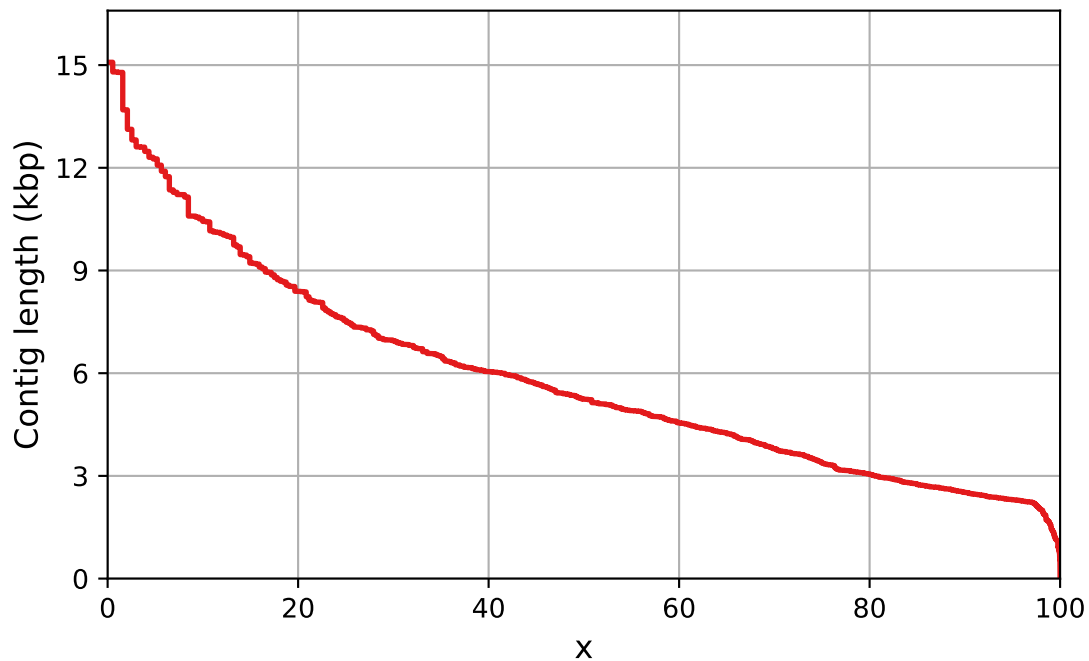


Report

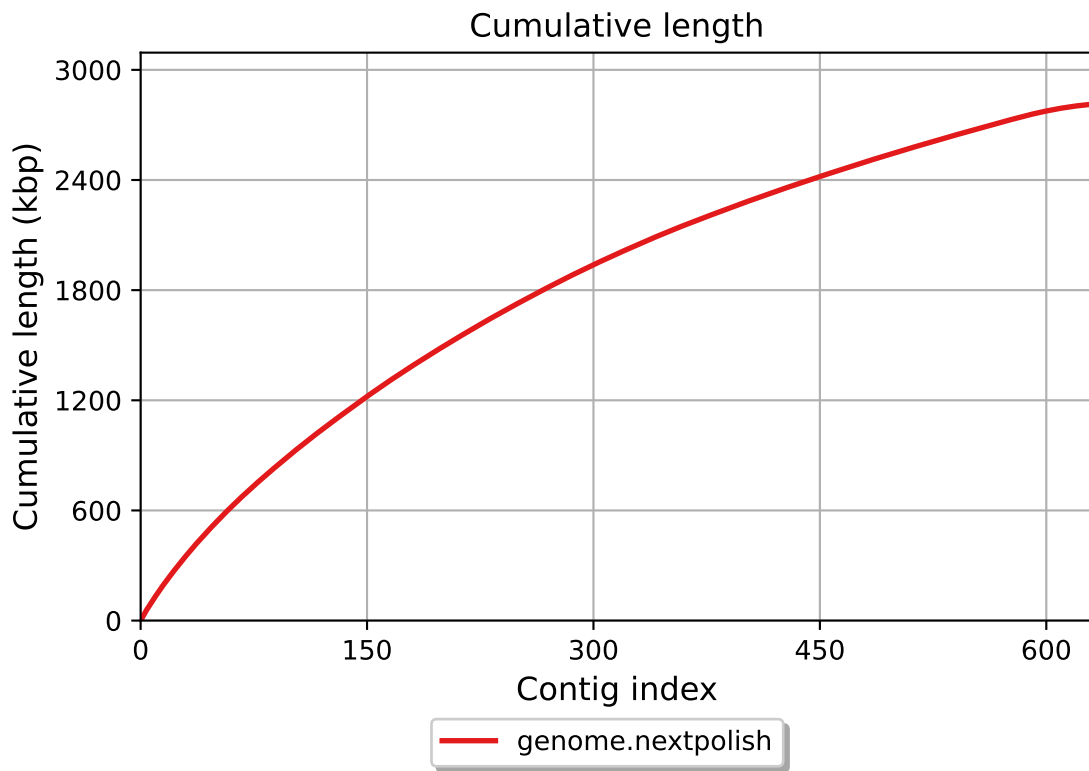
	genome.nextpolish
# contigs (>= 0 bp)	631
# contigs (>= 1000 bp)	622
# contigs (>= 5000 bp)	202
# contigs (>= 10000 bp)	30
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2812530
Total length (>= 1000 bp)	2805777
Total length (>= 5000 bp)	1500990
Total length (>= 10000 bp)	352101
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	631
Largest contig	15082
Total length	2812530
GC (%)	56.27
N50	5242
N90	2519
auN	5892.2
L50	184
L90	493
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

