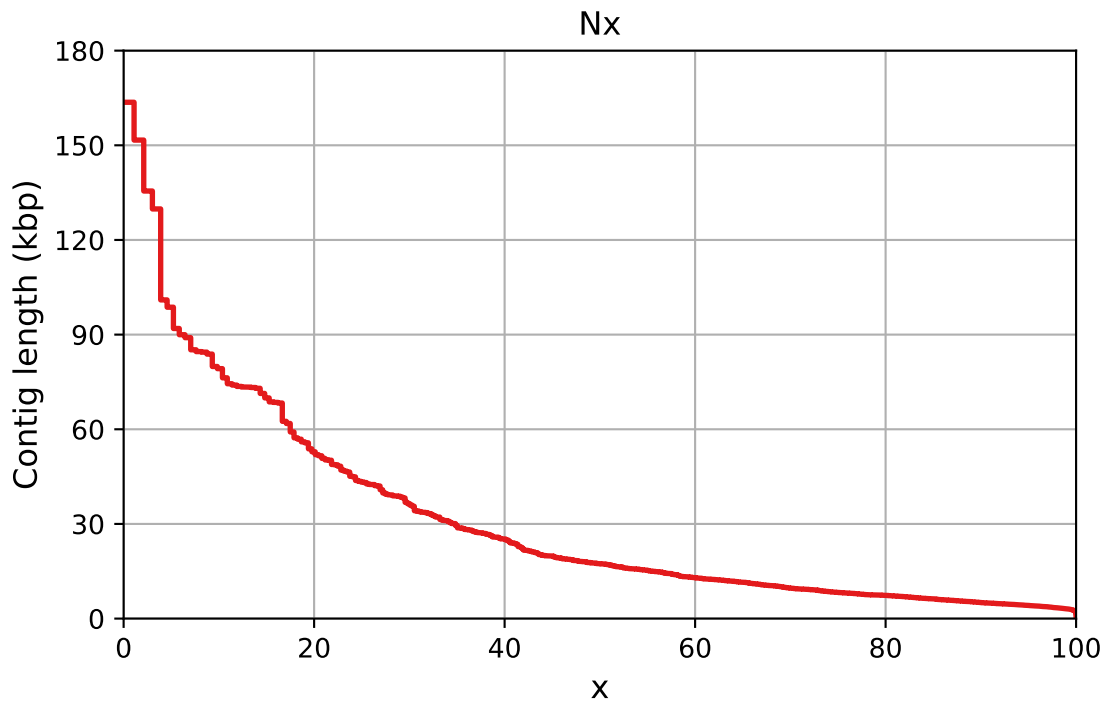


Report

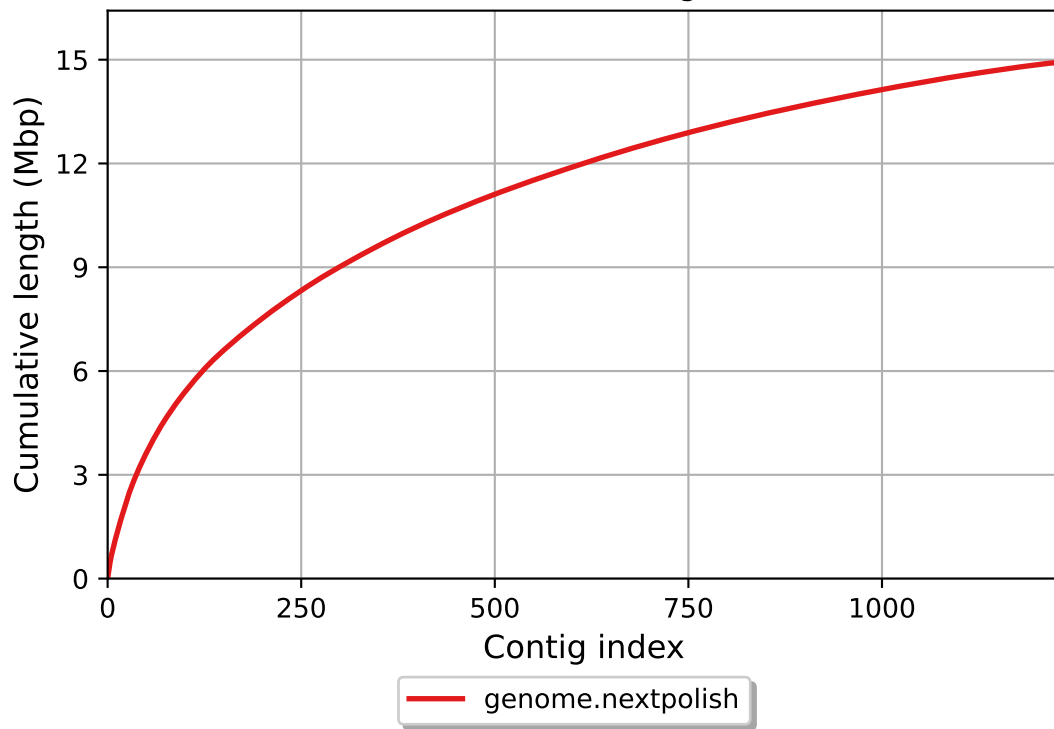
	genome.nextpolish
# contigs (>= 0 bp)	1230
# contigs (>= 1000 bp)	1228
# contigs (>= 5000 bp)	858
# contigs (>= 10000 bp)	415
# contigs (>= 25000 bp)	122
# contigs (>= 50000 bp)	42
Total length (>= 0 bp)	14925709
Total length (>= 1000 bp)	14924472
Total length (>= 5000 bp)	13481517
Total length (>= 10000 bp)	10331461
Total length (>= 25000 bp)	5986625
Total length (>= 50000 bp)	3257391
# contigs	1230
Largest contig	163636
Total length	14925709
GC (%)	45.03
N50	17377
N90	5098
auN	31846.5
L50	197
L90	849
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	31 + 5 part

