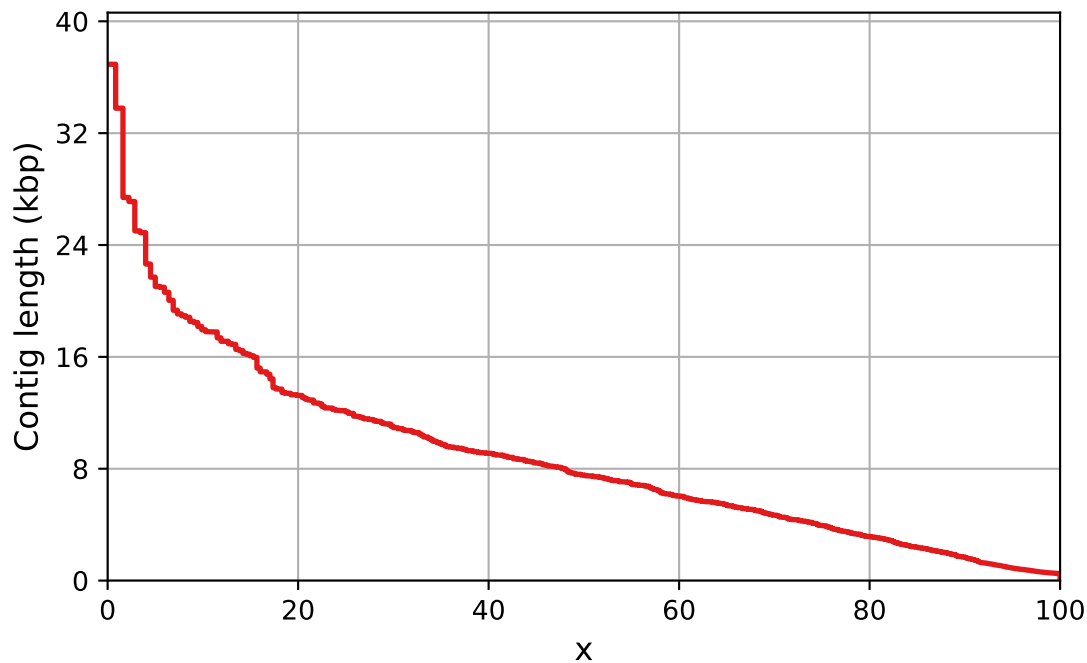


Report

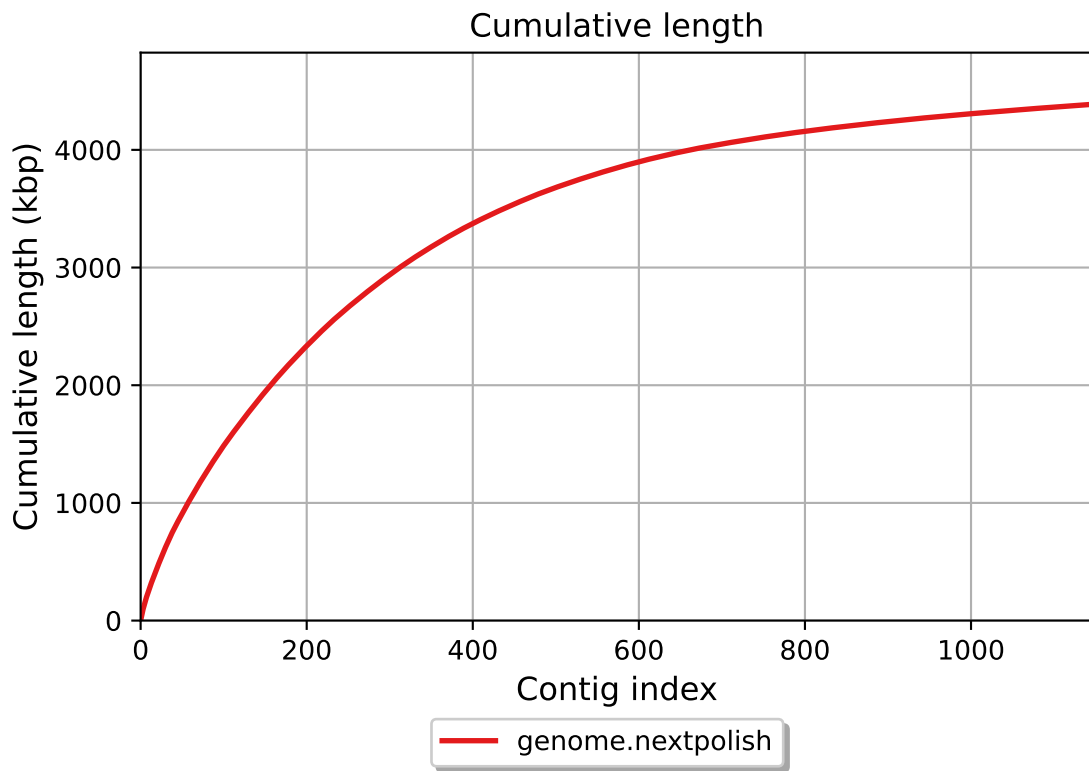
	genome.nextpolish
# contigs (>= 0 bp)	1175
# contigs (>= 1000 bp)	776
# contigs (>= 5000 bp)	310
# contigs (>= 10000 bp)	101
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4397339
Total length (>= 1000 bp)	4134800
Total length (>= 5000 bp)	2990067
Total length (>= 10000 bp)	1495929
Total length (>= 25000 bp)	150227
Total length (>= 50000 bp)	0
# contigs	1147
Largest contig	36926
Total length	4387407
GC (%)	58.54
N50	7547
N90	1673
auN	8951.4
L50	181
L90	629
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 3 part

