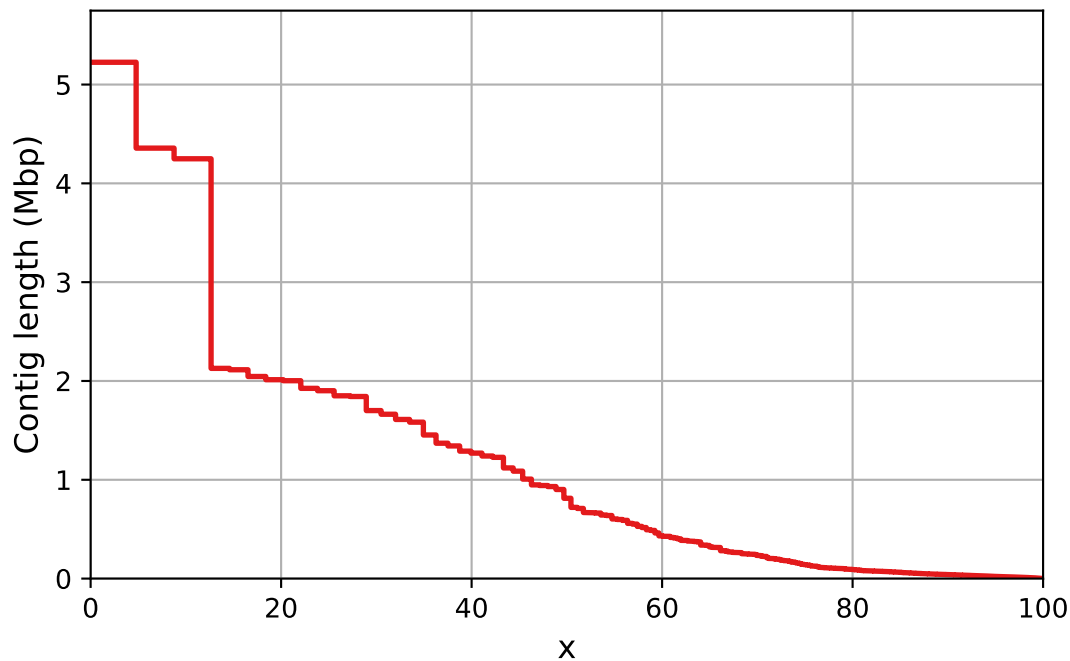


Report

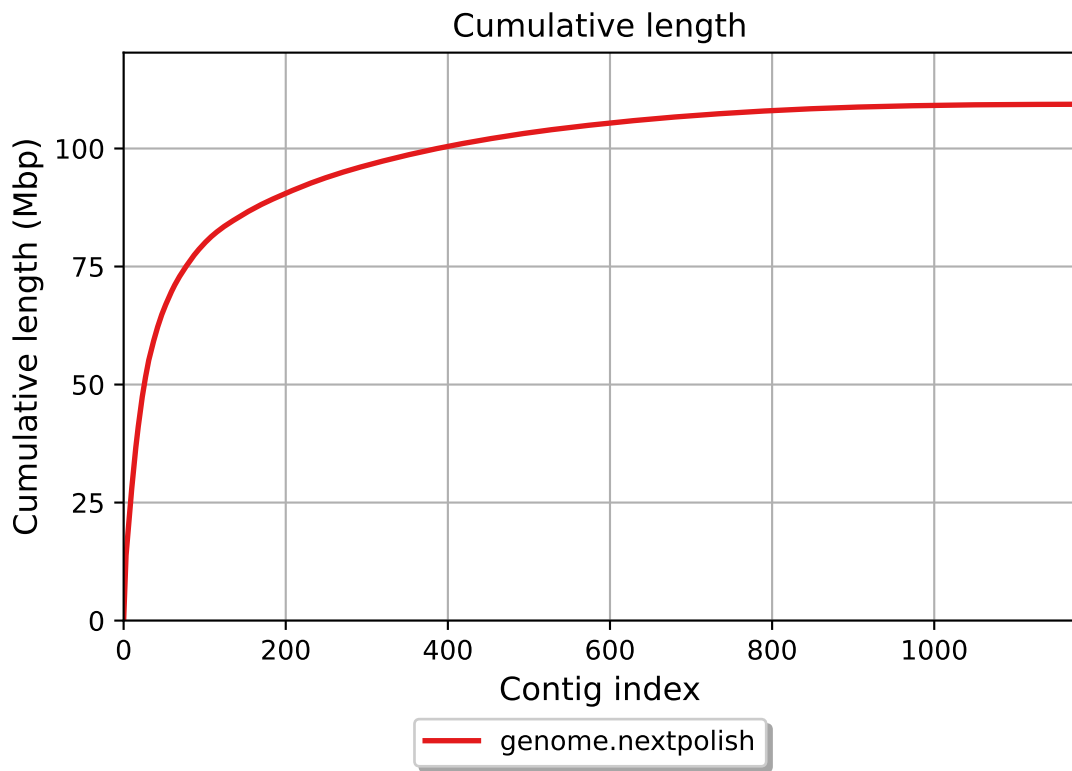
	genome.nextpolish
# contigs (>= 0 bp)	1191
# contigs (>= 1000 bp)	1092
# contigs (>= 5000 bp)	910
# contigs (>= 10000 bp)	770
# contigs (>= 25000 bp)	497
# contigs (>= 50000 bp)	280
Total length (>= 0 bp)	109374931
Total length (>= 1000 bp)	109311686
Total length (>= 5000 bp)	108798490
Total length (>= 10000 bp)	107746476
Total length (>= 25000 bp)	103269167
Total length (>= 50000 bp)	95478216
# contigs	1175
Largest contig	5225517
Total length	109369354
GC (%)	46.12
N50	812711
N90	40586
auN	1303945.7
L50	31
L90	346
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	273 + 14 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).