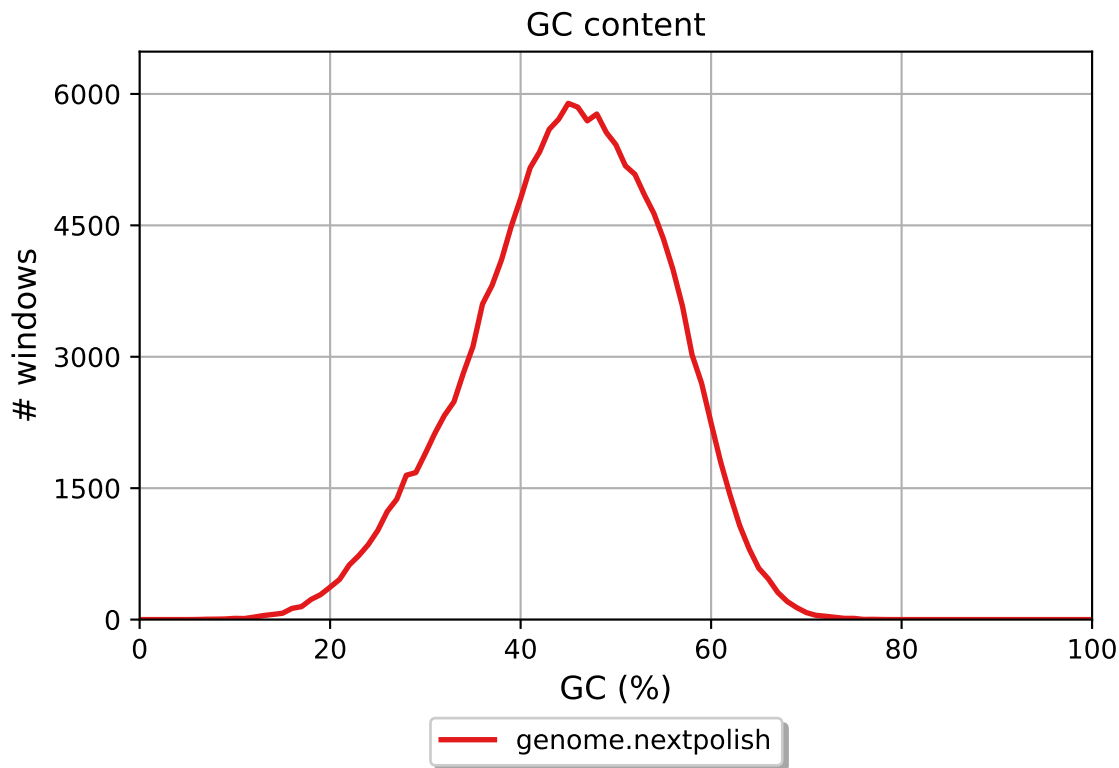
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

