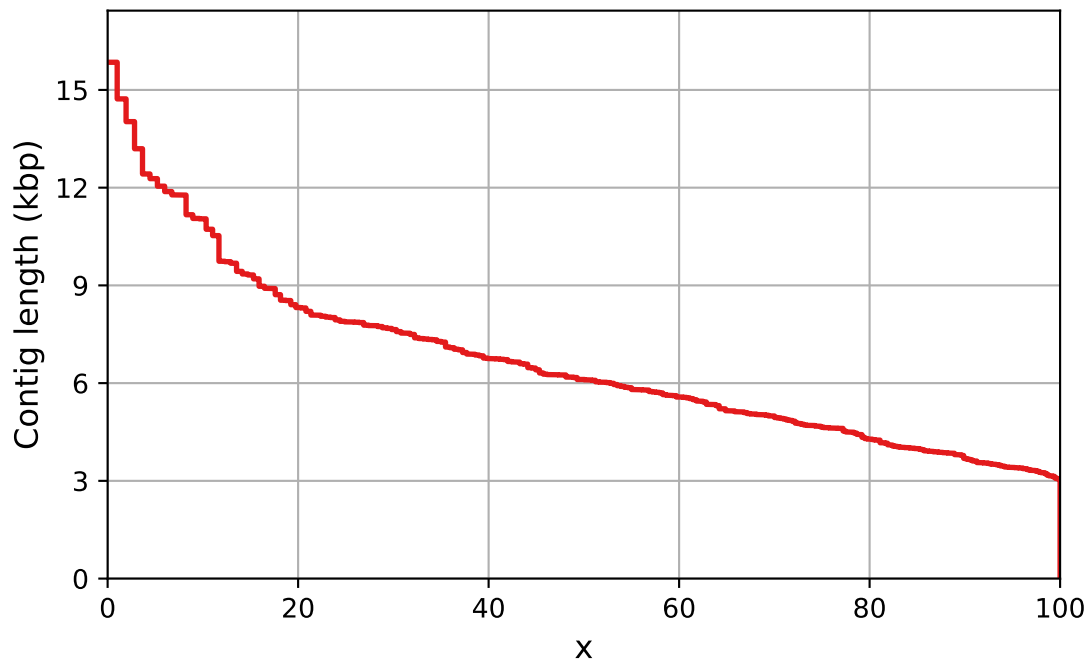


Report

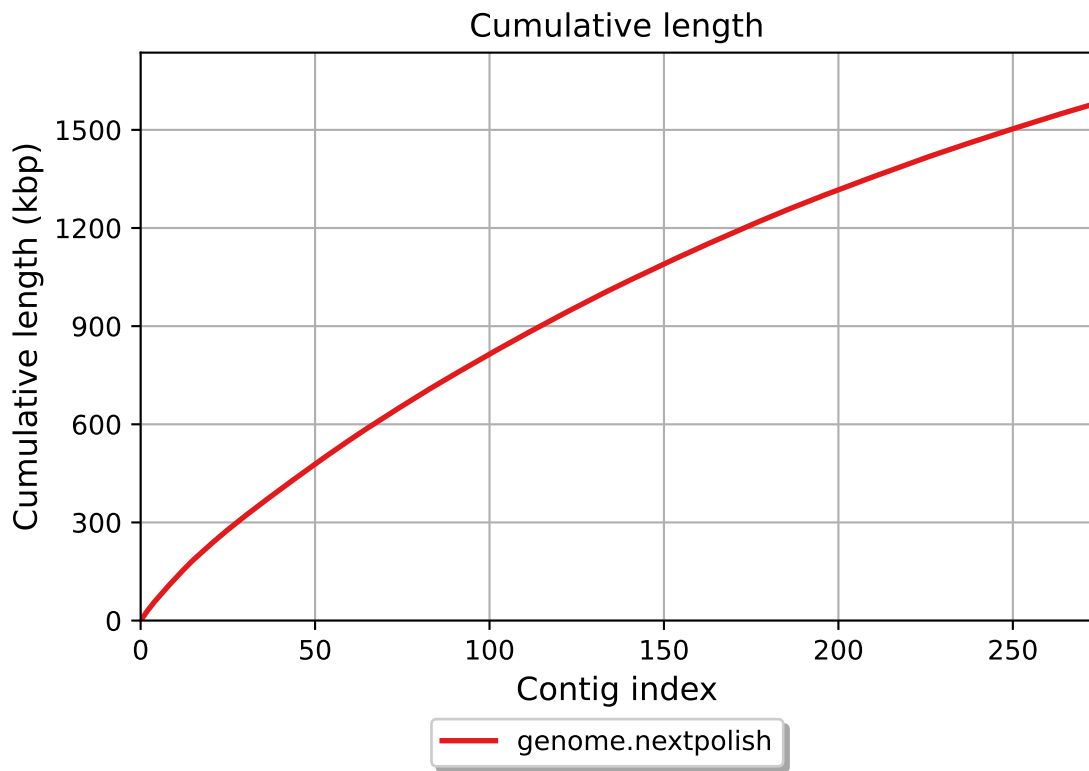
	genome.nextpolish
# contigs (>= 0 bp)	273
# contigs (>= 1000 bp)	273
# contigs (>= 5000 bp)	151
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1578243
Total length (>= 1000 bp)	1578243
Total length (>= 5000 bp)	1094946
Total length (>= 10000 bp)	184487
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	273
Largest contig	15849
Total length	1578243
GC (%)	51.44
N50	6112
N90	3690
auN	6686.9
L50	96
L90	227
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

