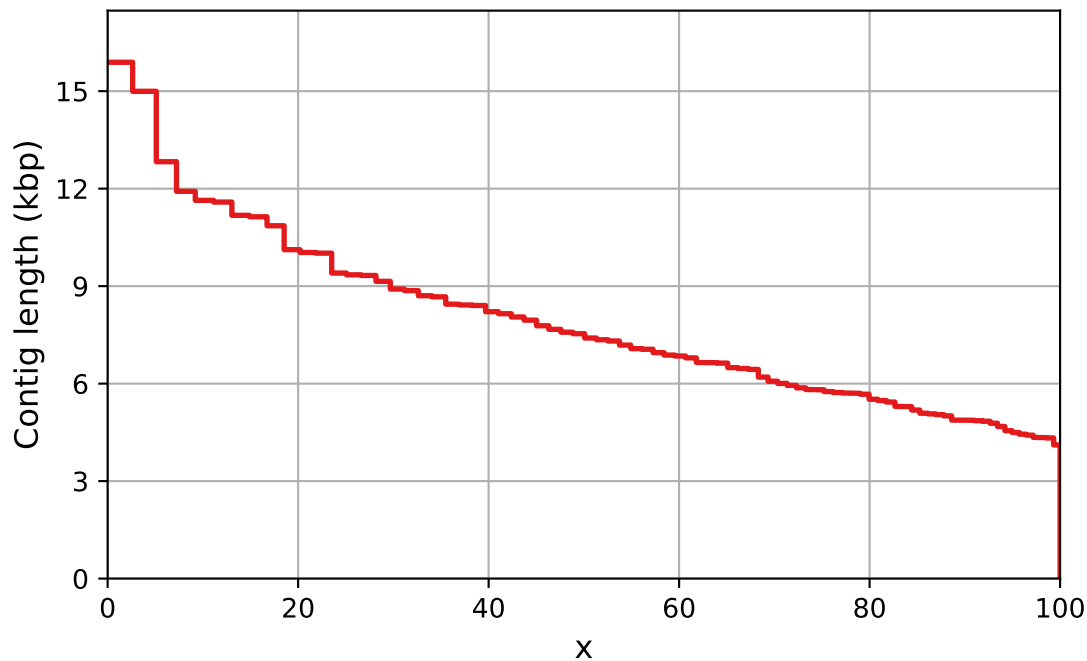


Report

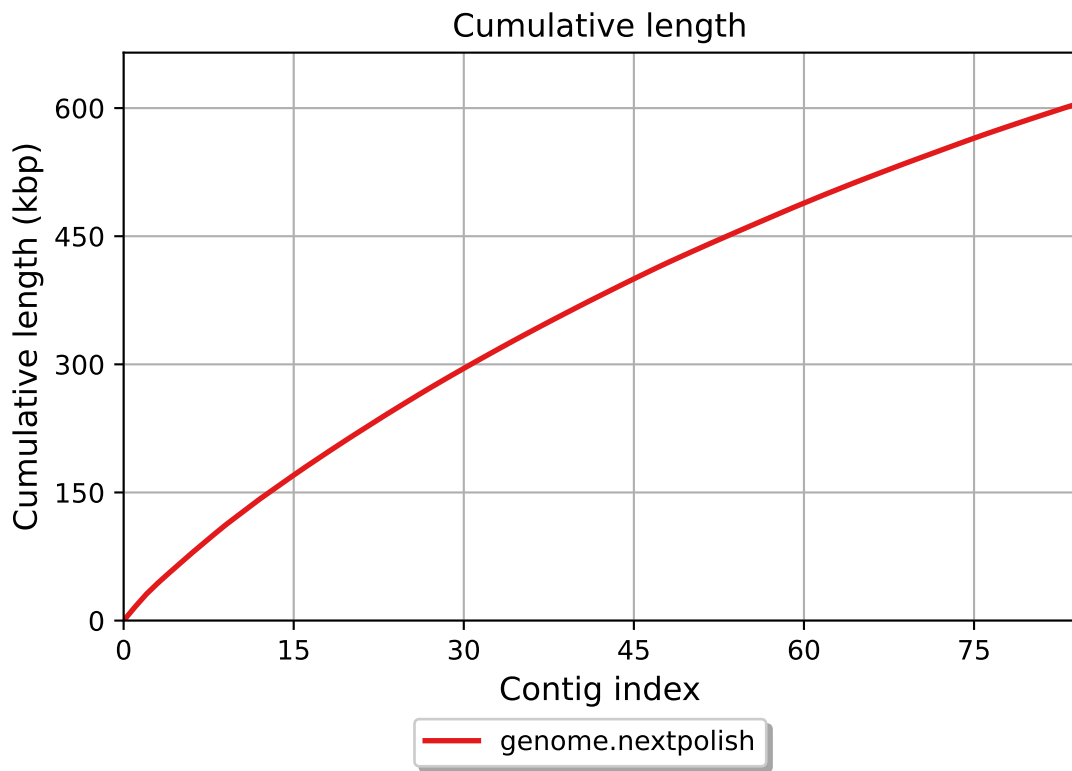
	genome.nextpolish
# contigs (>= 0 bp)	84
# contigs (>= 1000 bp)	84
# contigs (>= 5000 bp)	69
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	604481
Total length (>= 1000 bp)	604481
Total length (>= 5000 bp)	535664
Total length (>= 10000 bp)	142201
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	84
Largest contig	15889
Total length	604481
GC (%)	51.21
N50	7536
N90	4874
auN	8026.3
L50	31
L90	71
# 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).

