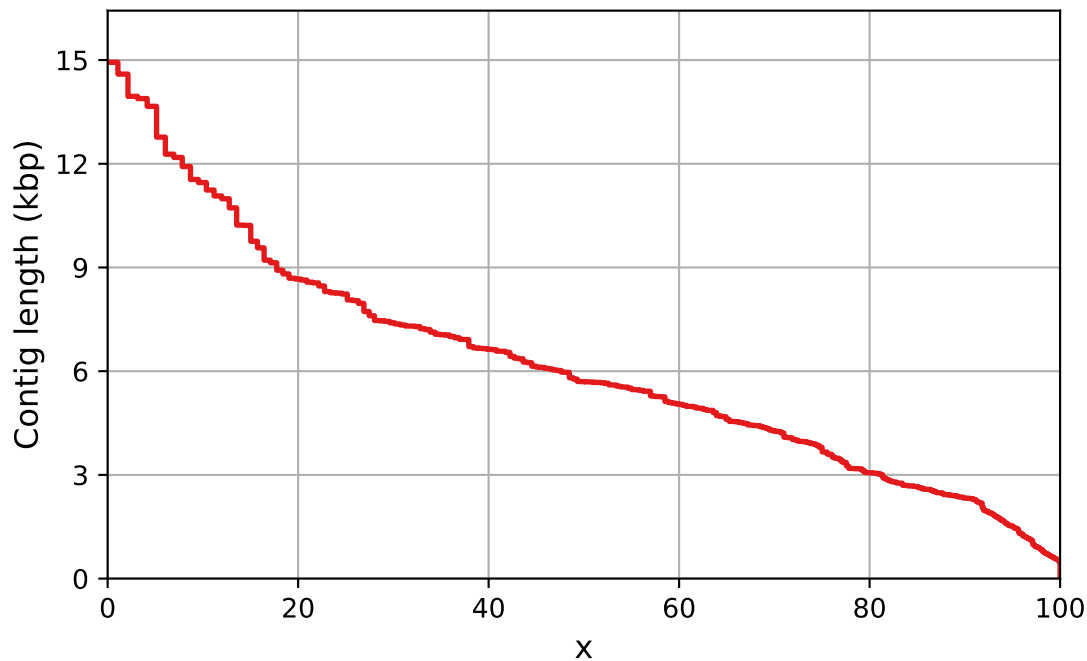


Report

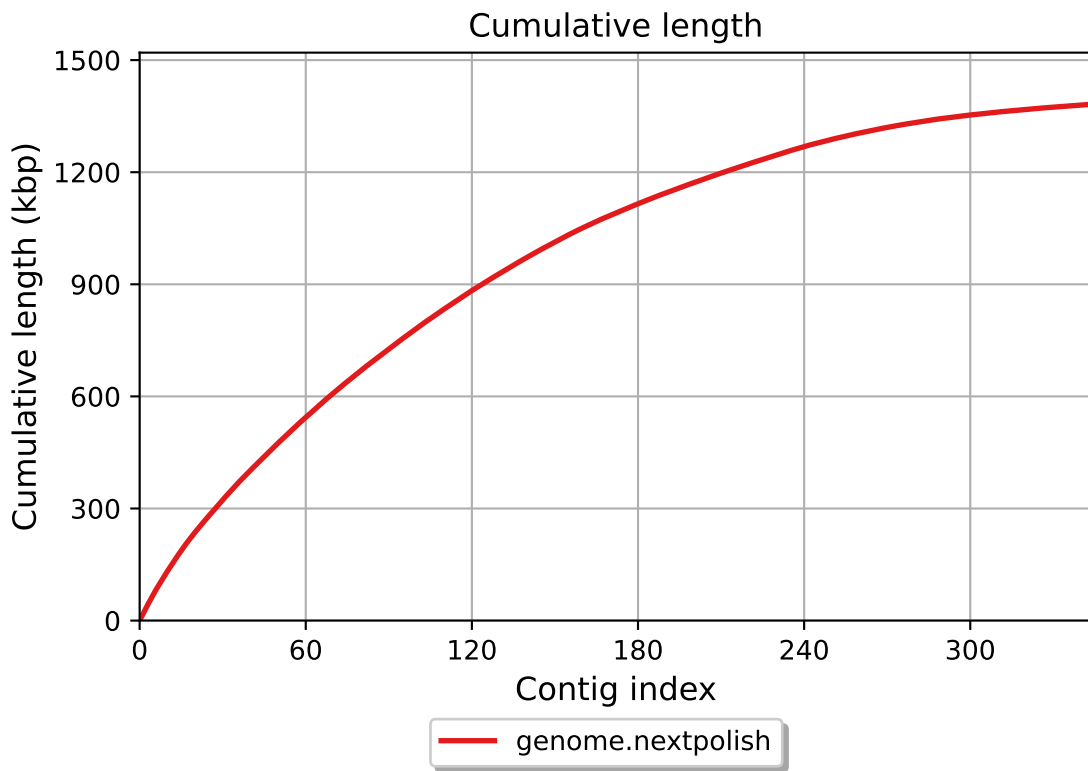
	genome.nextpolish
# contigs (>= 0 bp)	344
# contigs (>= 1000 bp)	288
# contigs (>= 5000 bp)	111
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1381810
Total length (>= 1000 bp)	1341963
Total length (>= 5000 bp)	838903
Total length (>= 10000 bp)	207652
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	344
Largest contig	14936
Total length	1381810
GC (%)	61.59
N50	5694
N90	2329
auN	6240.6
L50	84
L90	229
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