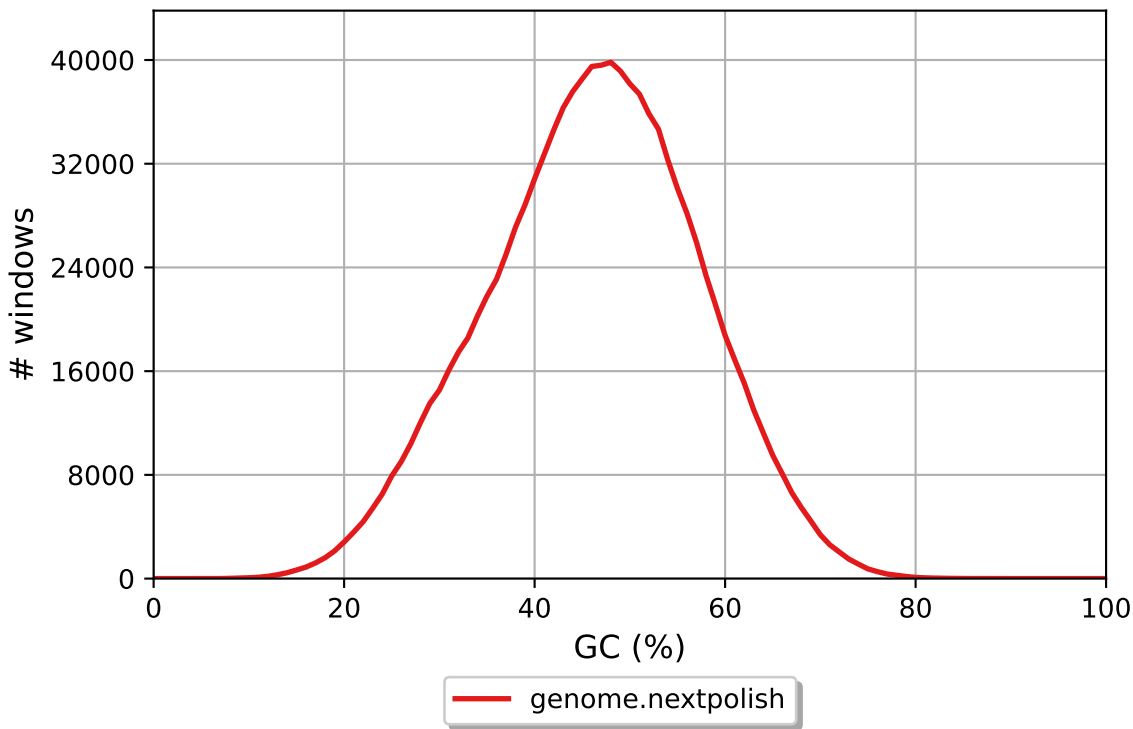
Nx



genome.nextpolish



GC content



genome.nextpolish GC content

