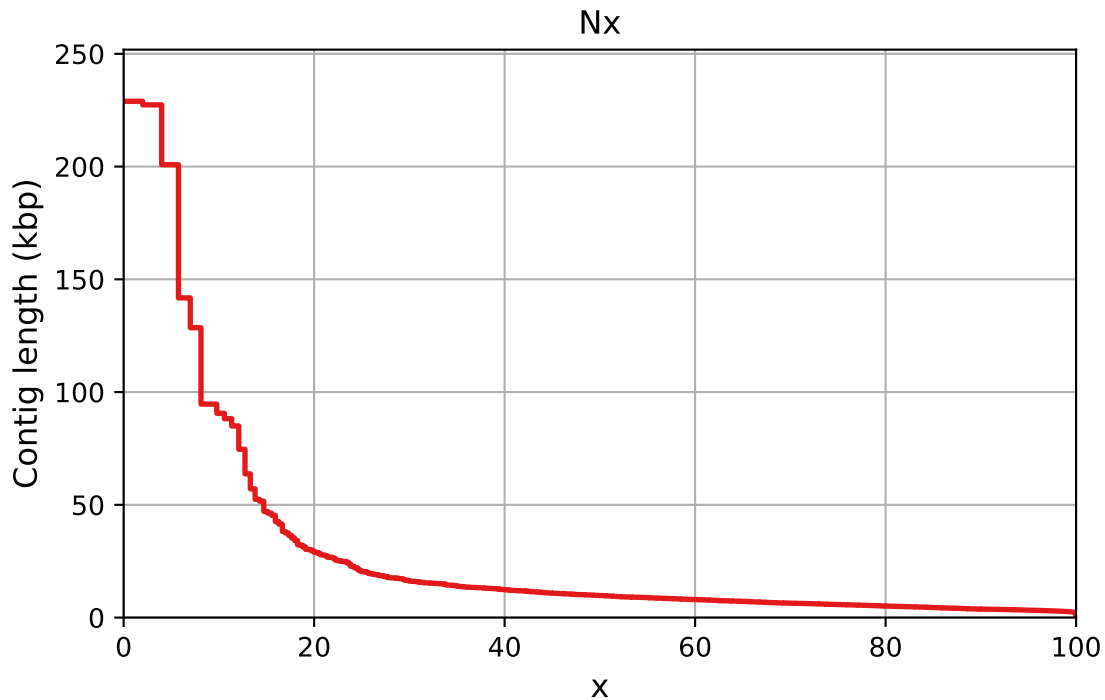


Report

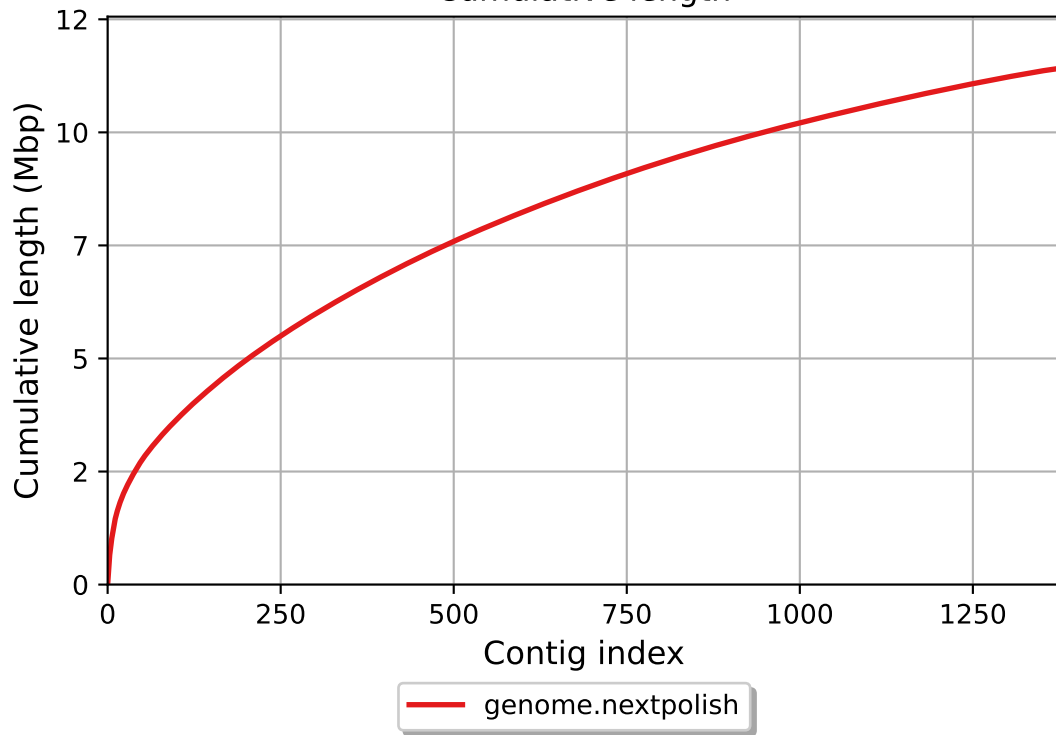
	genome.nextpolish
# contigs (>= 0 bp)	1376
# contigs (>= 1000 bp)	1375
# contigs (>= 5000 bp)	779
# contigs (>= 10000 bp)	258
# contigs (>= 25000 bp)	44
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	11417289
Total length (>= 1000 bp)	11416463
Total length (>= 5000 bp)	9236831
Total length (>= 10000 bp)	5578962
Total length (>= 25000 bp)	2621504
Total length (>= 50000 bp)	1680059
# contigs	1376
Largest contig	228953
Total length	11417289
GC (%)	41.33
N50	9764
N90	3766
auN	30700.8
L50	272
L90	1018
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 1 part