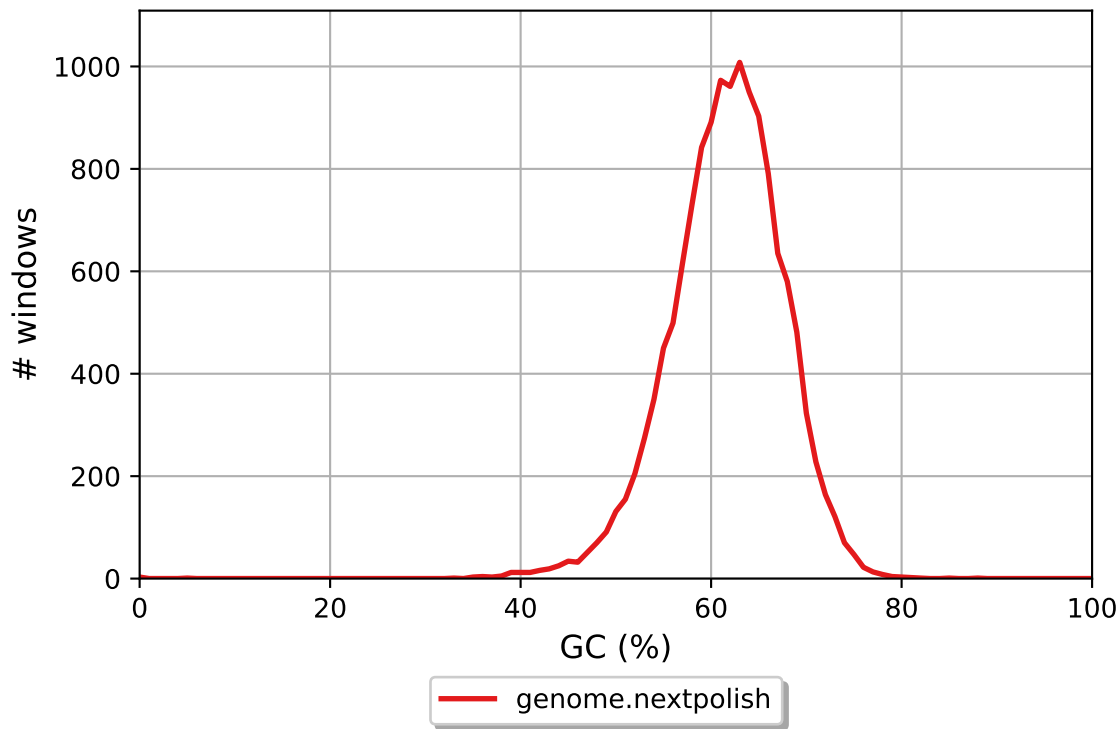
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

