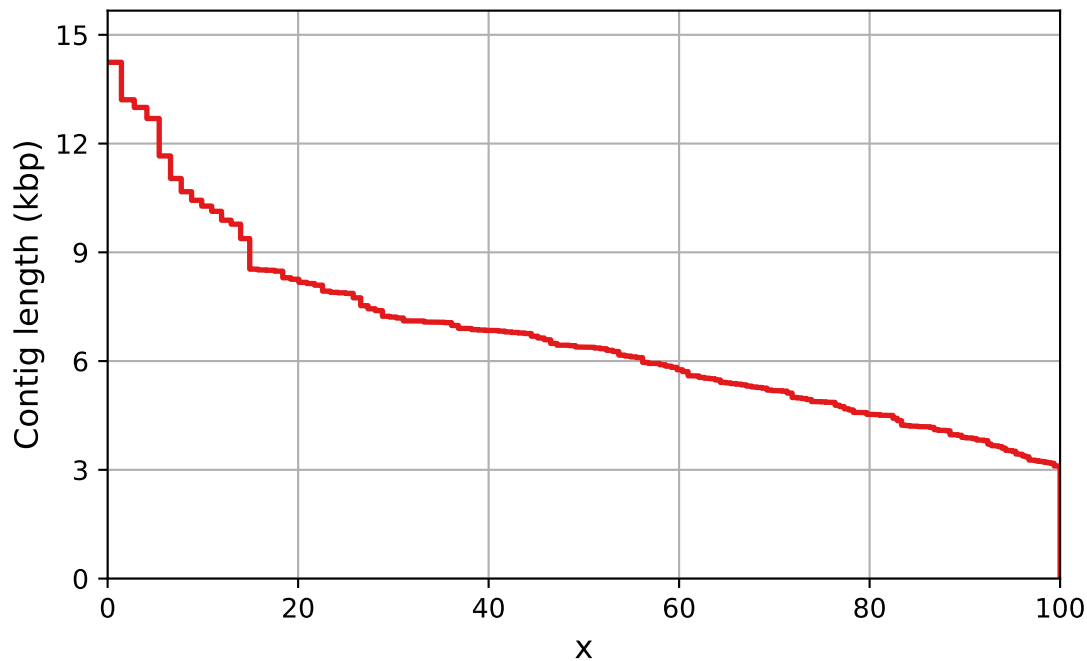


Report

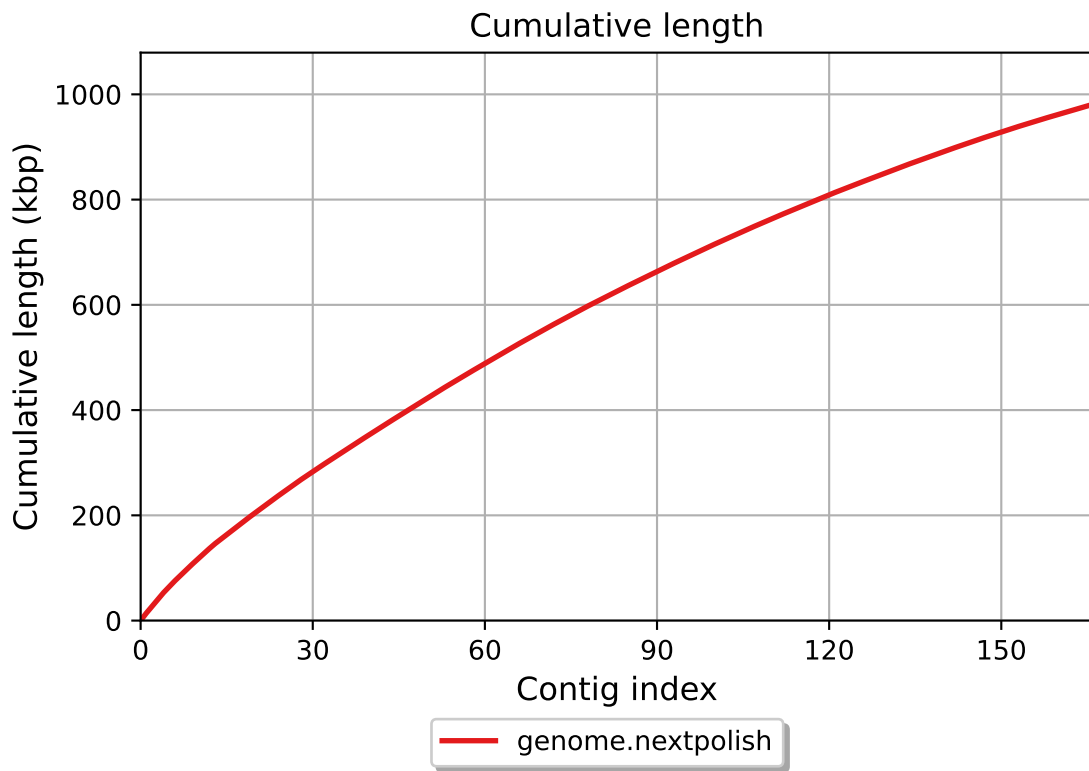
	genome.nextpolish
# contigs (>= 0 bp)	166
# contigs (>= 1000 bp)	166
# contigs (>= 5000 bp)	98
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	981141
Total length (>= 1000 bp)	981141
Total length (>= 5000 bp)	705093
Total length (>= 10000 bp)	117347
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	166
Largest contig	14242
Total length	981141
GC (%)	51.32
N50	6384
N90	3897
auN	6696.7
L50	61
L90	138
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

