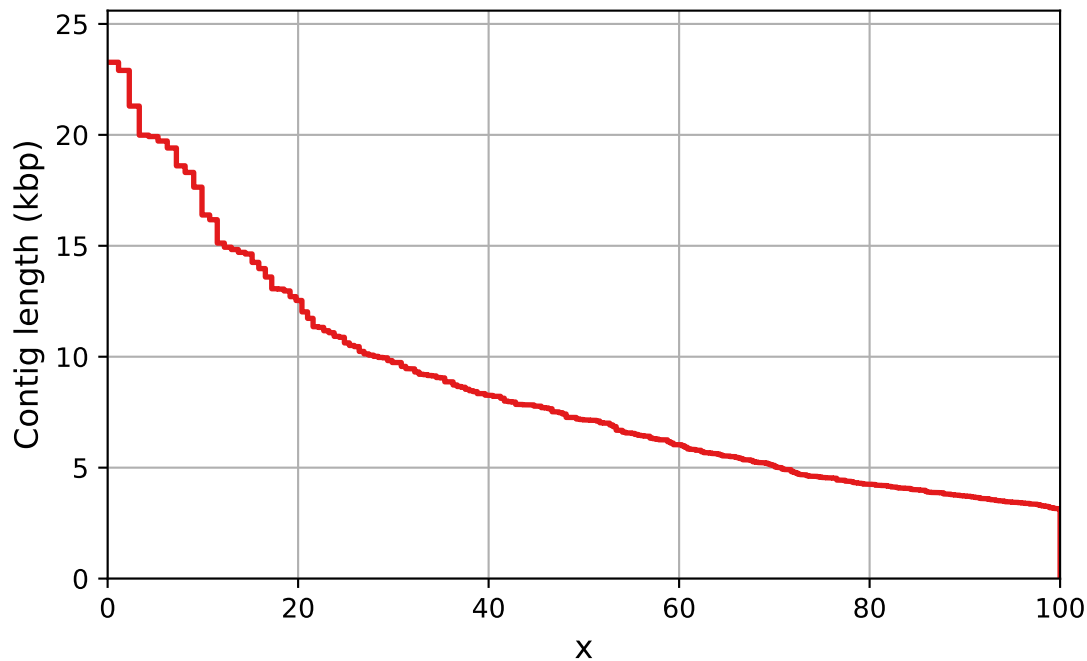


Report

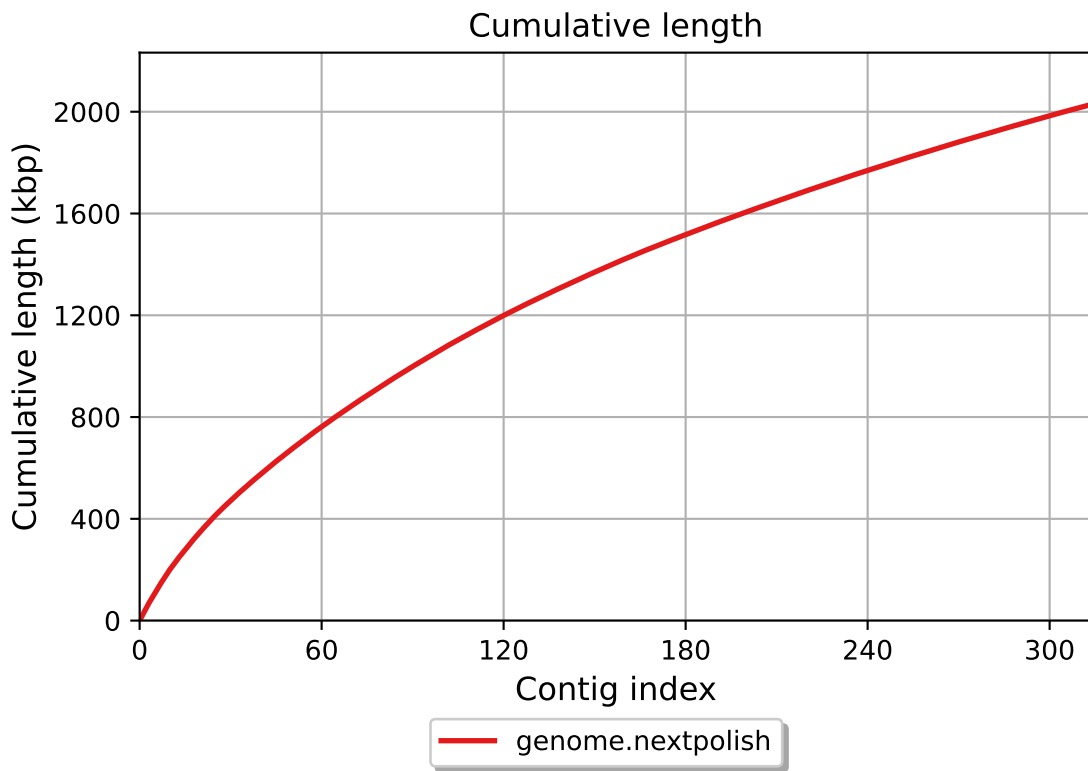
	genome.nextpolish
# contigs (>= 0 bp)	314
# contigs (>= 1000 bp)	314
# contigs (>= 5000 bp)	162
# contigs (>= 10000 bp)	40
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2029374
Total length (>= 1000 bp)	2029374
Total length (>= 5000 bp)	1431692
Total length (>= 10000 bp)	576500
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	314
Largest contig	23271
Total length	2029374
GC (%)	57.89
N50	7153
N90	3718
auN	8622.6
L50	93
L90	256
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 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).