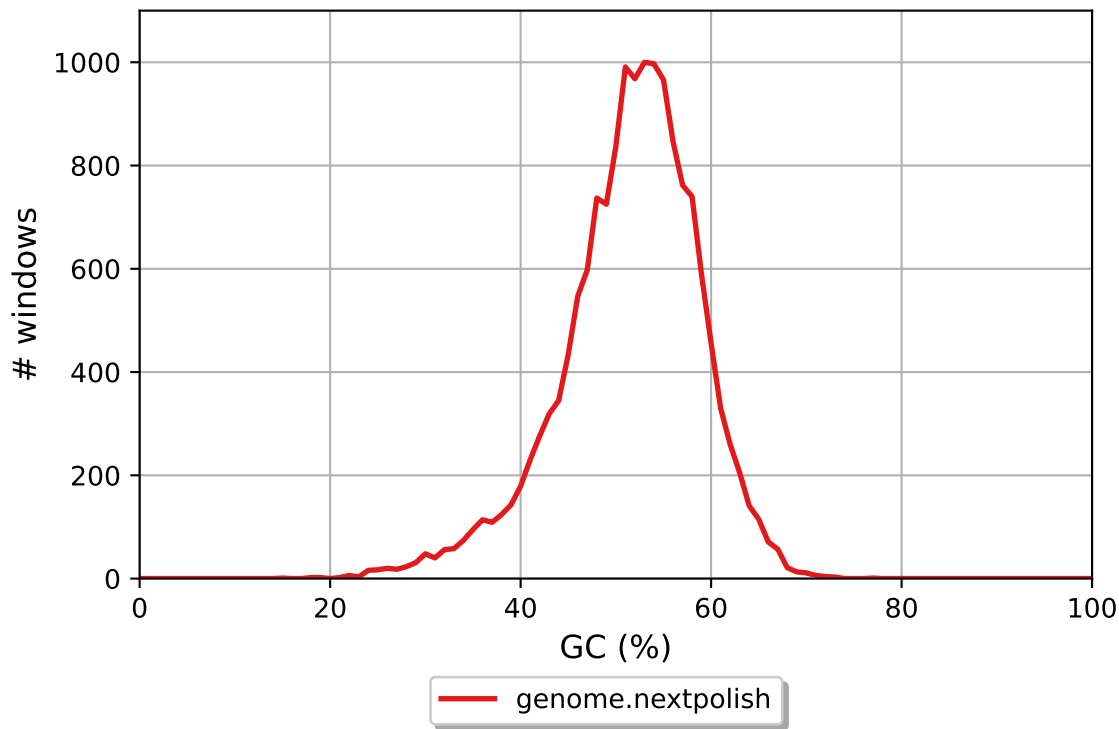
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

