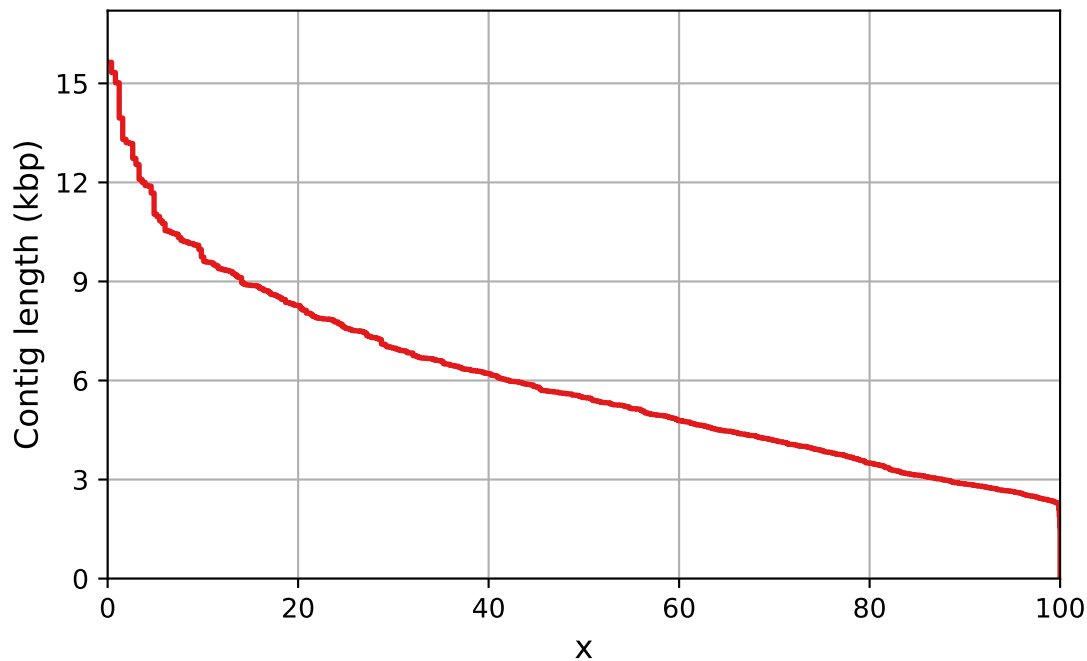


Report

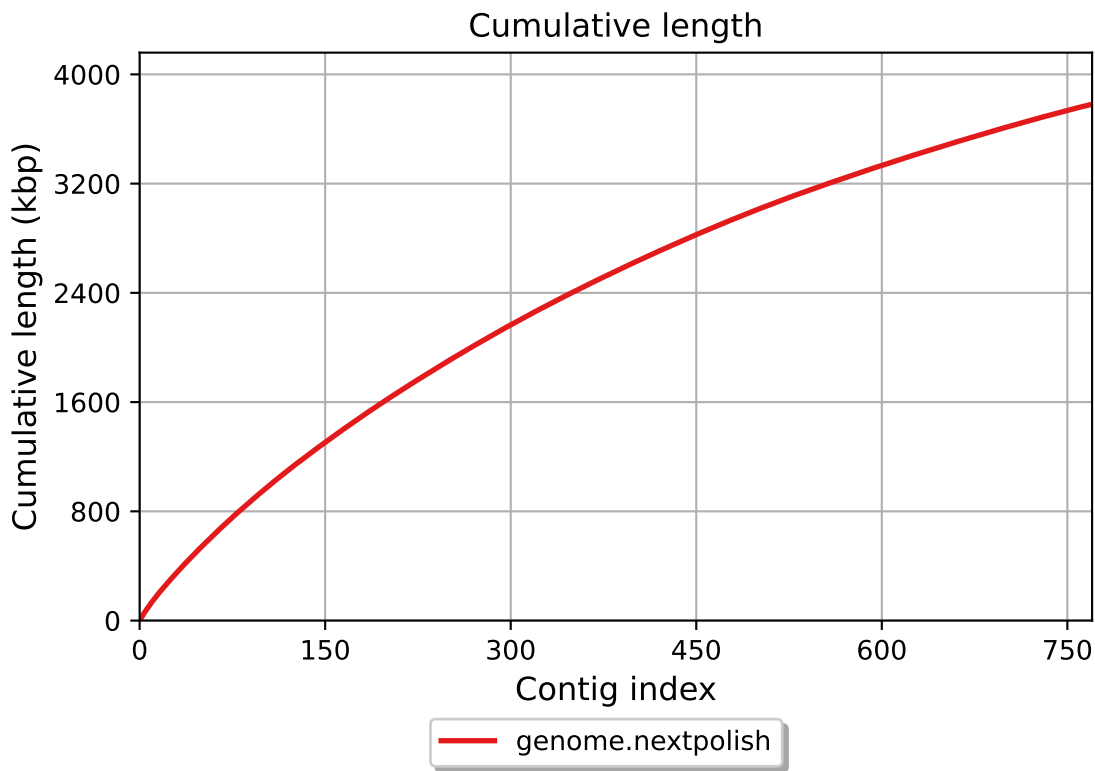
	genome.nextpolish
# contigs (>= 0 bp)	770
# contigs (>= 1000 bp)	770
# contigs (>= 5000 bp)	296
# contigs (>= 10000 bp)	31
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3781035
Total length (>= 1000 bp)	3781035
Total length (>= 5000 bp)	2145621
Total length (>= 10000 bp)	361951
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	770
Largest contig	15639
Total length	3781035
GC (%)	42.57
N50	5484
N90	2870
auN	6001.3
L50	248
L90	624
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 1 part