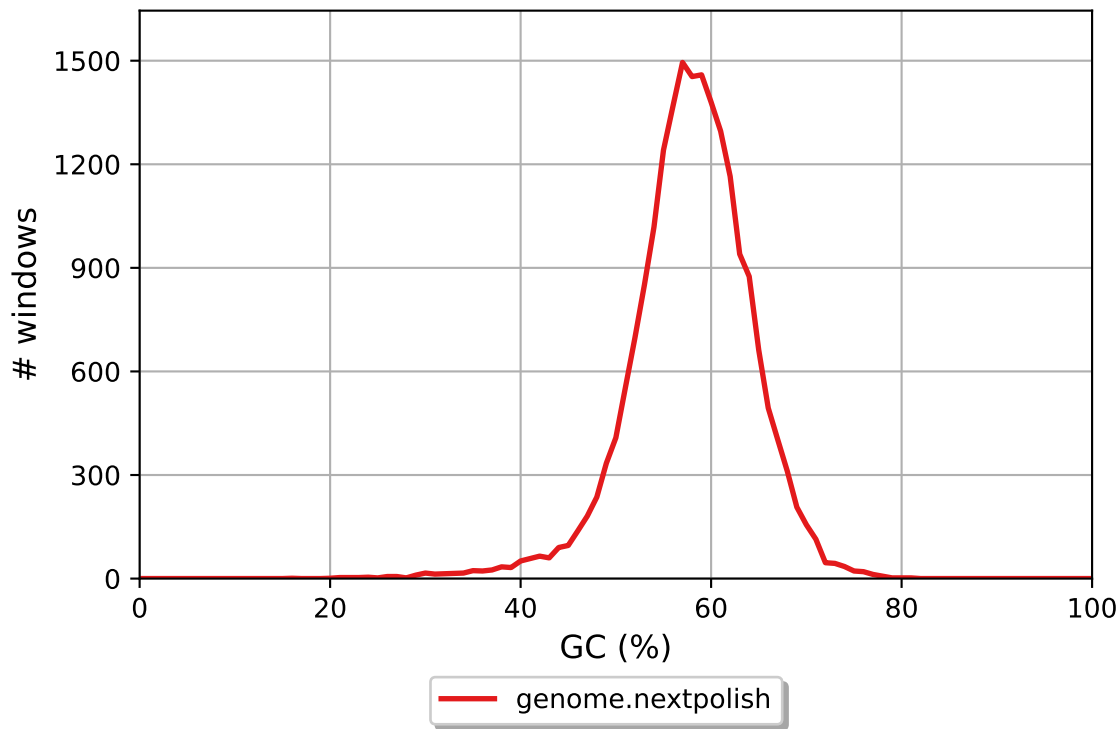
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