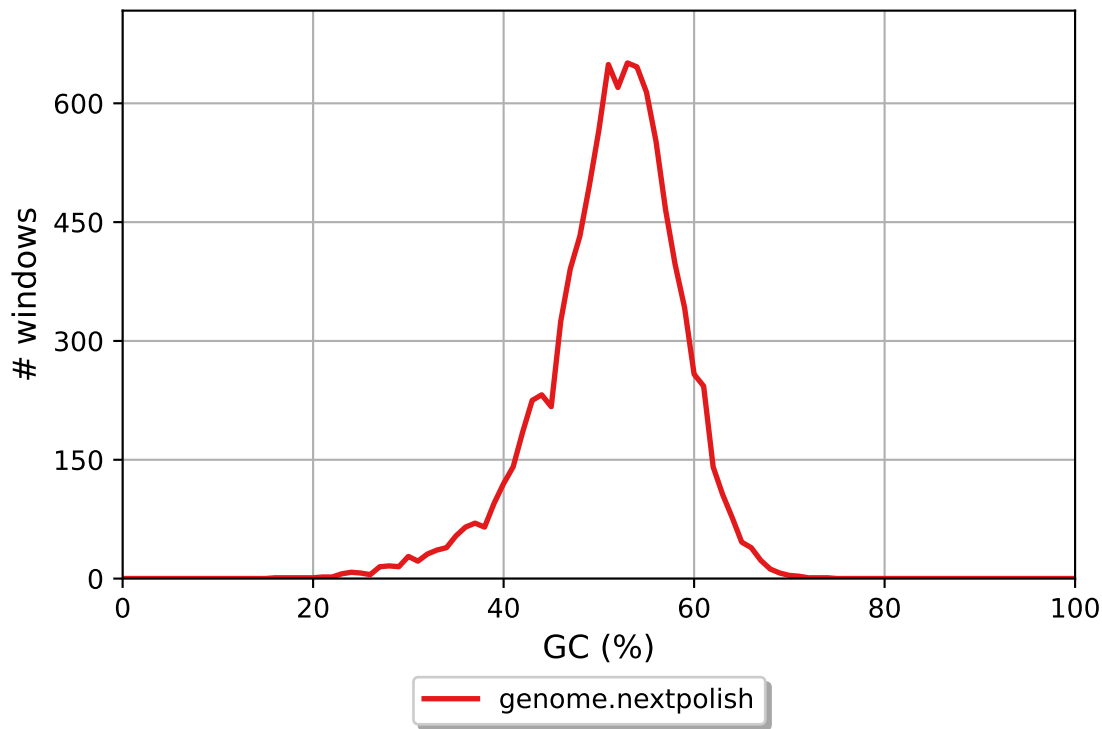
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

