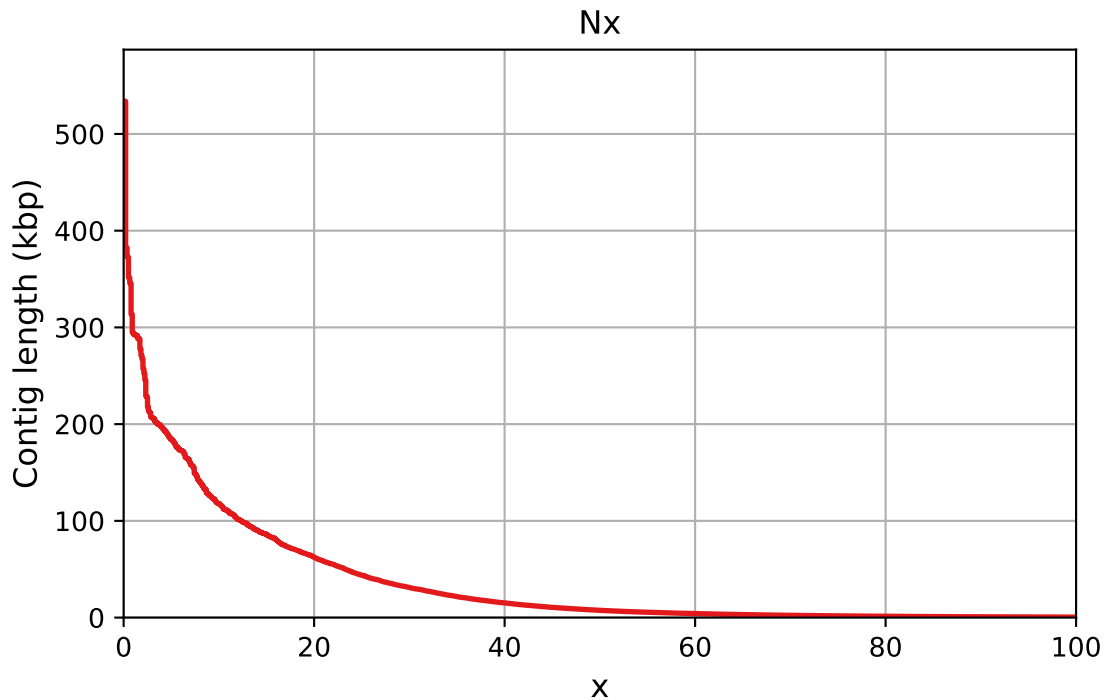


Report

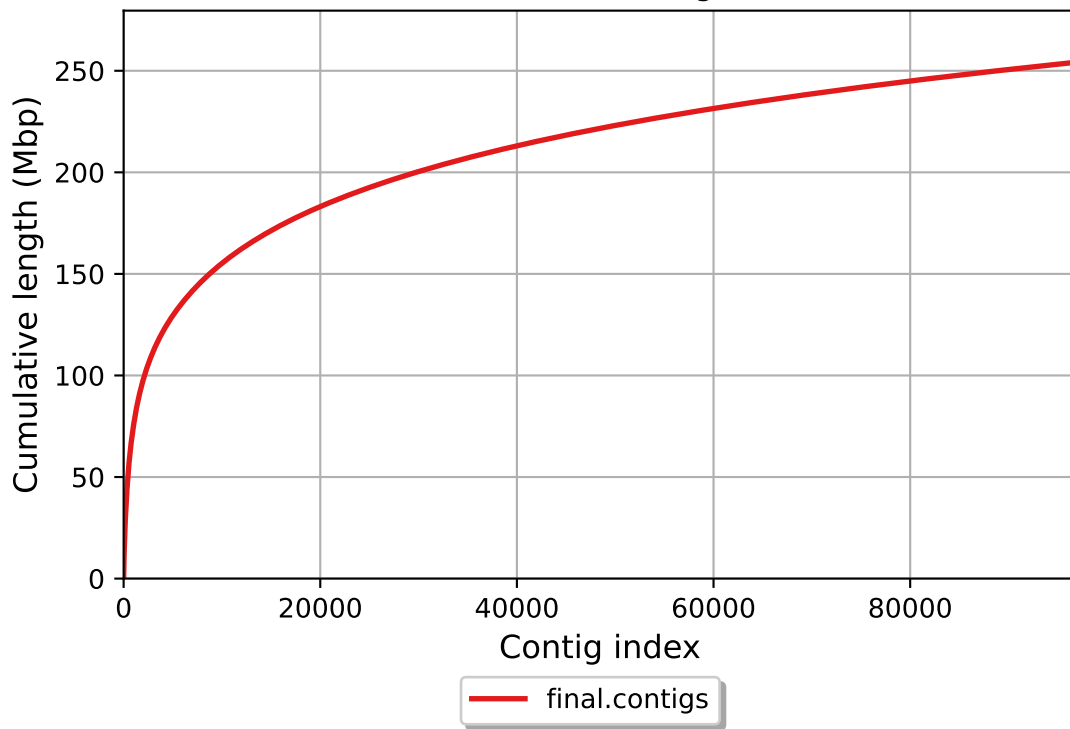
	final.contigs
# contigs (>= 0 bp)	164368
# contigs (>= 1000 bp)	44989
# contigs (>= 5000 bp)	7509
# contigs (>= 10000 bp)	3441
# contigs (>= 25000 bp)	1317
# contigs (>= 50000 bp)	591
Total length (>= 0 bp)	279115600
Total length (>= 1000 bp)	218313388
Total length (>= 5000 bp)	144281871
Total length (>= 10000 bp)	116473543
Total length (>= 25000 bp)	84182573
Total length (>= 50000 bp)	59114209
# contigs	96876
Largest contig	533782
Total length	254186606
GC (%)	46.43
N50	7517
N90	813
auN	38107.1
L50	4670
L90	56611
# N's per 100 kbp	0.00
# predicted genes (unique)	192861
# predicted genes (>= 0 bp)	174126 + 18842 part
# predicted genes (>= 300 bp)	155732 + 17476 part
# predicted genes (>= 1500 bp)	29108 + 1084 part
# predicted genes (>= 3000 bp)	5682 + 125 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).

