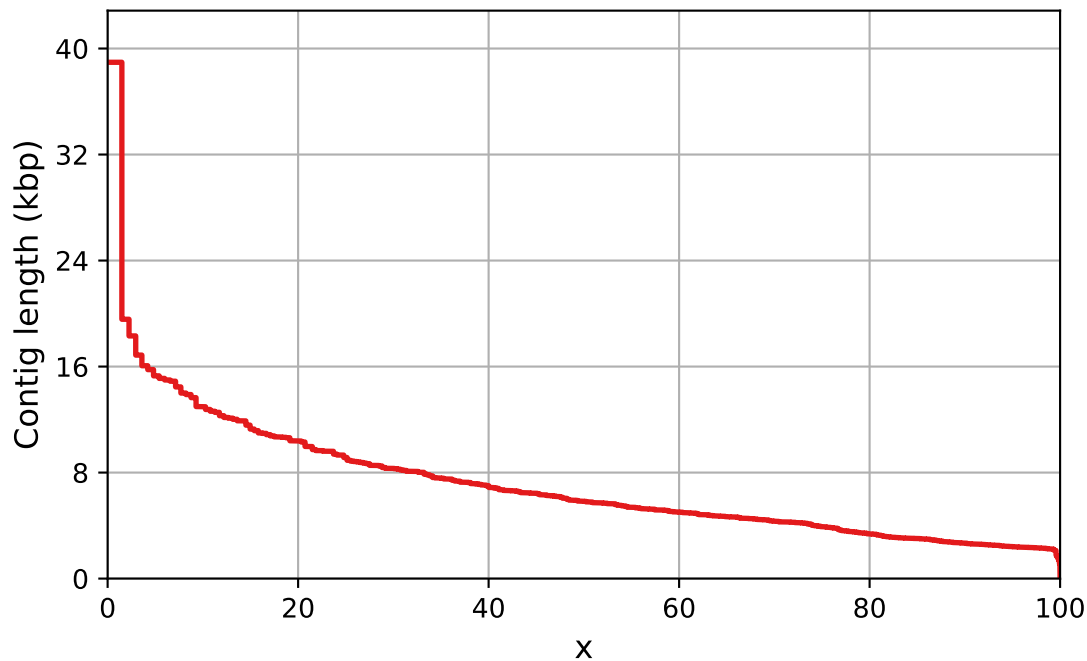


Report

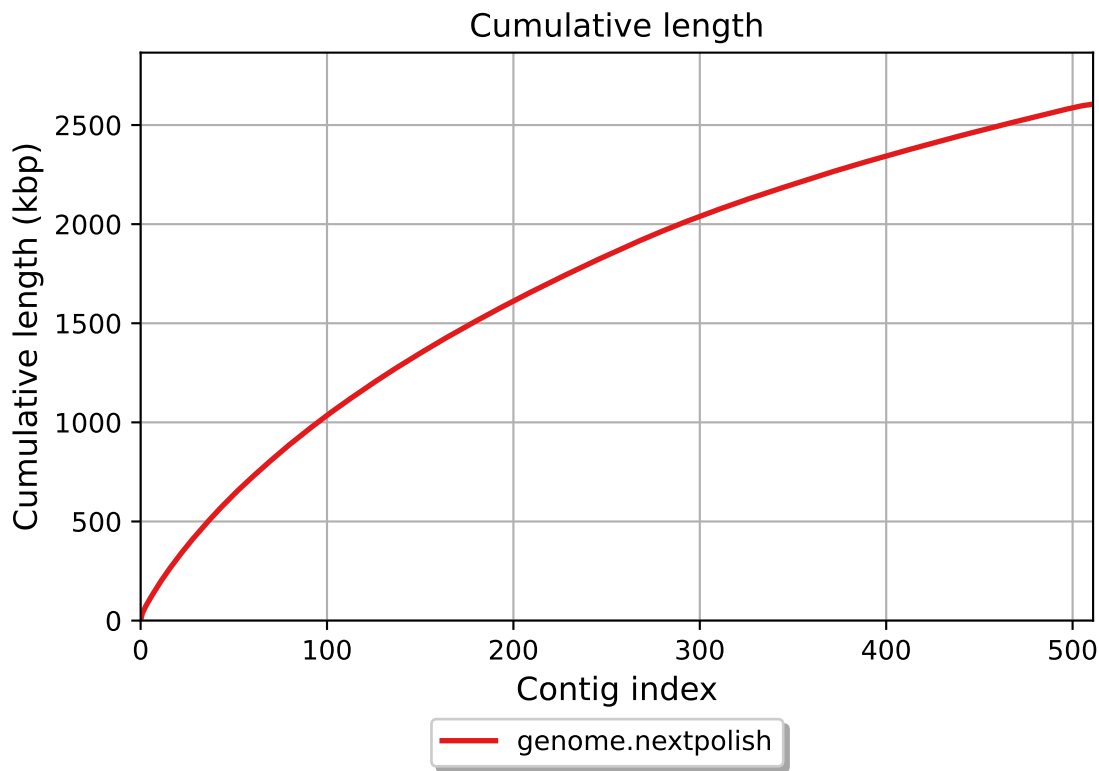
	genome.nextpolish
# contigs (>= 0 bp)	511
# contigs (>= 1000 bp)	510
# contigs (>= 5000 bp)	192
# contigs (>= 10000 bp)	40
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2604803
Total length (>= 1000 bp)	2604222
Total length (>= 5000 bp)	1572542
Total length (>= 10000 bp)	539851
Total length (>= 25000 bp)	38962
Total length (>= 50000 bp)	0
# contigs	511
Largest contig	38962
Total length	2604803
GC (%)	59.11
N50	5831
N90	2670
auN	7313.3
L50	142
L90	401
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

