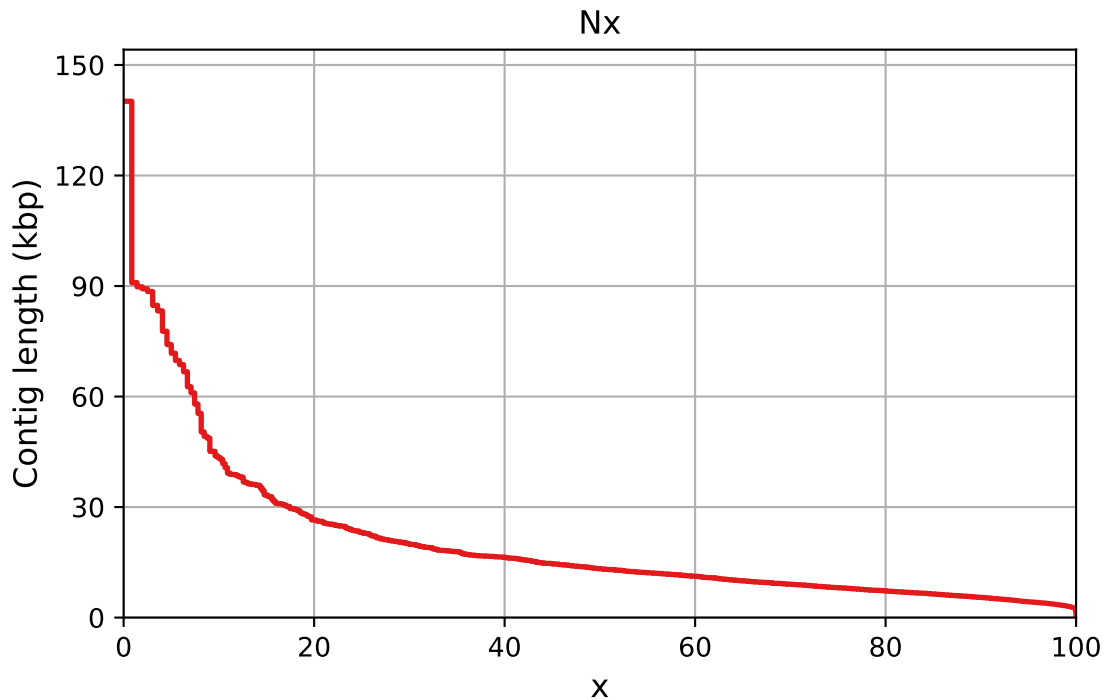


# Report

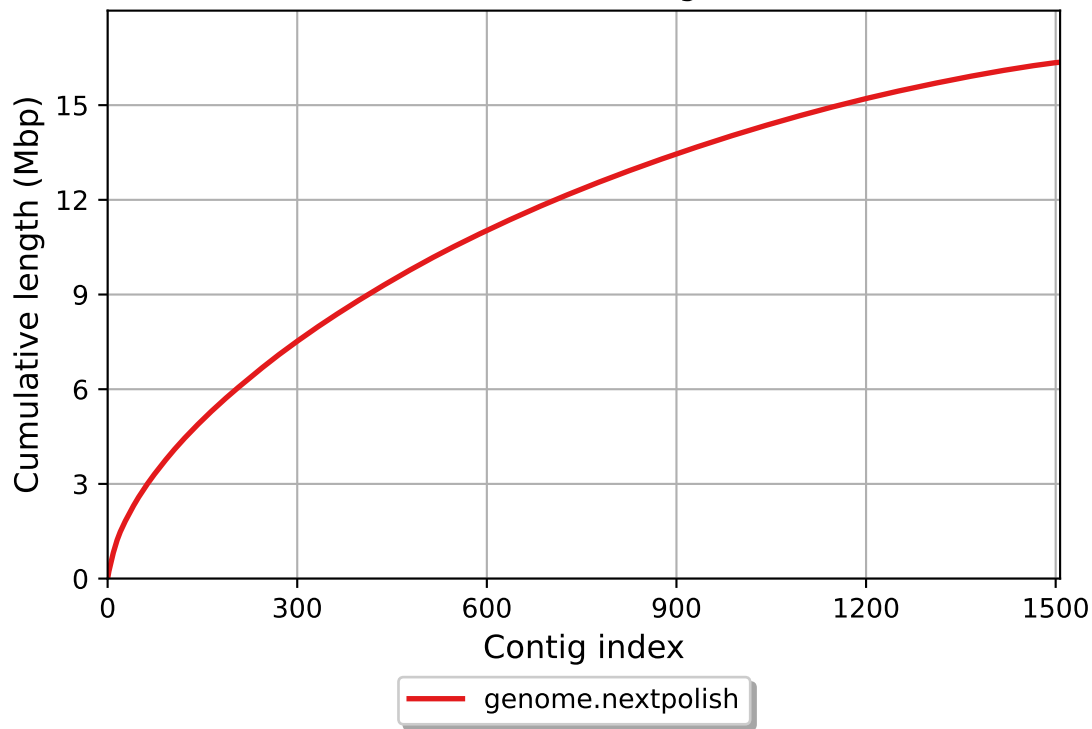
	genome.nextpolish
# contigs (>= 0 bp)	1507
# contigs (>= 1000 bp)	1505
# contigs (>= 5000 bp)	1174
# contigs (>= 10000 bp)	558
# contigs (>= 25000 bp)	88
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	16358664
Total length (>= 1000 bp)	16357531
Total length (>= 5000 bp)	15081417
Total length (>= 10000 bp)	10620693
Total length (>= 25000 bp)	3664876
Total length (>= 50000 bp)	1382820
# contigs	1507
Largest contig	140152
Total length	16358664
GC (%)	55.76
N50	13249
N90	5482
auN	20689.9
L50	348
L90	1106
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	25 + 4 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).

