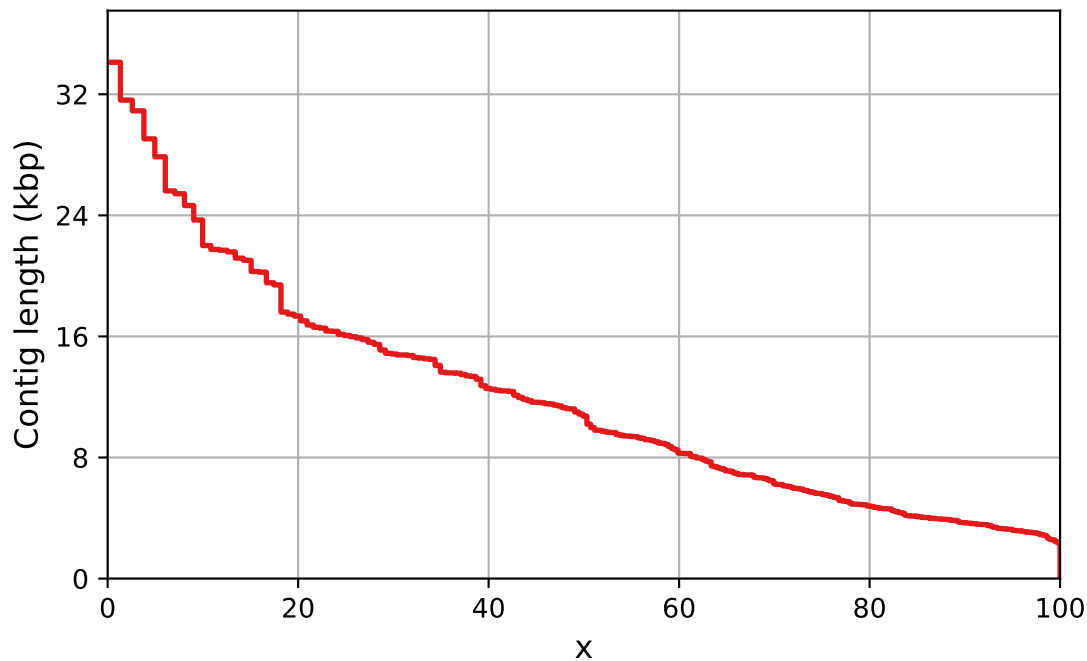


Report

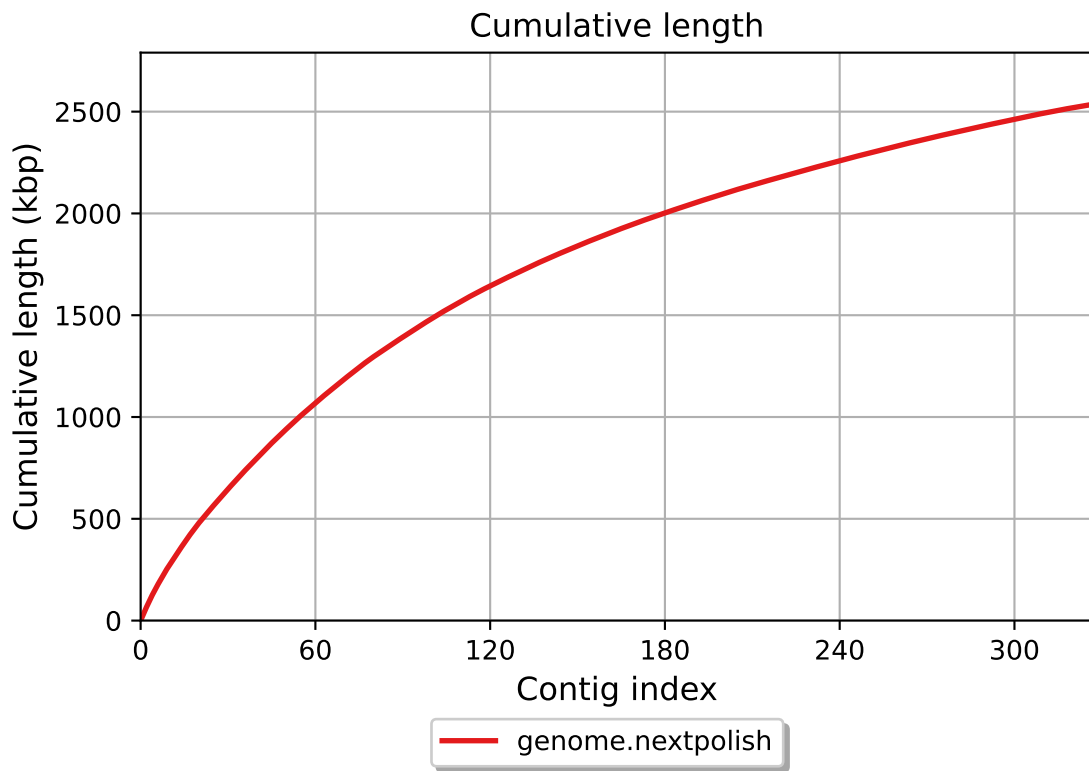
	genome.nextpolish
# contigs (>= 0 bp)	327
# contigs (>= 1000 bp)	327
# contigs (>= 5000 bp)	175
# contigs (>= 10000 bp)	79
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2536212
Total length (>= 1000 bp)	2536212
Total length (>= 5000 bp)	1977795
Total length (>= 10000 bp)	1286384
Total length (>= 25000 bp)	204603
Total length (>= 50000 bp)	0
# contigs	327
Largest contig	34114
Total length	2536212
GC (%)	34.44
N50	10734
N90	3695
auN	12015.5
L50	78
L90	247
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 3 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).