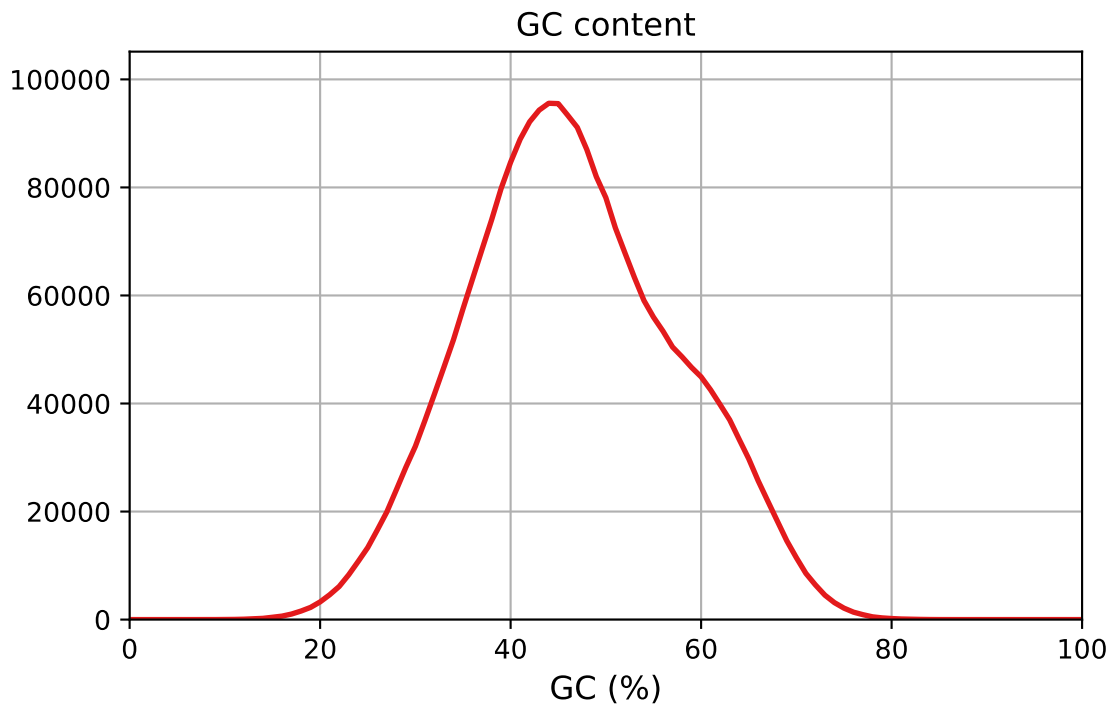


— final.contigs

Cumulative length



windows



— final.contigs

final.contigs GC content

