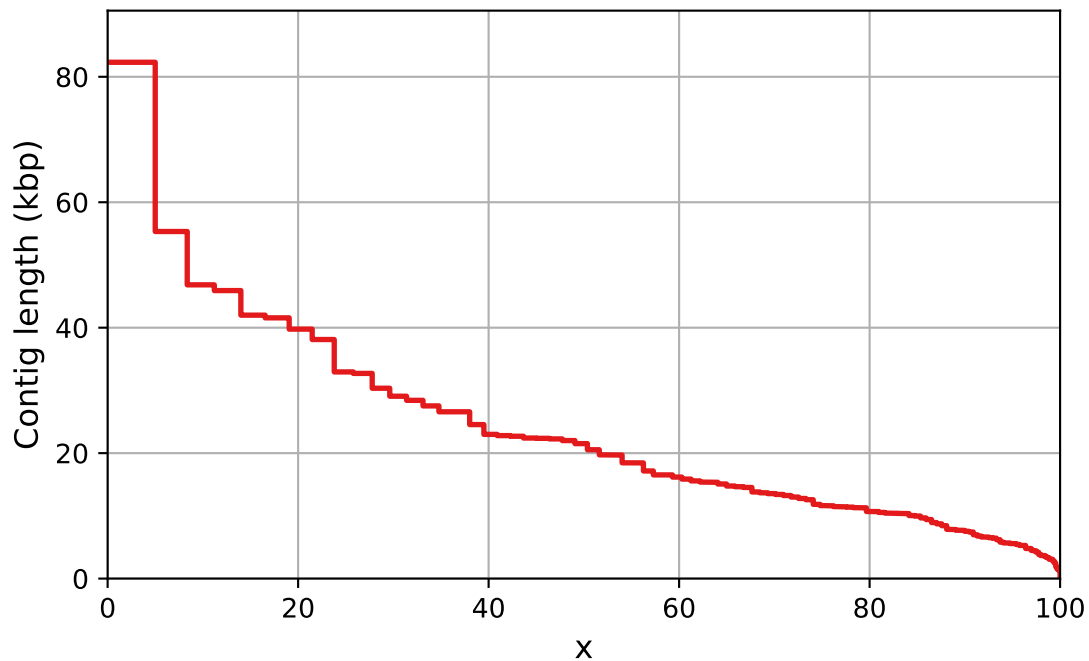


# Report

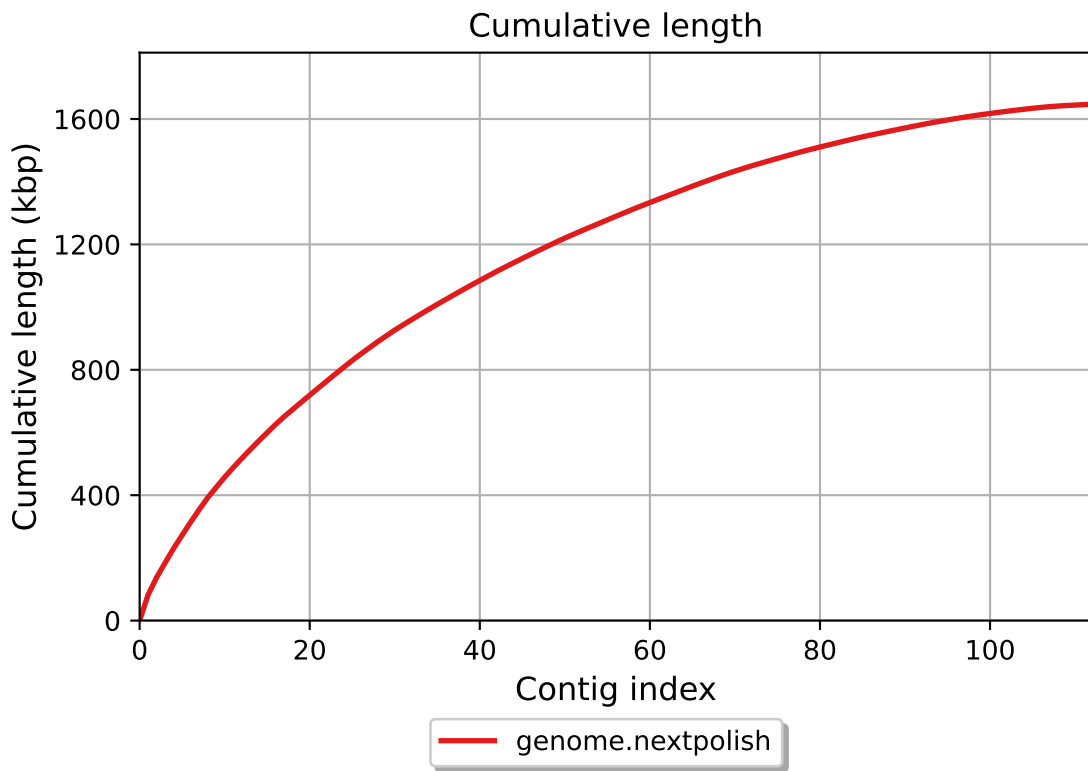
	genome.nextpolish
# contigs (>= 0 bp)	112
# contigs (>= 1000 bp)	112
# contigs (>= 5000 bp)	93
# contigs (>= 10000 bp)	66
# contigs (>= 25000 bp)	16
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	1646998
Total length (>= 1000 bp)	1646998
Total length (>= 5000 bp)	1587410
Total length (>= 10000 bp)	1395703
Total length (>= 25000 bp)	626062
Total length (>= 50000 bp)	137648
# contigs	112
Largest contig	82314
Total length	1646998
GC (%)	46.08
N50	21517
N90	7537
auN	25477.7
L50	25
L90	77
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 2 part

