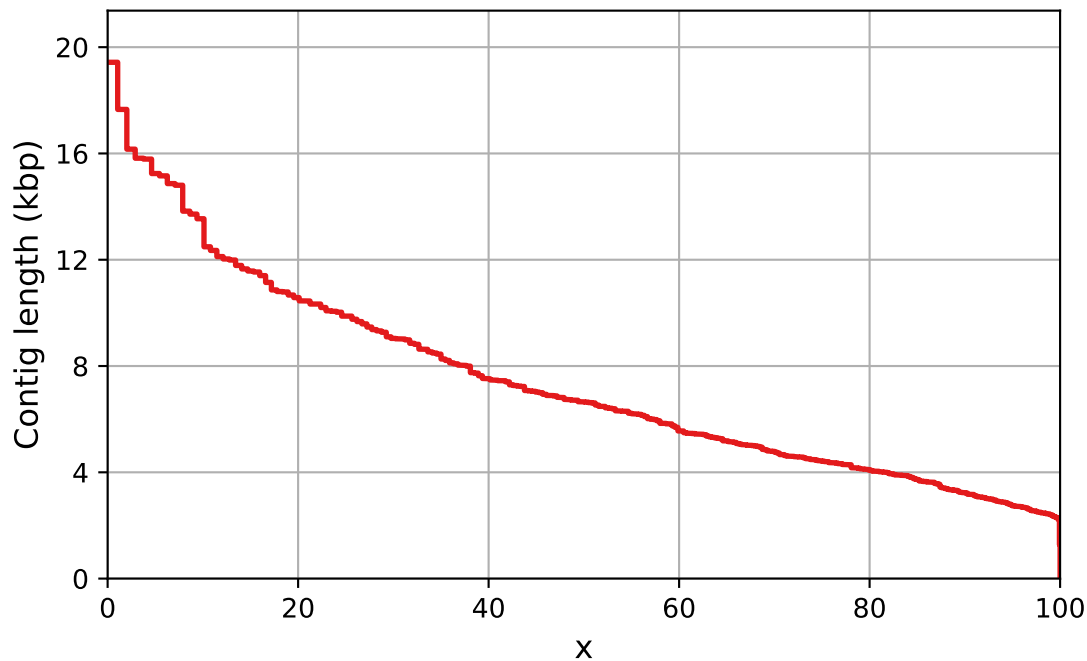


Report

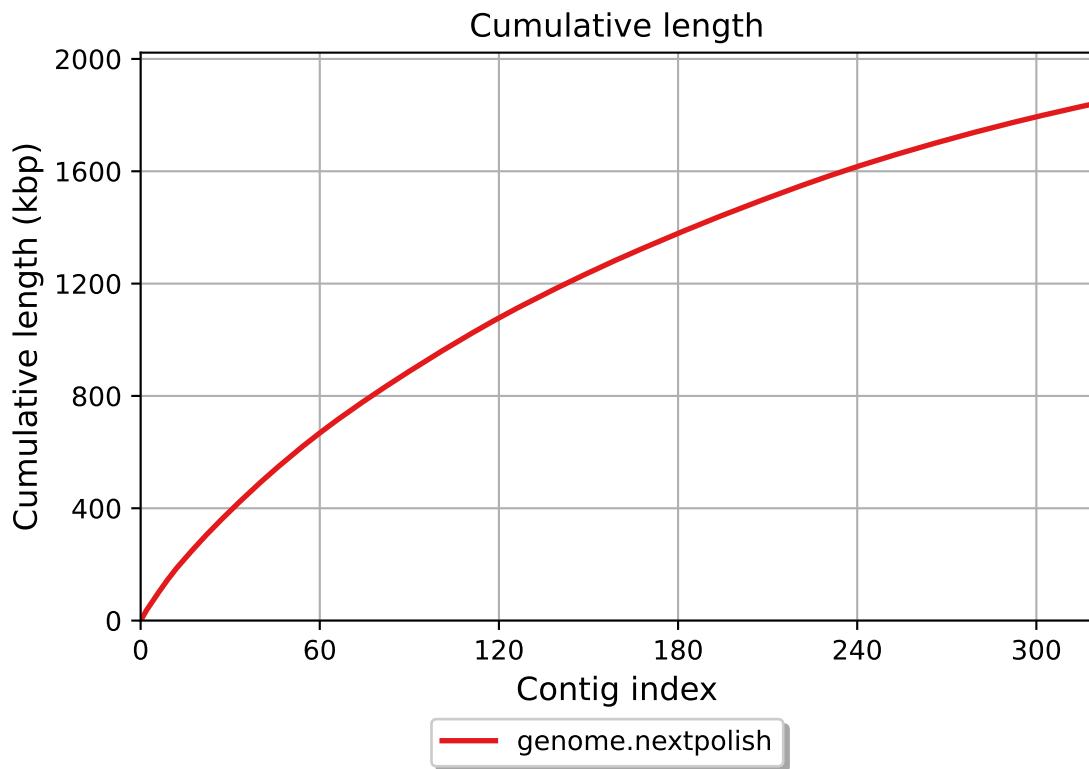
	genome.nextpolish
# contigs (>= 0 bp)	319
# contigs (>= 1000 bp)	319
# contigs (>= 5000 bp)	152
# contigs (>= 10000 bp)	36
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1838590
Total length (>= 1000 bp)	1838590
Total length (>= 5000 bp)	1248500
Total length (>= 10000 bp)	451722
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	319
Largest contig	19431
Total length	1838590
GC (%)	41.91
N50	6650
N90	3231
auN	7477.9
L50	95
L90	252
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

