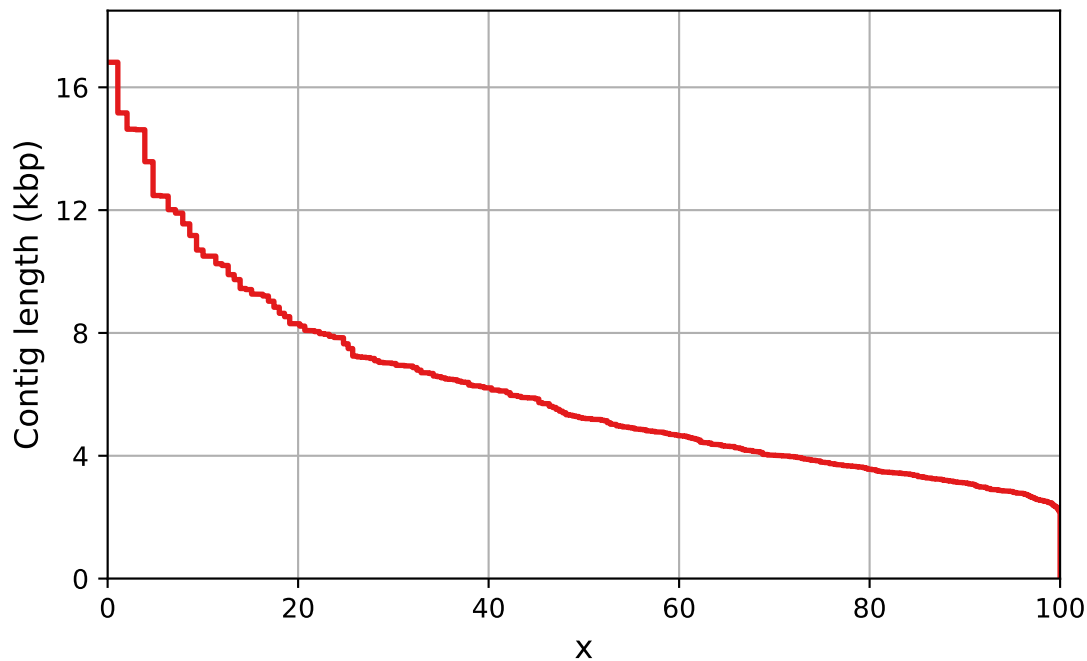


Report

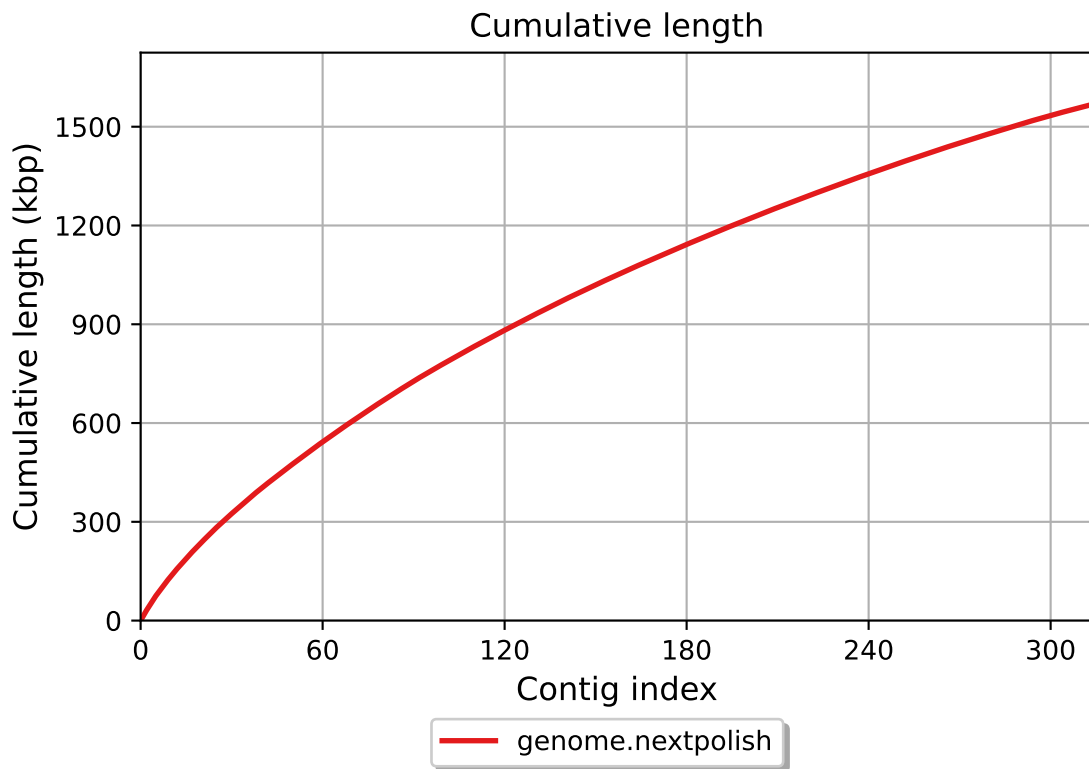
	genome.nextpolish
# contigs (>= 0 bp)	314
# contigs (>= 1000 bp)	314
# contigs (>= 5000 bp)	111
# contigs (>= 10000 bp)	16
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1568183
Total length (>= 1000 bp)	1568183
Total length (>= 5000 bp)	837706
Total length (>= 10000 bp)	198541
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	314
Largest contig	16815
Total length	1568183
GC (%)	57.08
N50	5221
N90	3114
auN	6188.8
L50	101
L90	258
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