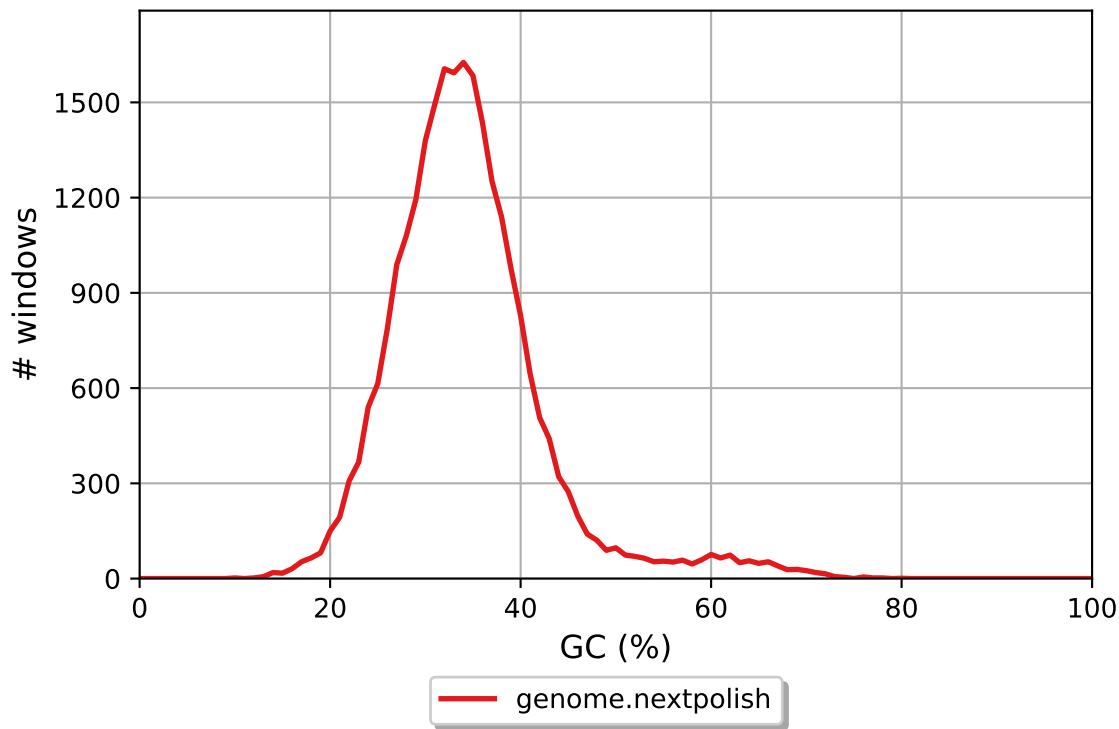
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

