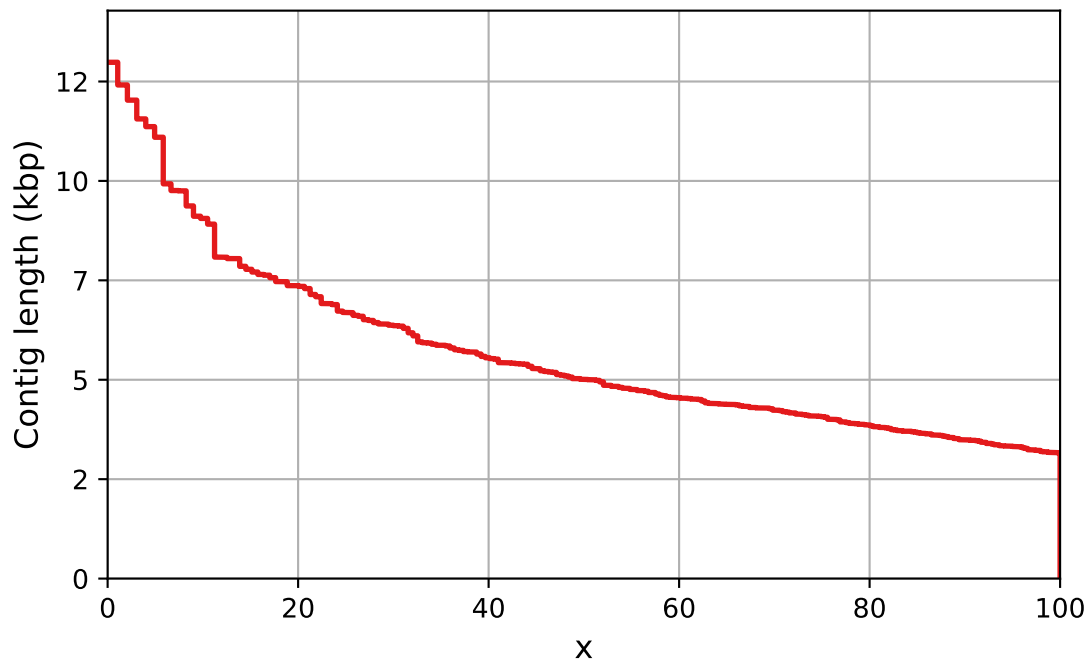


Report

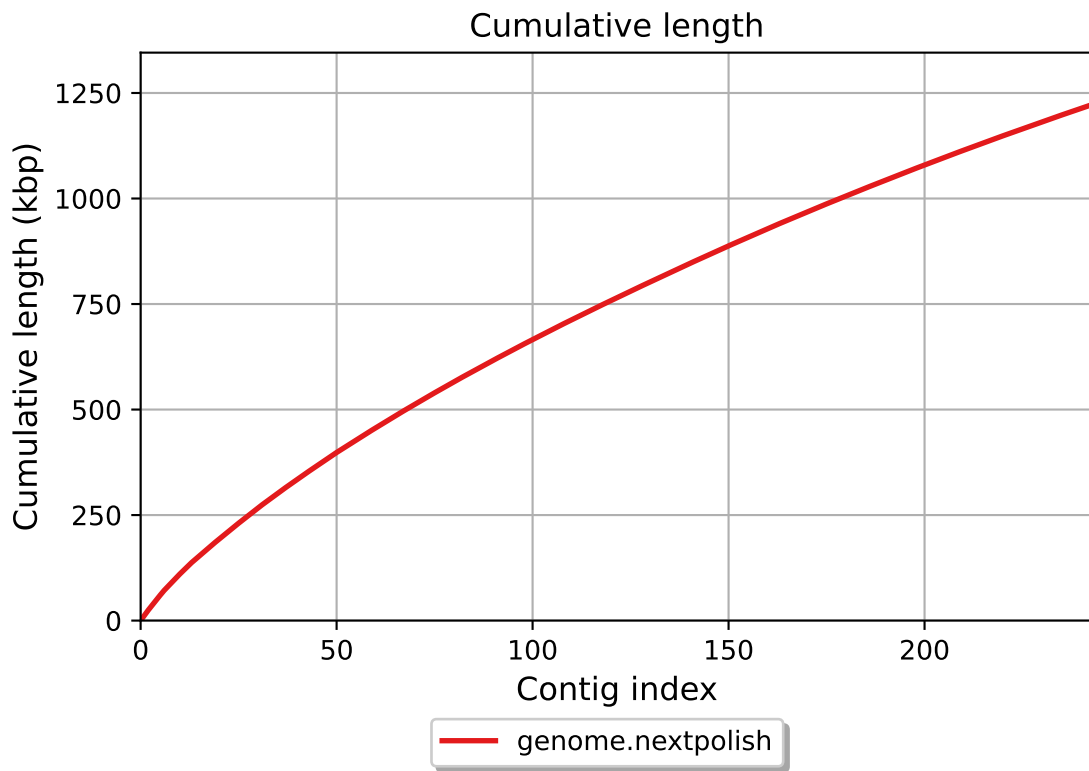
	genome.nextpolish
# contigs (>= 0 bp)	243
# contigs (>= 1000 bp)	243
# contigs (>= 5000 bp)	91
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1223273
Total length (>= 1000 bp)	1223273
Total length (>= 5000 bp)	621859
Total length (>= 10000 bp)	71449
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	243
Largest contig	12983
Total length	1223273
GC (%)	47.35
N50	5012
N90	3486
auN	5698.4
L50	89
L90	207
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

