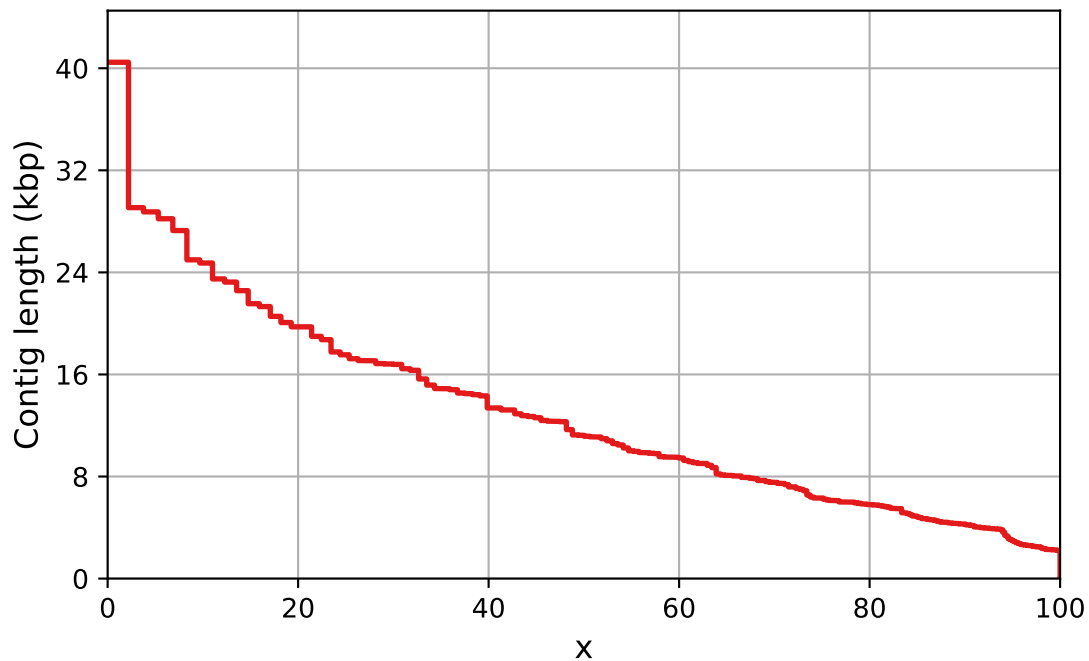


Report

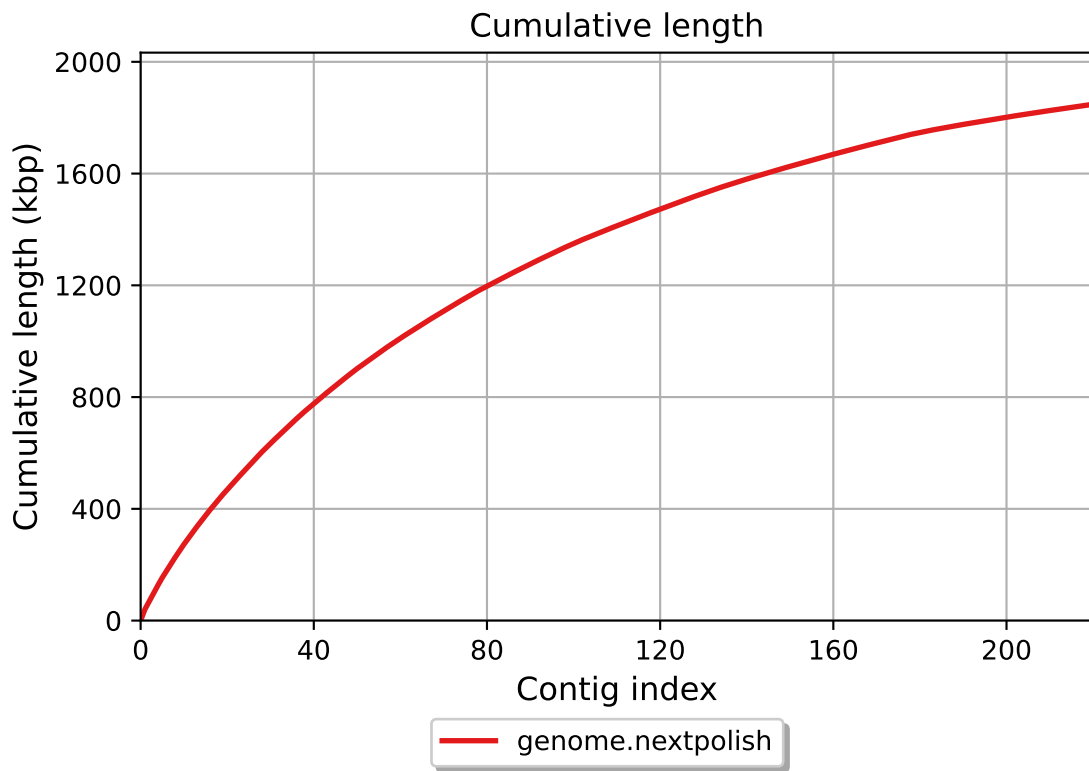
	genome.nextpolish
# contigs (>= 0 bp)	220
# contigs (>= 1000 bp)	220
# contigs (>= 5000 bp)	136
# contigs (>= 10000 bp)	61
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1848082
Total length (>= 1000 bp)	1848082
Total length (>= 5000 bp)	1560868
Total length (>= 10000 bp)	1020693
Total length (>= 25000 bp)	153771
Total length (>= 50000 bp)	0
# contigs	220
Largest contig	40469
Total length	1848082
GC (%)	38.53
N50	11230
N90	4274
auN	13136.6
L50	52
L90	159
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 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).

