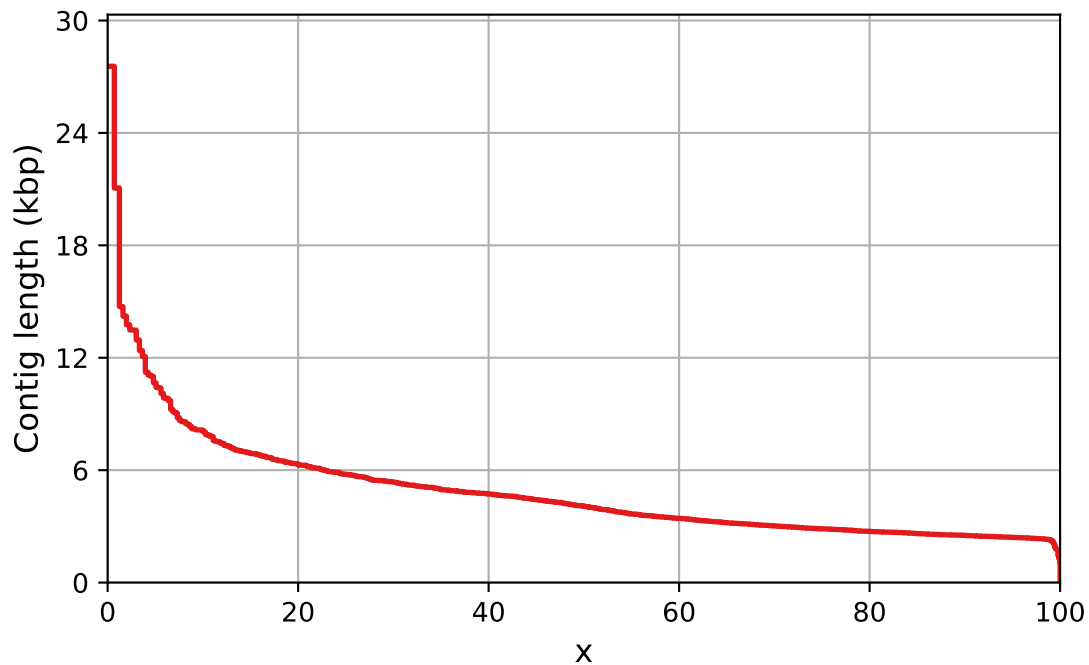


Report

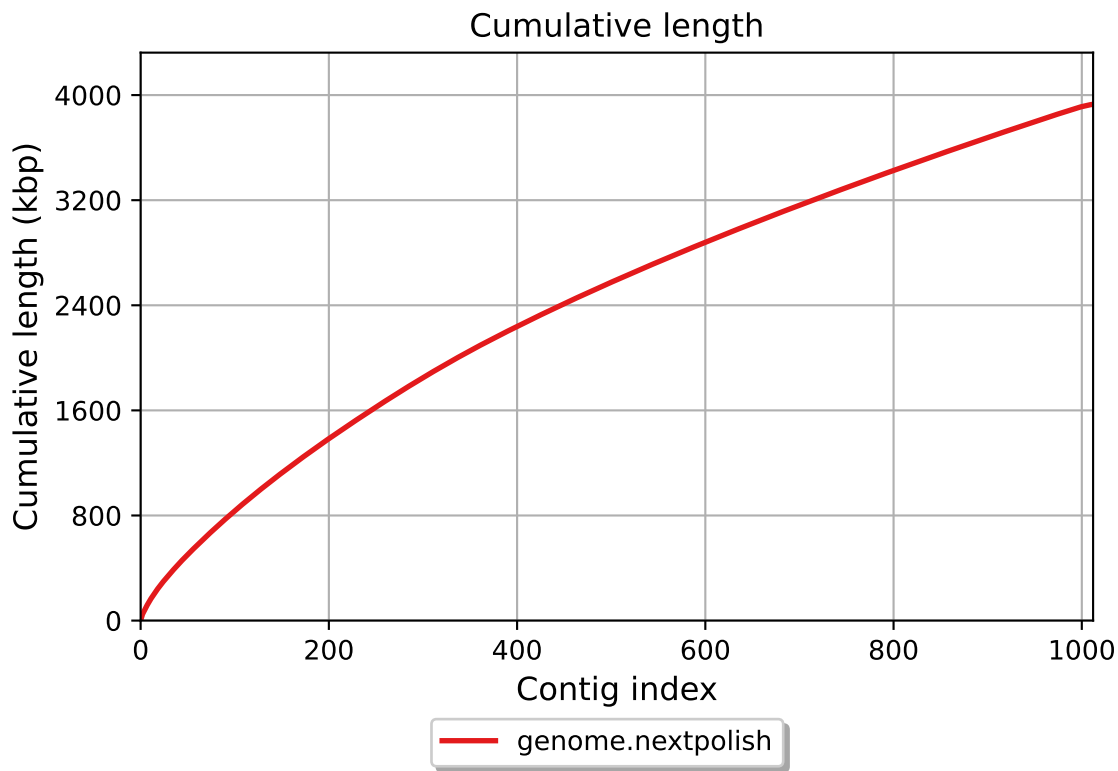
	genome.nextpolish
# contigs (≥ 0 bp)	1012
# contigs (≥ 1000 bp)	1012
# contigs (≥ 5000 bp)	197
# contigs (≥ 10000 bp)	17
# contigs (≥ 25000 bp)	1
# contigs (≥ 50000 bp)	0
Total length (≥ 0 bp)	3930081
Total length (≥ 1000 bp)	3930081
Total length (≥ 5000 bp)	1369799
Total length (≥ 10000 bp)	230550
Total length (≥ 25000 bp)	27557
Total length (≥ 50000 bp)	0
# contigs	1012
Largest contig	27557
Total length	3930081
GC (%)	45.65
N50	4080
N90	2518
auN	4936.8
L50	328
L90	844
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 11 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).