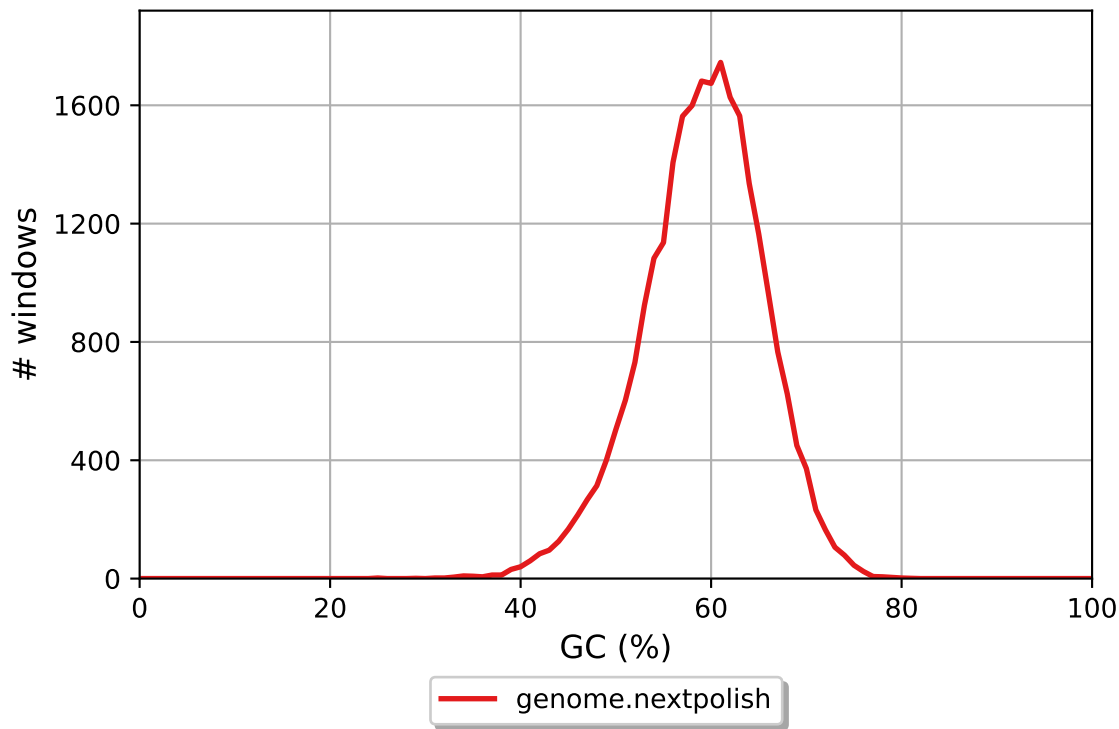
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