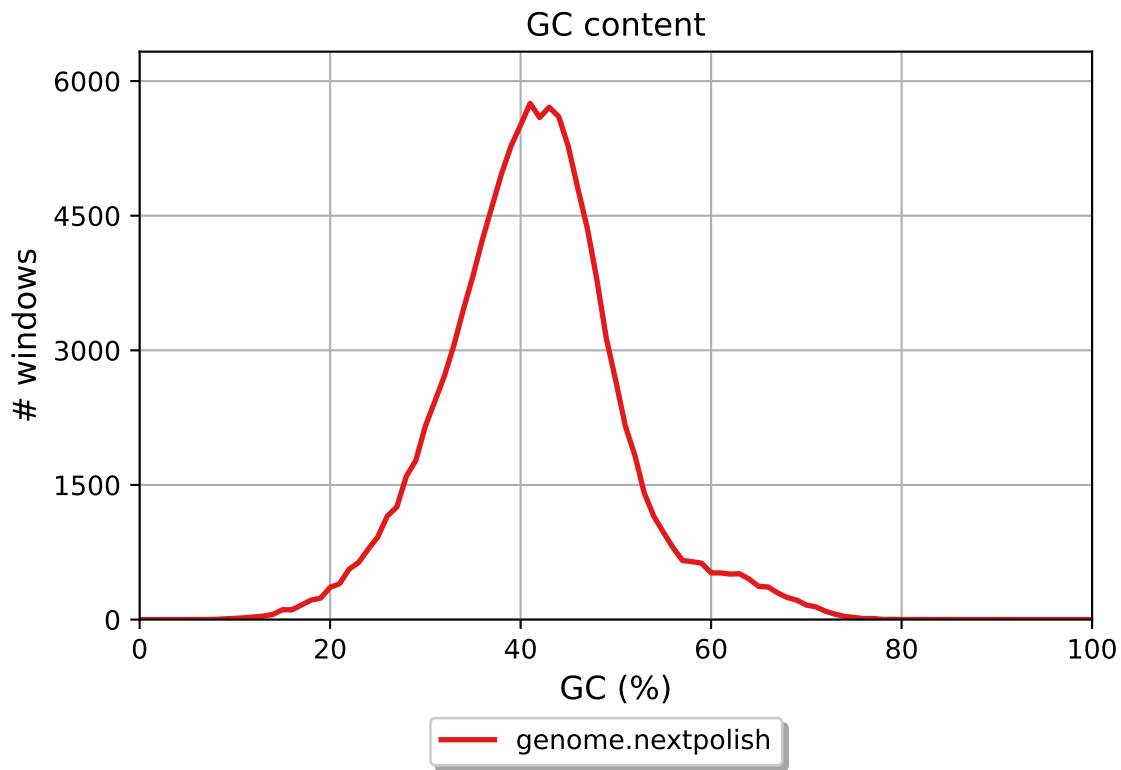
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

