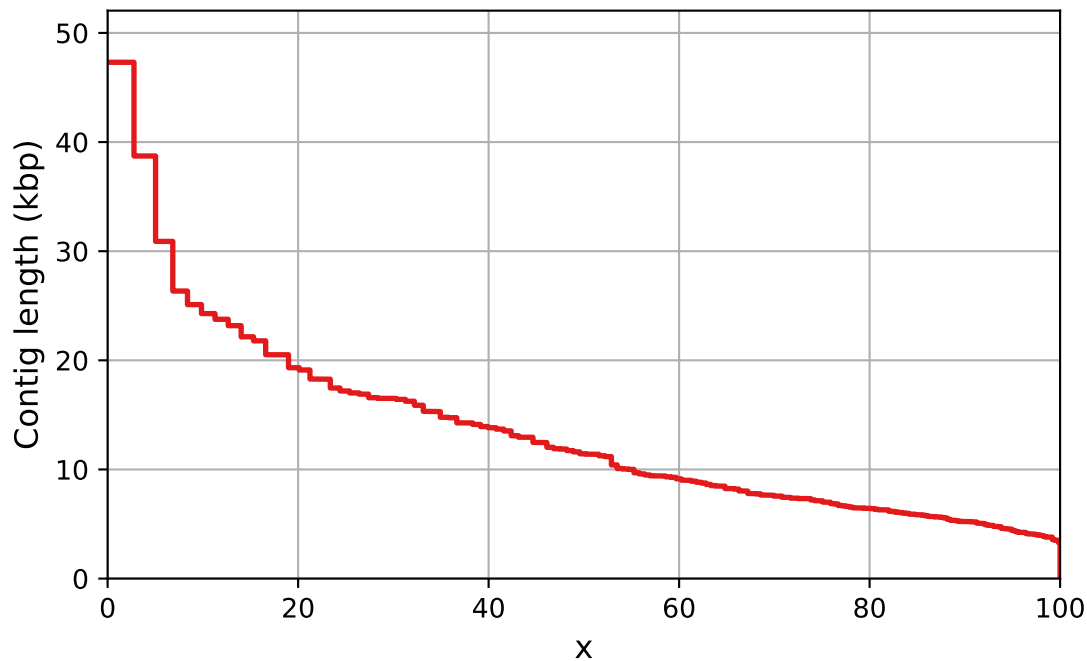


# Report

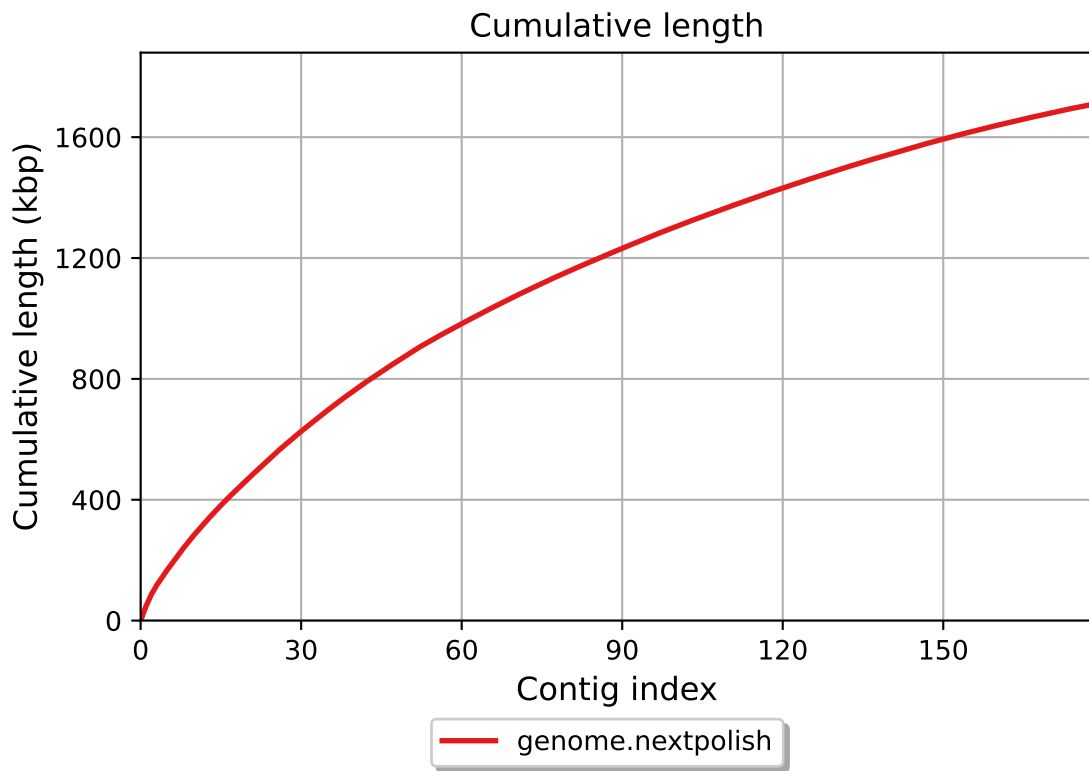
	genome.nextpolish
# contigs (>= 0 bp)	178
# contigs (>= 1000 bp)	178
# contigs (>= 5000 bp)	146
# contigs (>= 10000 bp)	56
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1708880
Total length (>= 1000 bp)	1708880
Total length (>= 5000 bp)	1574348
Total length (>= 10000 bp)	944367
Total length (>= 25000 bp)	168385
Total length (>= 50000 bp)	0
# contigs	178
Largest contig	47311
Total length	1708880
GC (%)	46.47
N50	11434
N90	5240
auN	13860.0
L50	48
L90	139
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 2 part