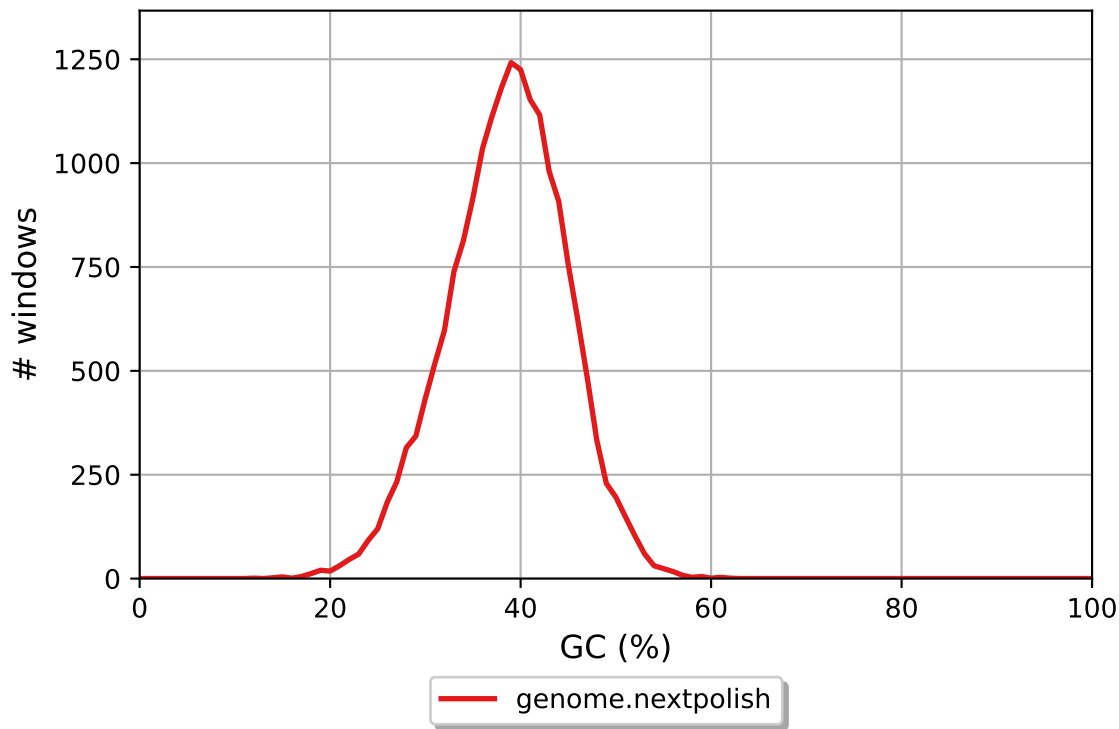
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