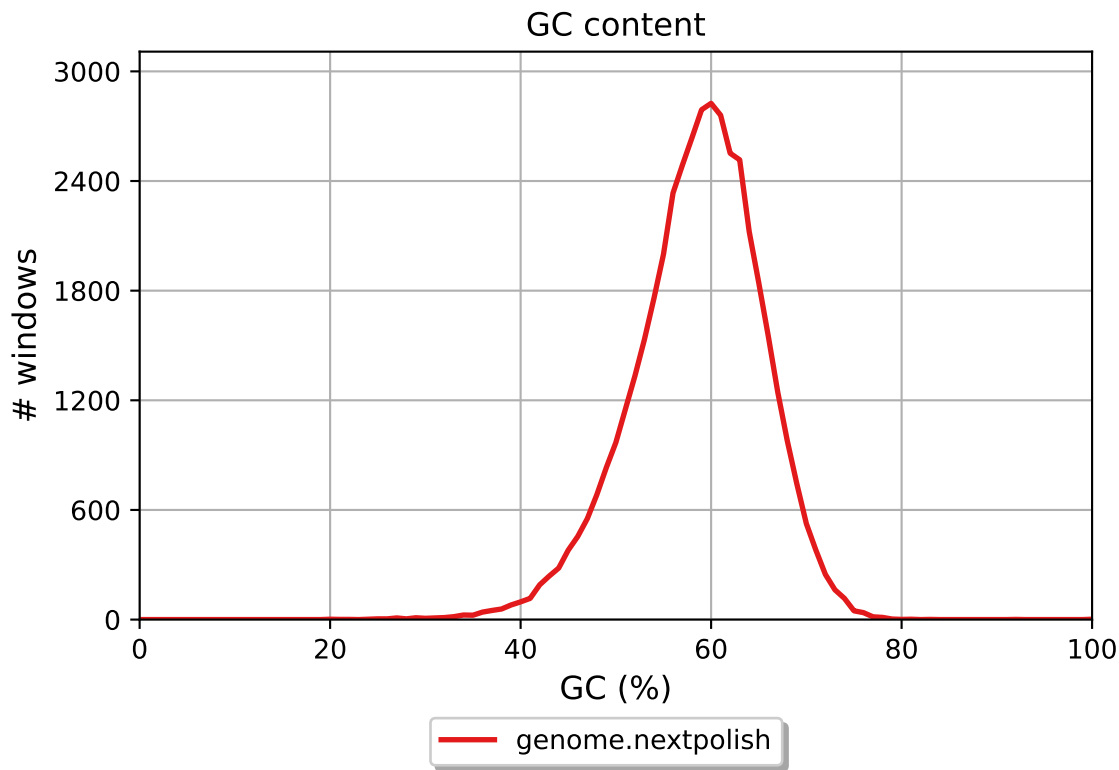
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

