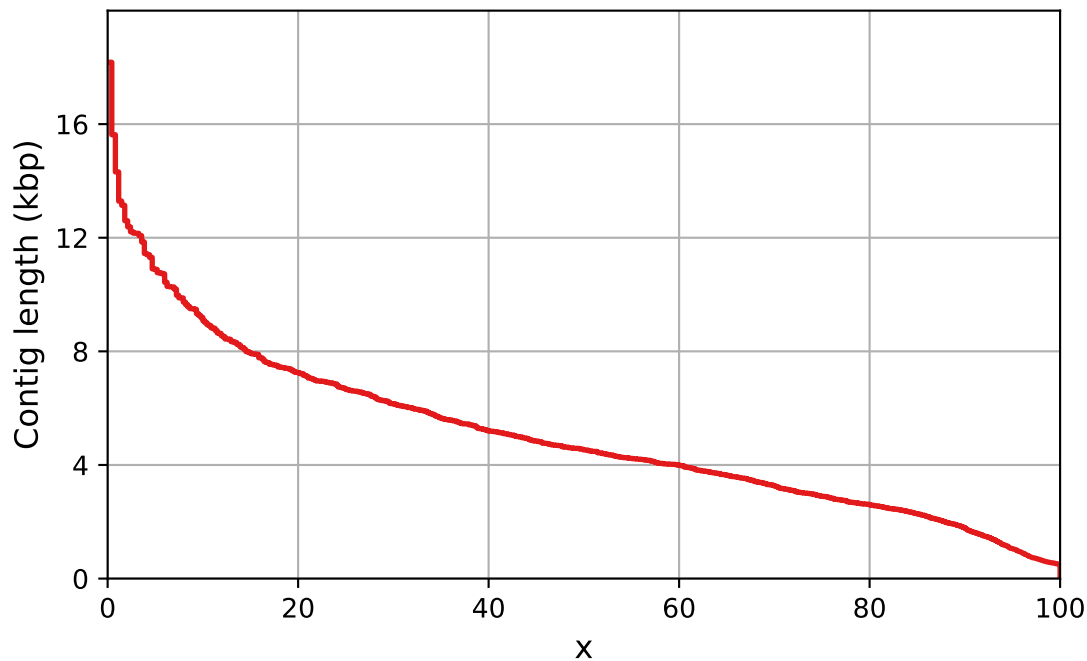


Report

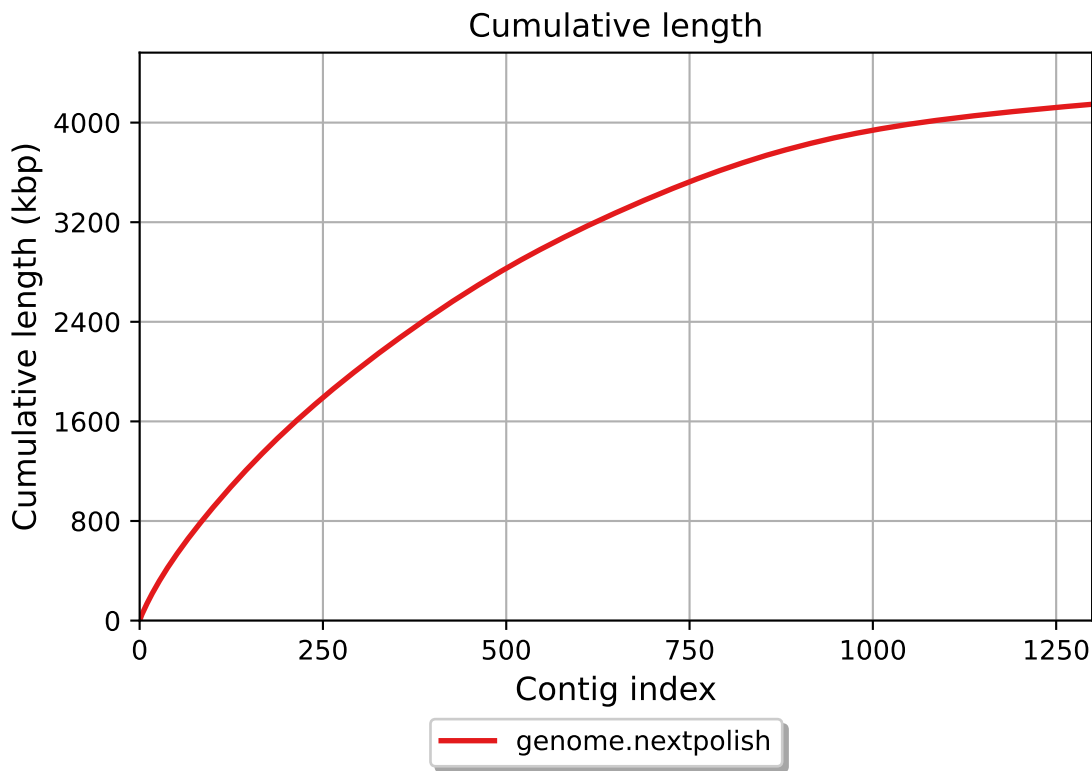
	genome.nextpolish
# contigs (>= 0 bp)	1329
# contigs (>= 1000 bp)	1017
# contigs (>= 5000 bp)	249
# contigs (>= 10000 bp)	25
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4158622
Total length (>= 1000 bp)	3956597
Total length (>= 5000 bp)	1786591
Total length (>= 10000 bp)	299617
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1299
Largest contig	18177
Total length	4147296
GC (%)	37.93
N50	4533
N90	1772
auN	5089.4
L50	310
L90	853
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