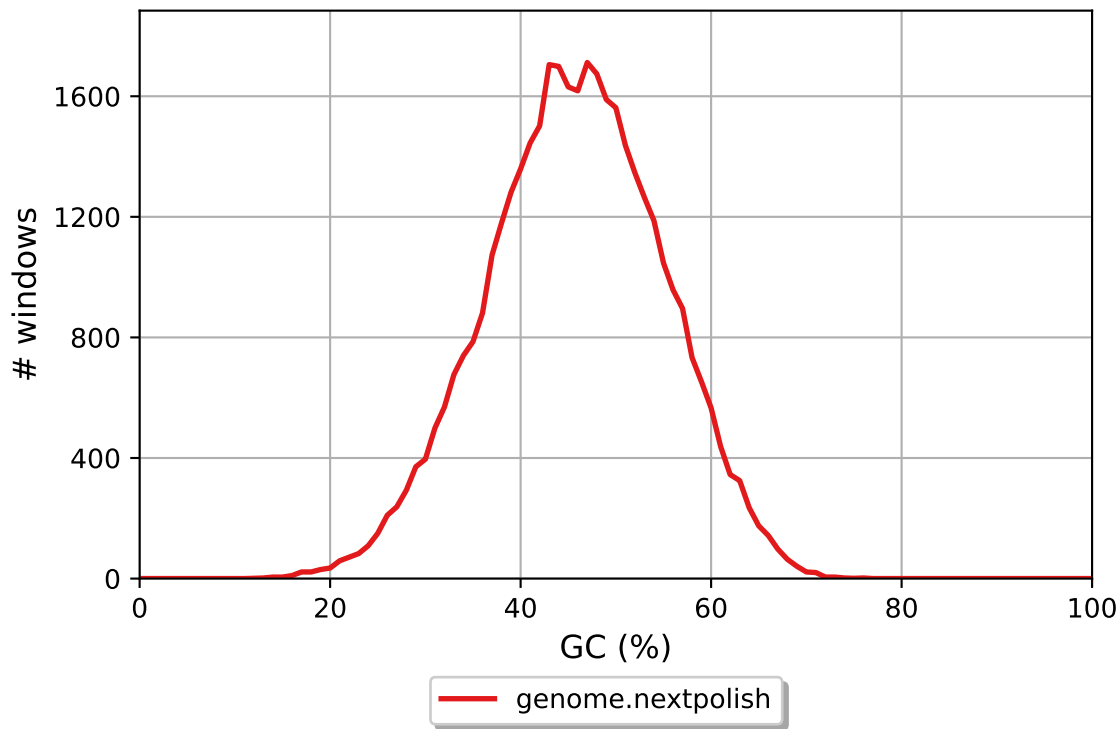
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

