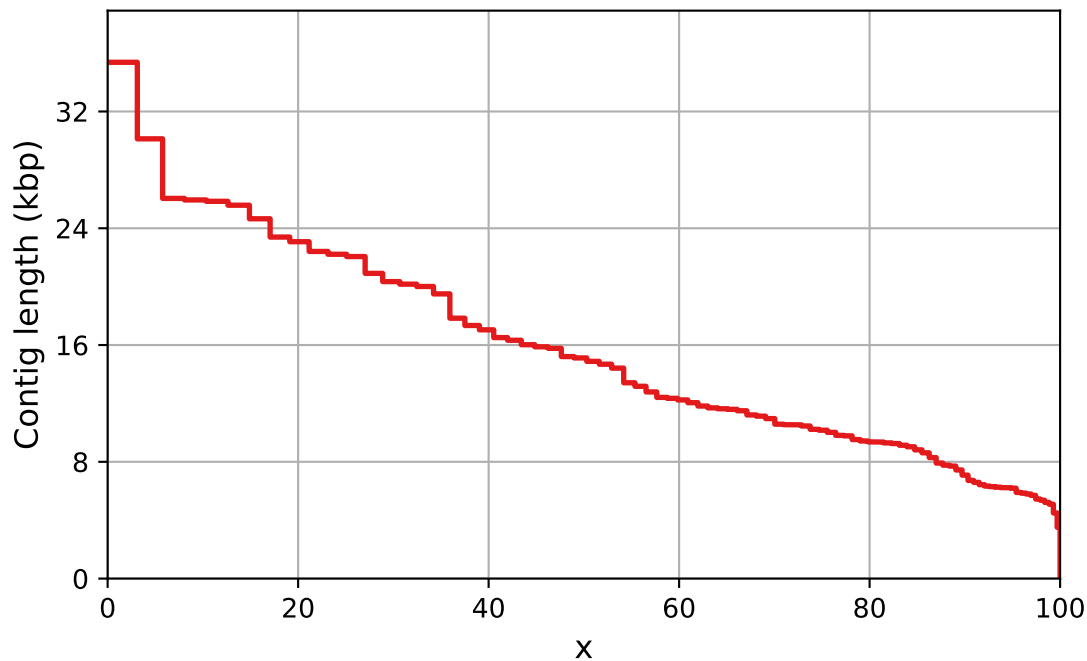


Report

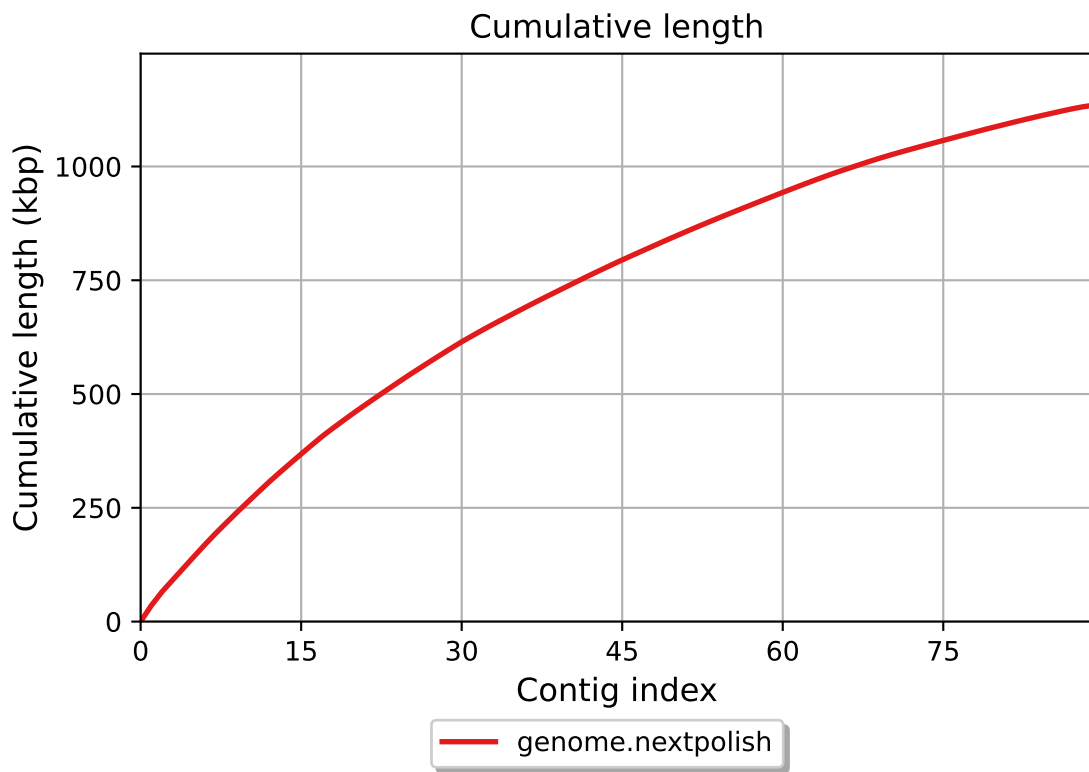
	genome.nextpolish
# contigs (>= 0 bp)	89
# contigs (>= 1000 bp)	89
# contigs (>= 5000 bp)	87
# contigs (>= 10000 bp)	52
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1134504
Total length (>= 1000 bp)	1134504
Total length (>= 5000 bp)	1126494
Total length (>= 10000 bp)	867206
Total length (>= 25000 bp)	168909
Total length (>= 50000 bp)	0
# contigs	89
Largest contig	35371
Total length	1134504
GC (%)	58.24
N50	15117
N90	7087
auN	16137.2
L50	27
L90	70
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 part