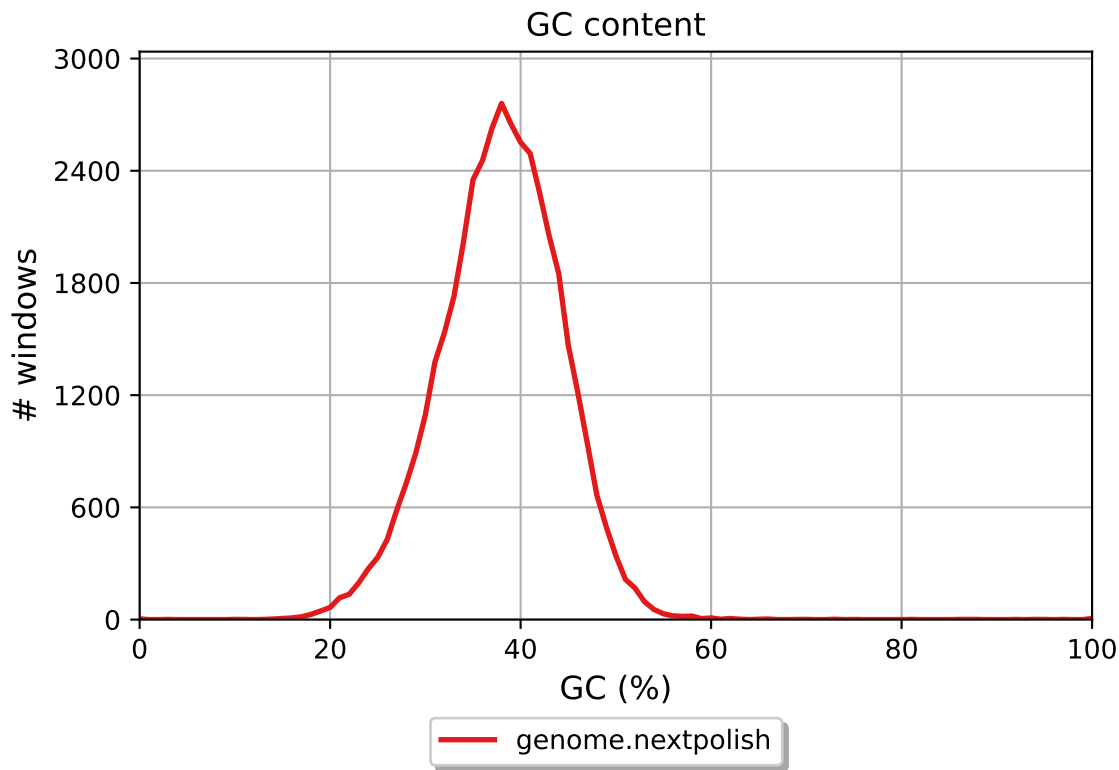
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