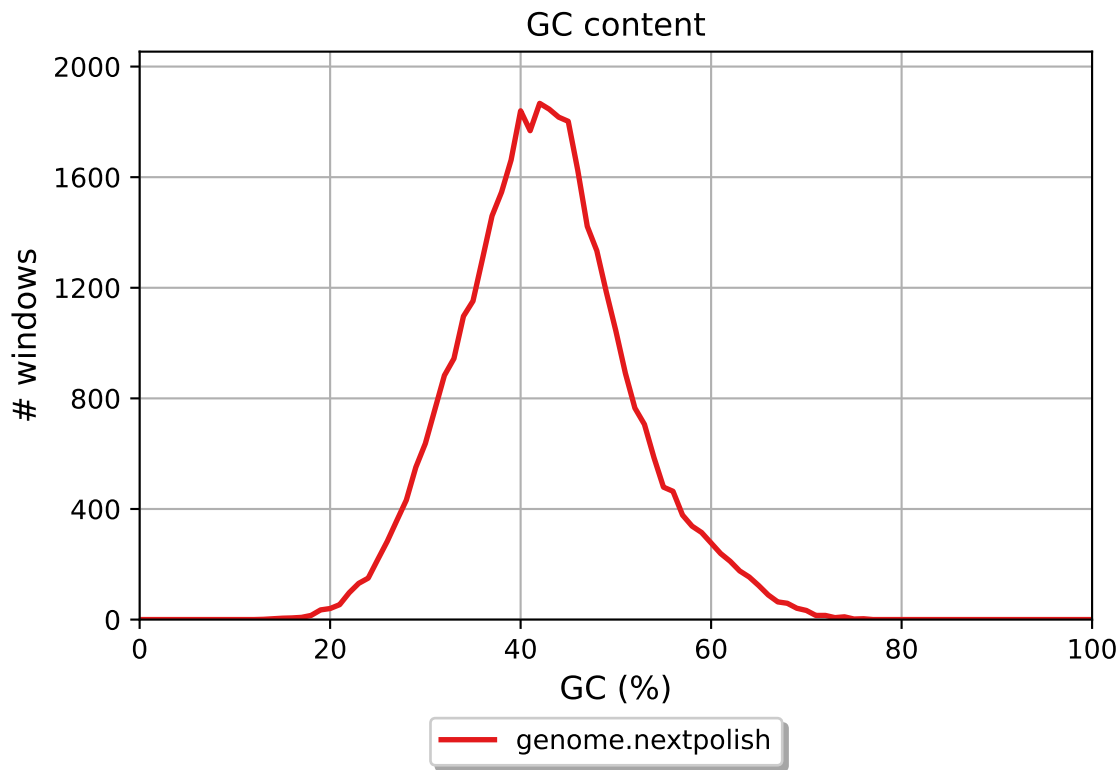
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

