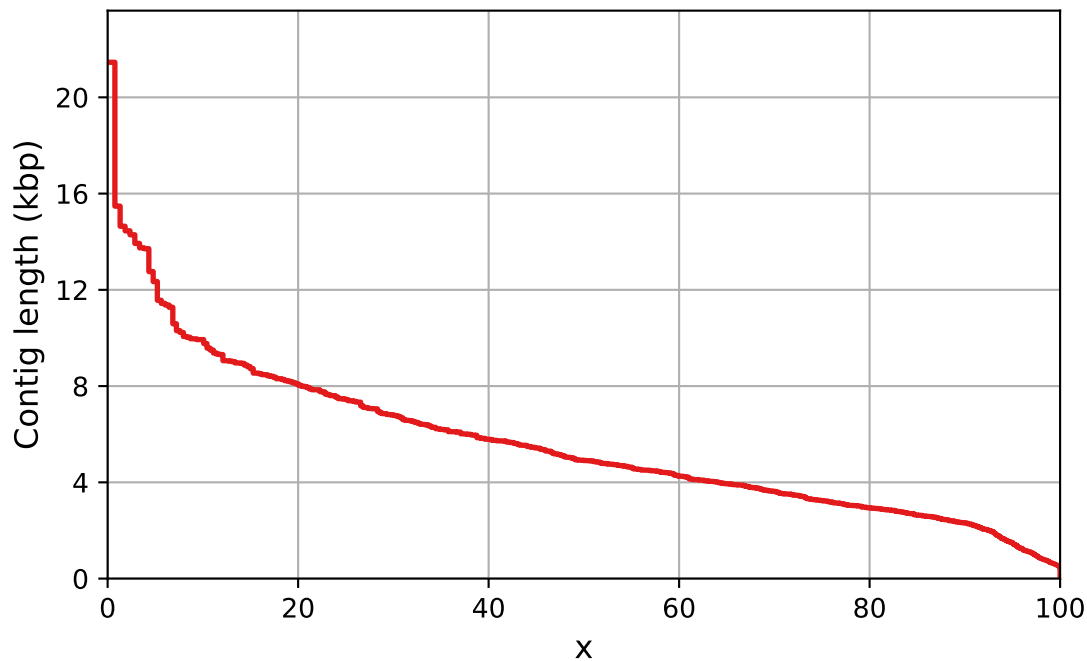


Report

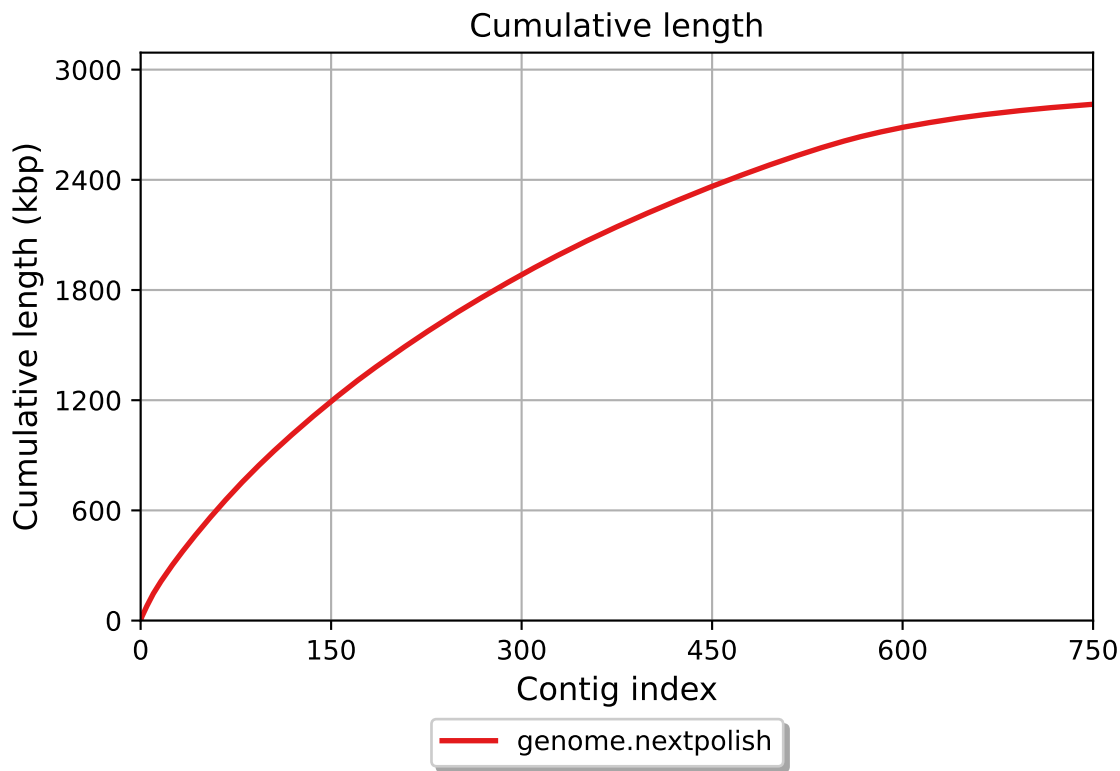
	genome.nextpolish
# contigs (>= 0 bp)	757
# contigs (>= 1000 bp)	643
# contigs (>= 5000 bp)	184
# contigs (>= 10000 bp)	19
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2813312
Total length (>= 1000 bp)	2735505
Total length (>= 5000 bp)	1374759
Total length (>= 10000 bp)	243673
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	750
Largest contig	21457
Total length	2811415
GC (%)	51.59
N50	4909
N90	2315
auN	5667.5
L50	191
L90	517
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 2 part

