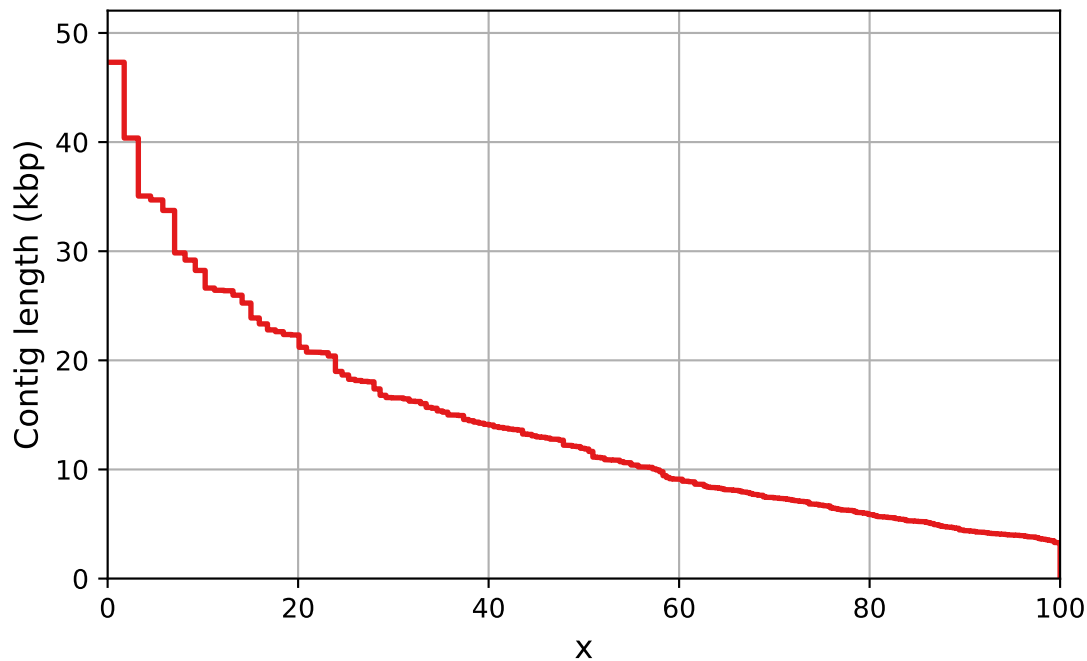


Report

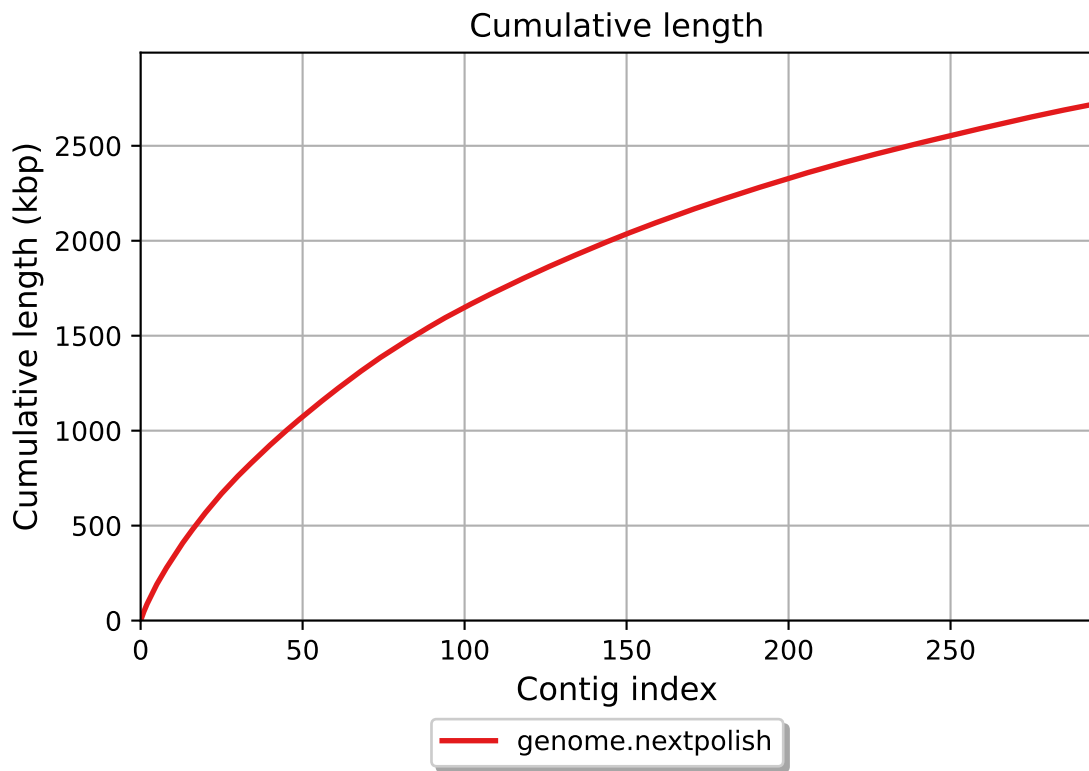
	genome.nextpolish
# contigs (>= 0 bp)	294
# contigs (>= 1000 bp)	294
# contigs (>= 5000 bp)	206
# contigs (>= 10000 bp)	91
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2718672
Total length (>= 1000 bp)	2718672
Total length (>= 5000 bp)	2359056
Total length (>= 10000 bp)	1565505
Total length (>= 25000 bp)	409081
Total length (>= 50000 bp)	0
# contigs	294
Largest contig	47314
Total length	2718672
GC (%)	57.27
N50	11942
N90	4394
auN	14258.7
L50	72
L90	225
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