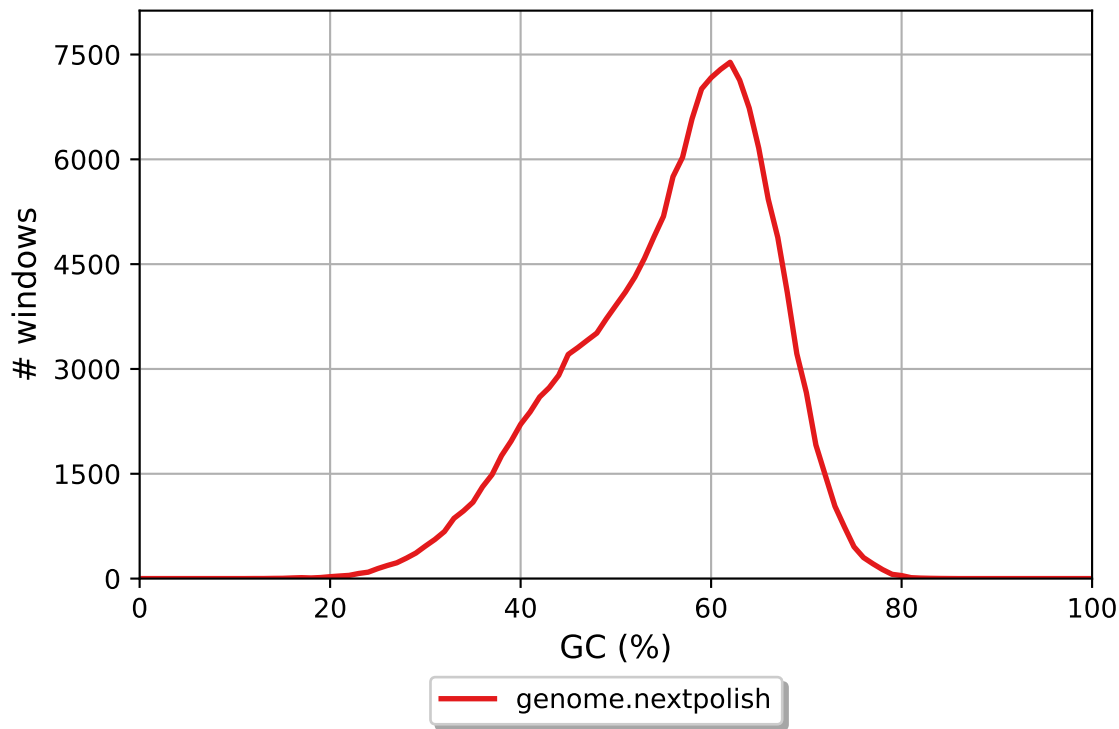


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