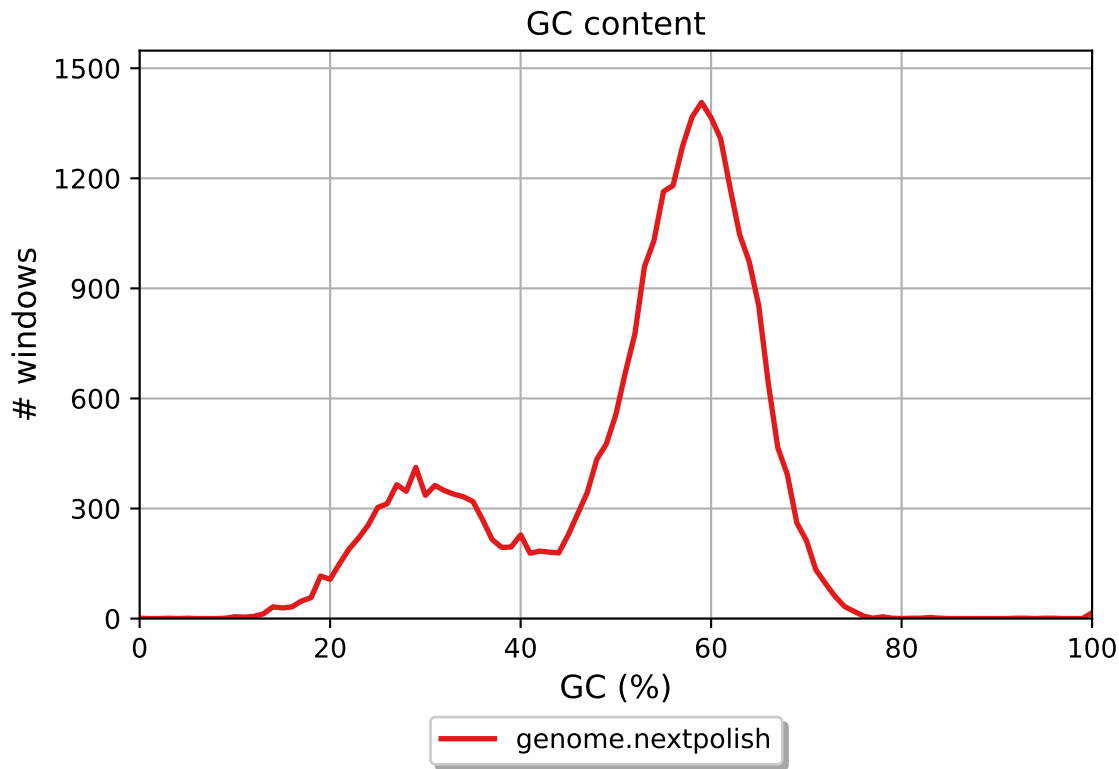
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