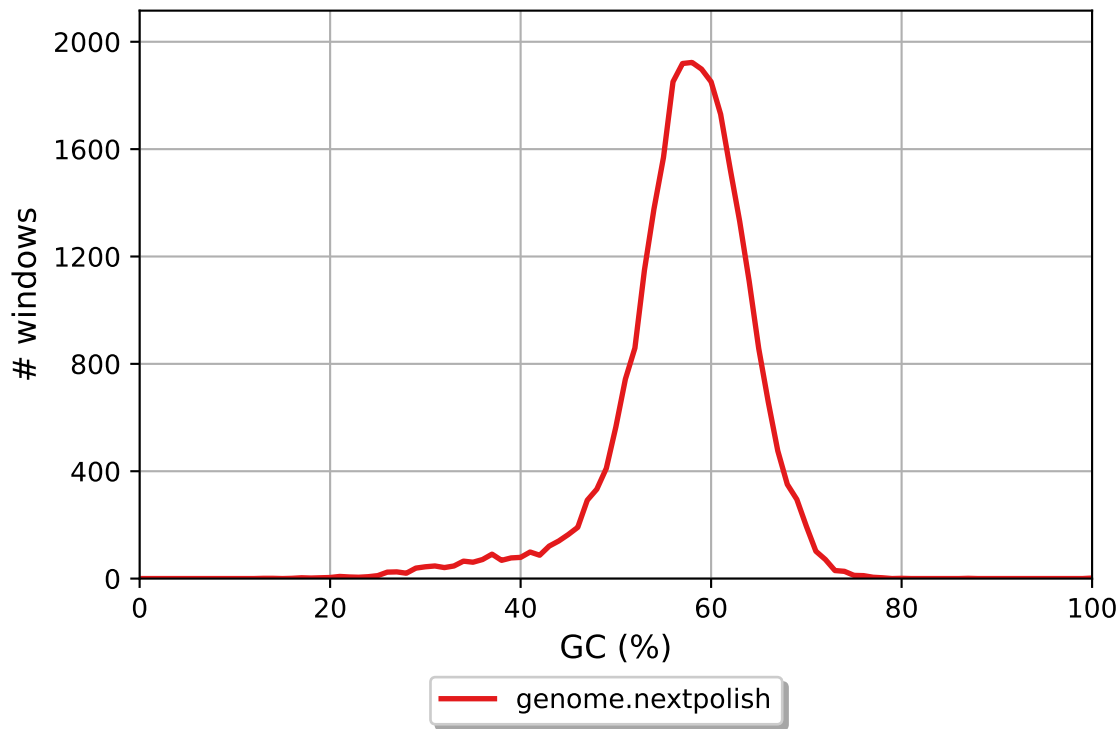
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