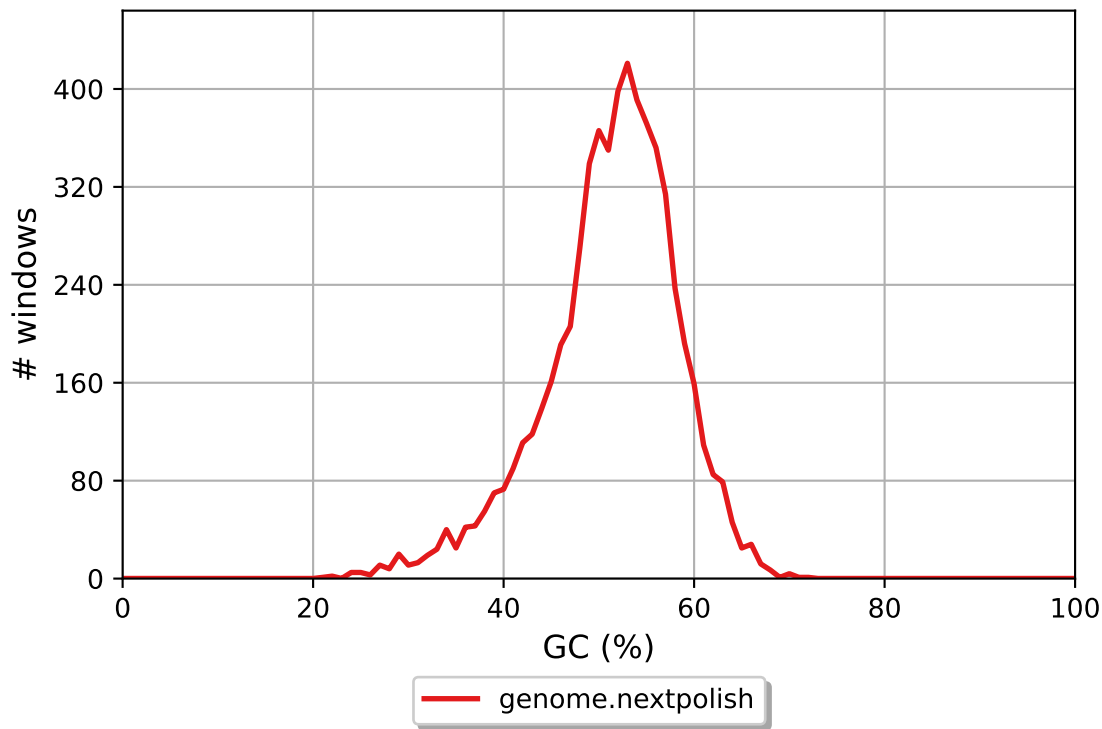
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