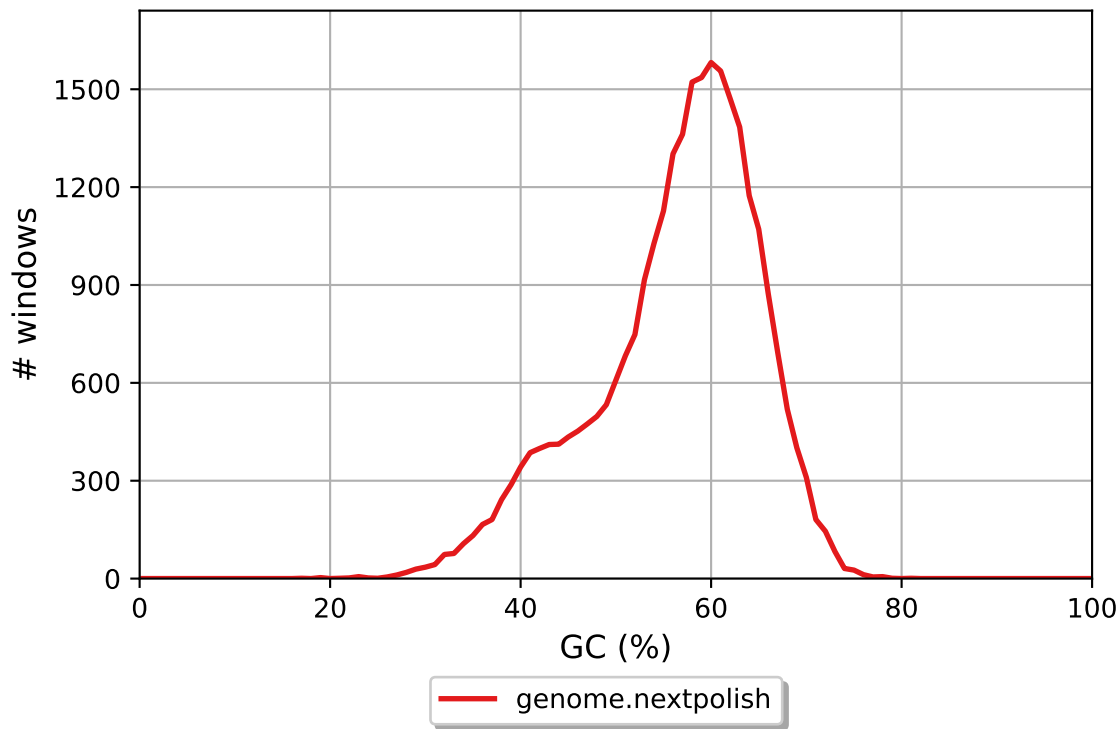
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