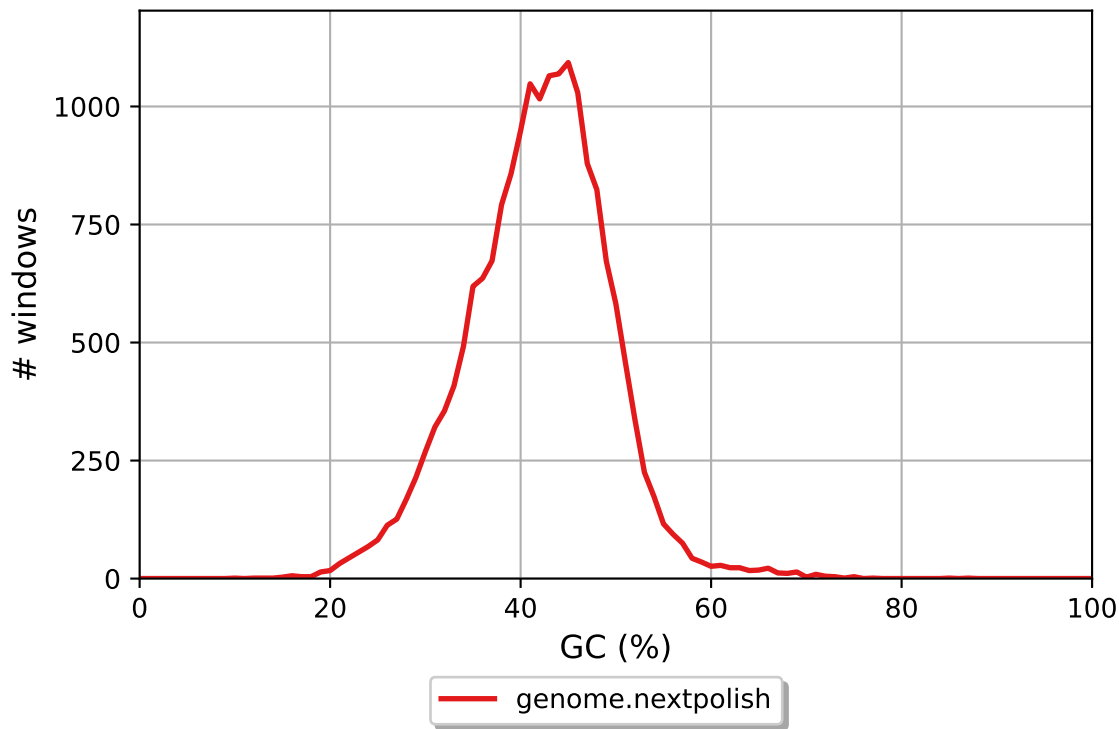
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