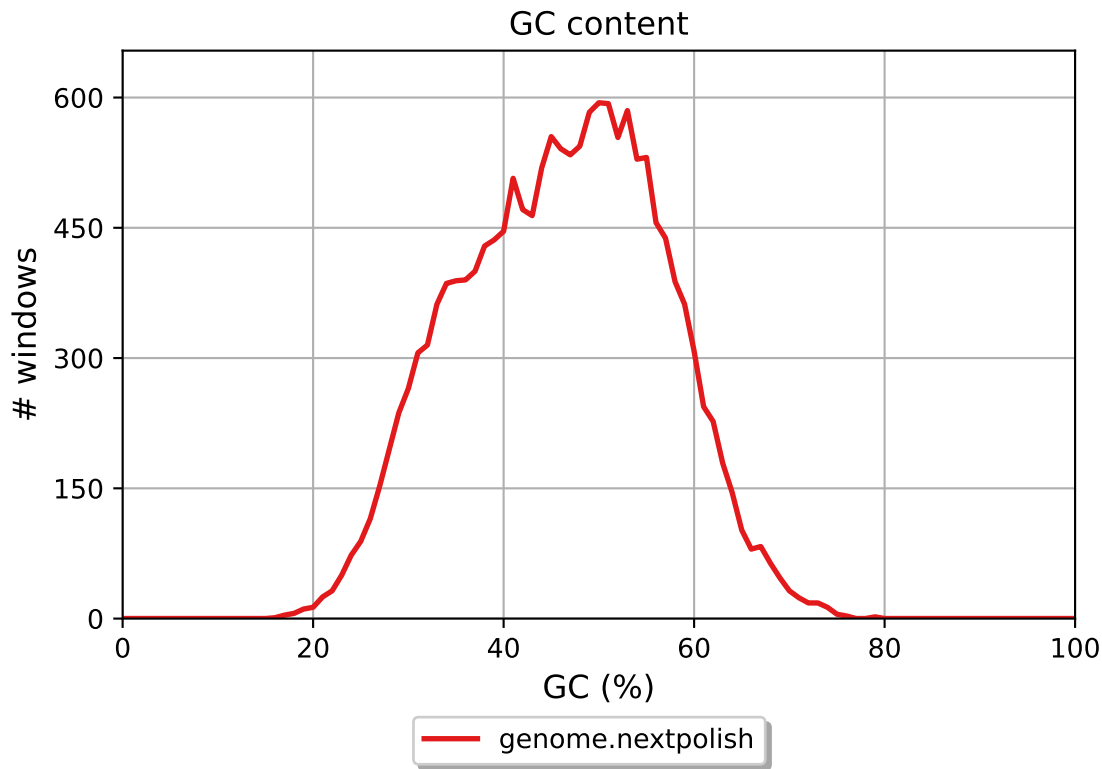
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