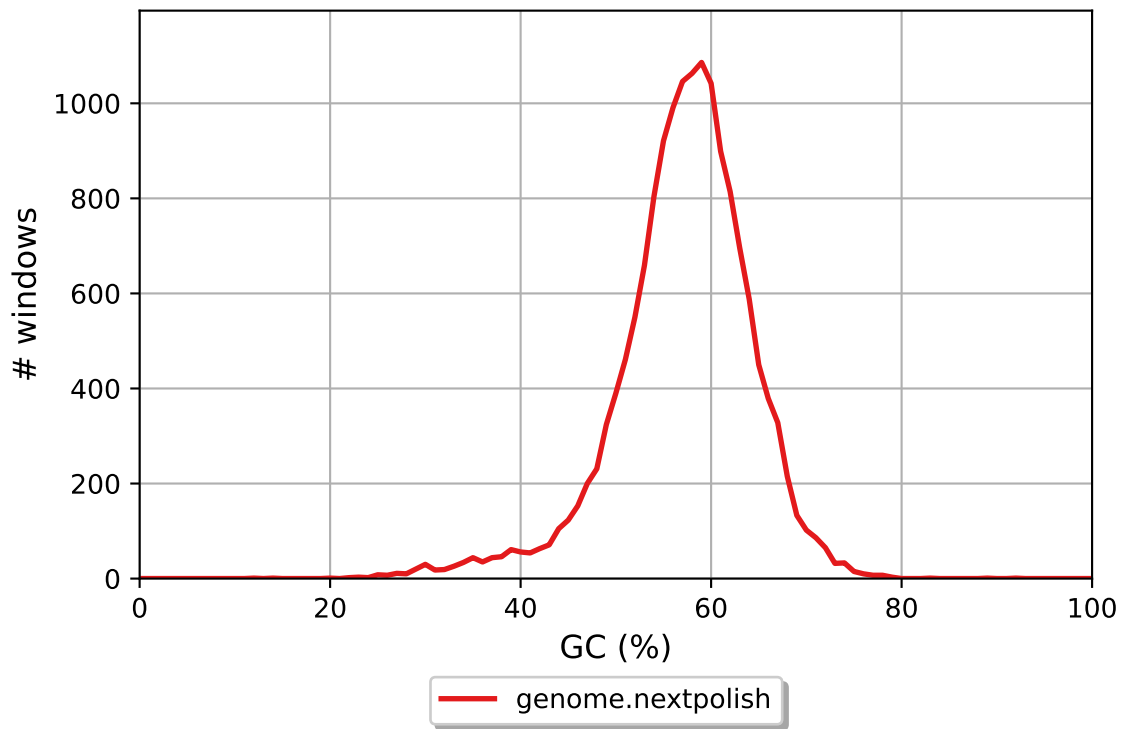
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