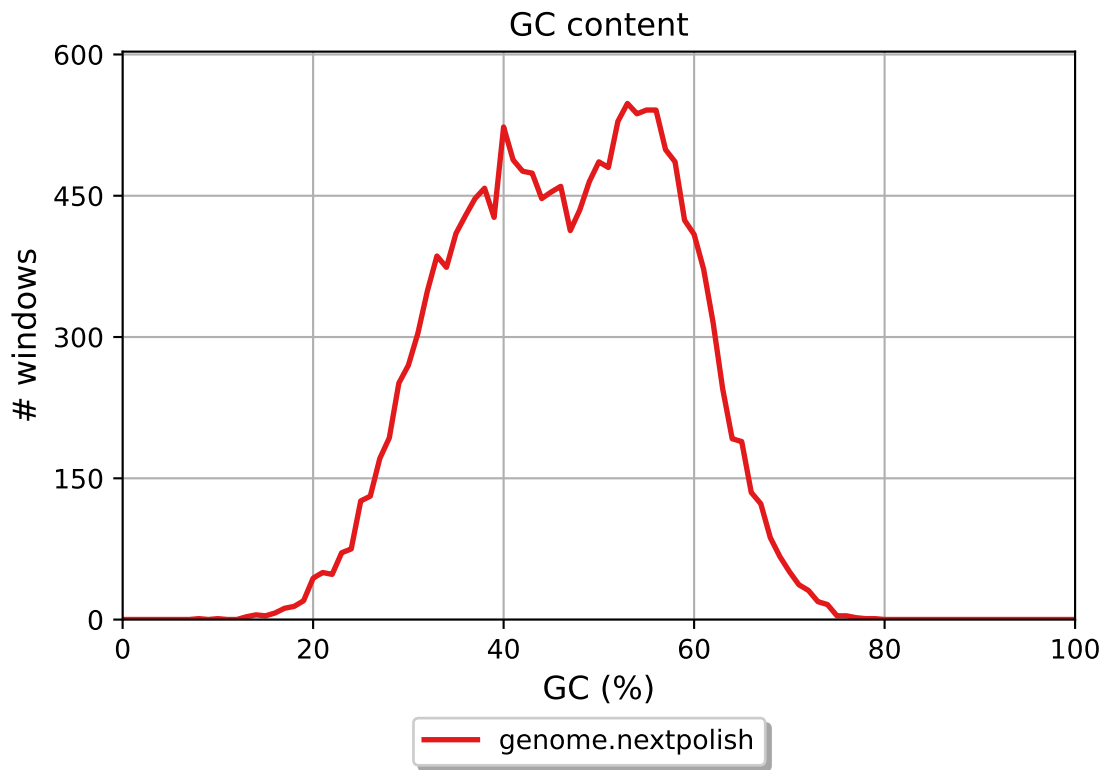
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