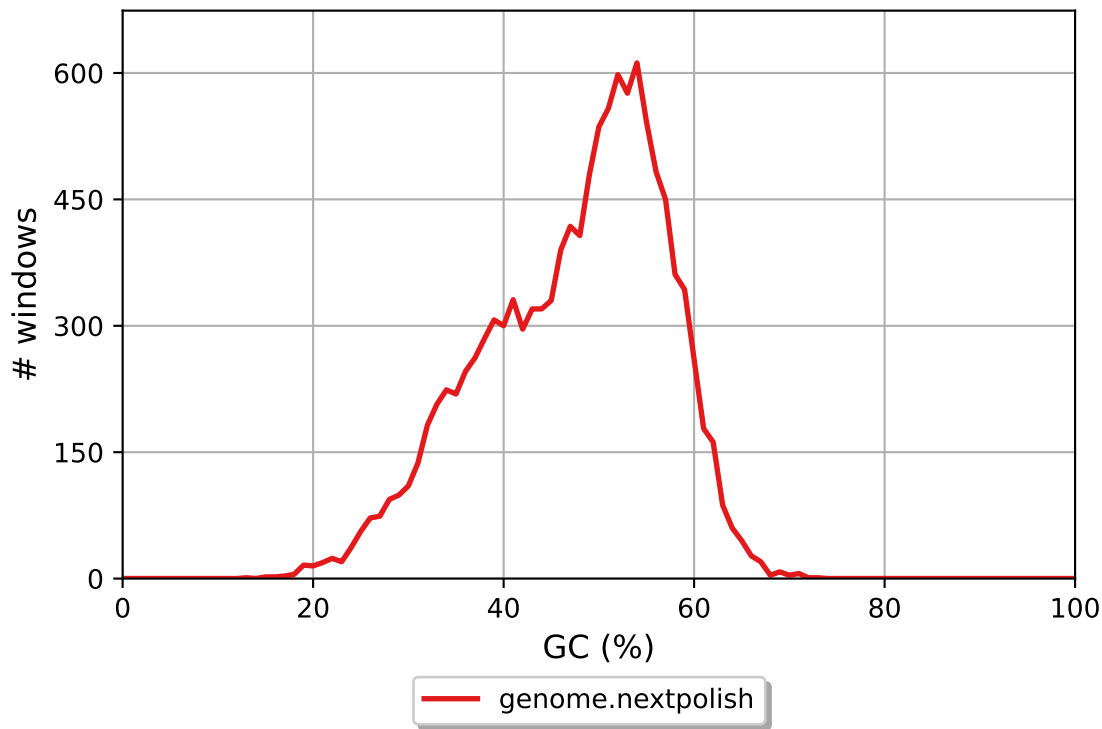
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