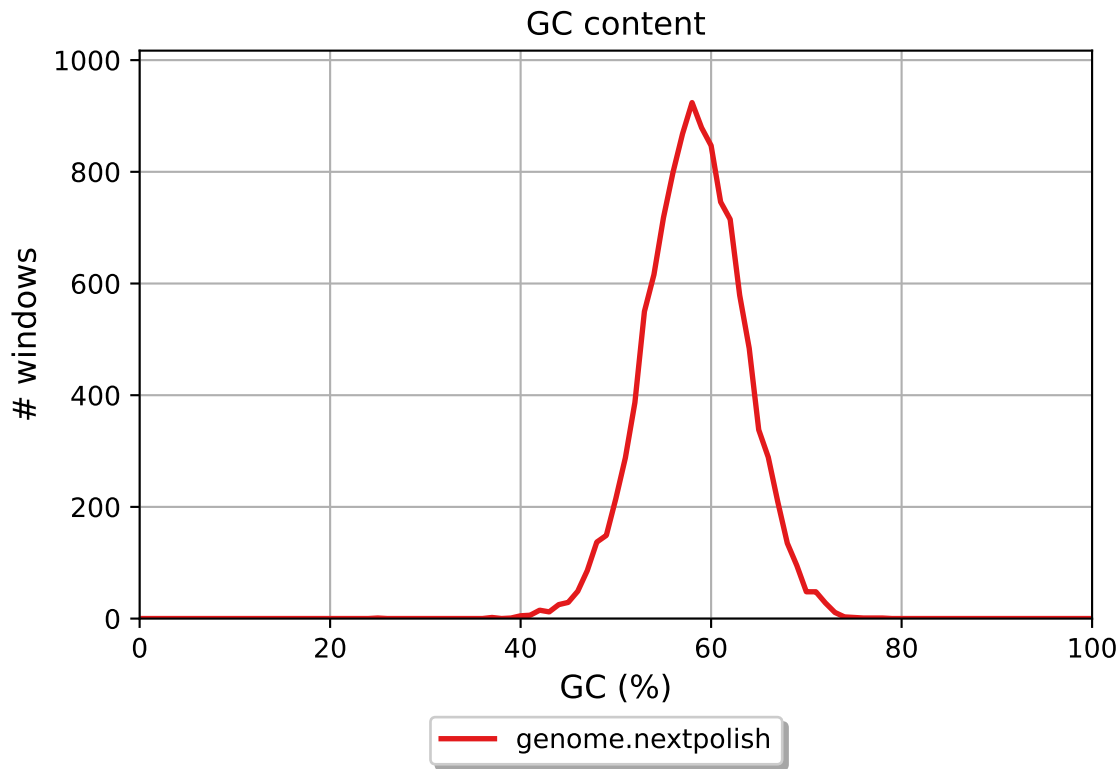
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

