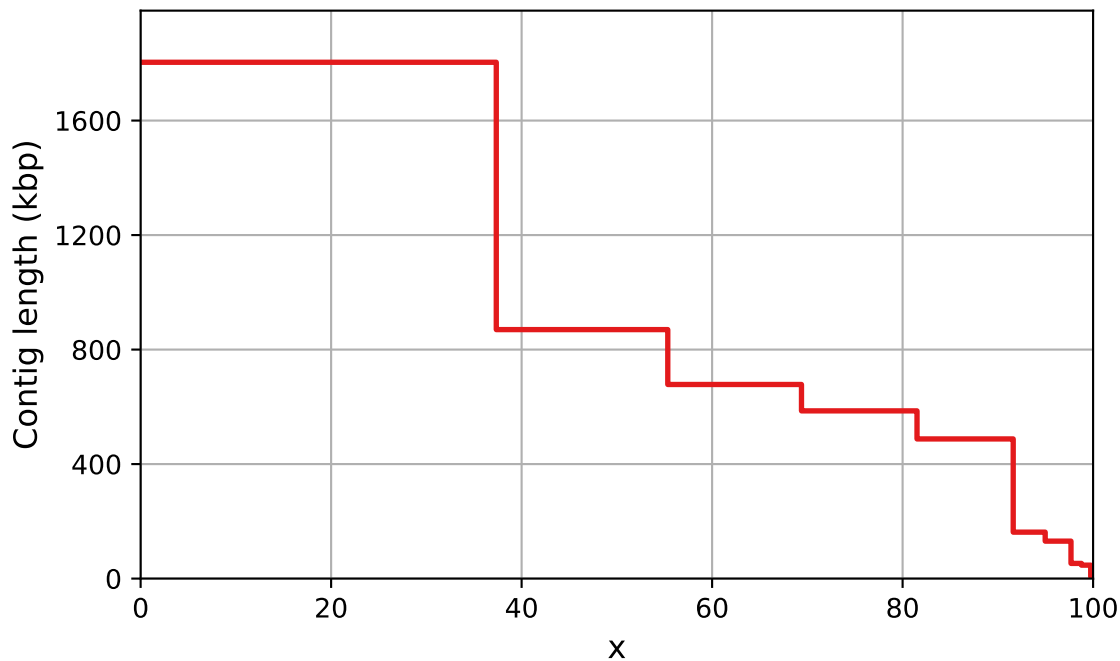


Report

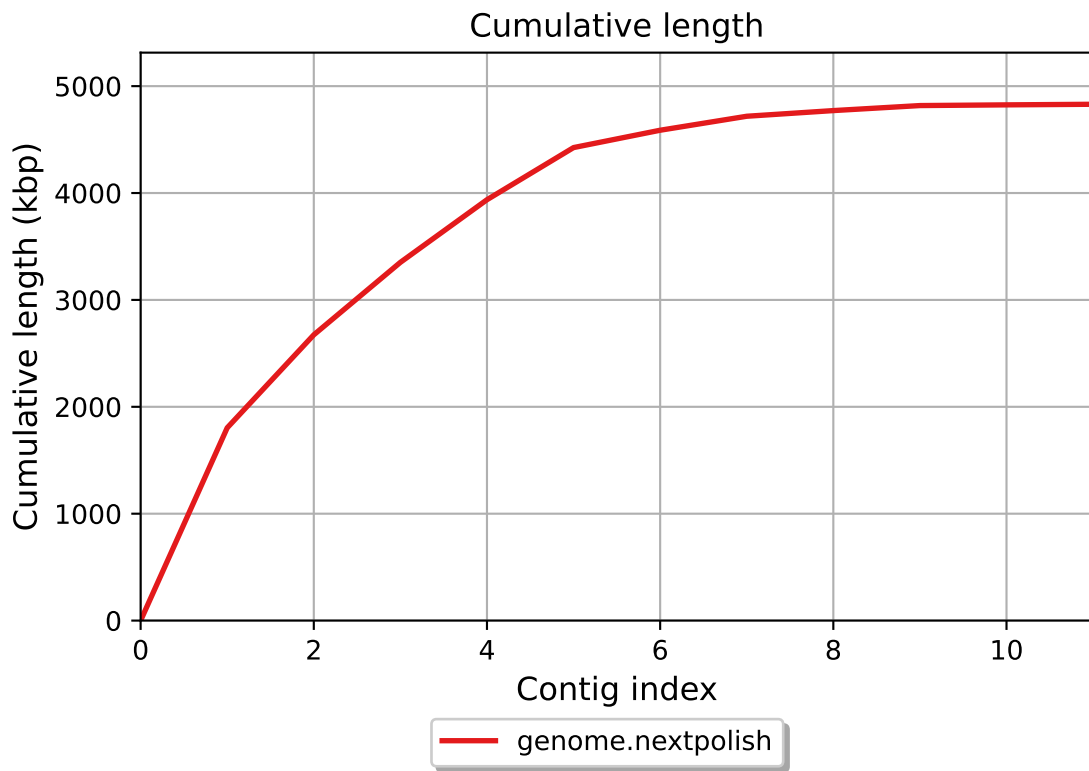
	genome.nextpolish
# contigs (>= 0 bp)	11
# contigs (>= 1000 bp)	11
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	9
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	4830368
Total length (>= 1000 bp)	4830368
Total length (>= 5000 bp)	4830368
Total length (>= 10000 bp)	4818221
Total length (>= 25000 bp)	4818221
Total length (>= 50000 bp)	4771280
# contigs	11
Largest contig	1803648
Total length	4830368
GC (%)	53.71
N50	869768
N90	487859
auN	1055654.8
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	22 + 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).

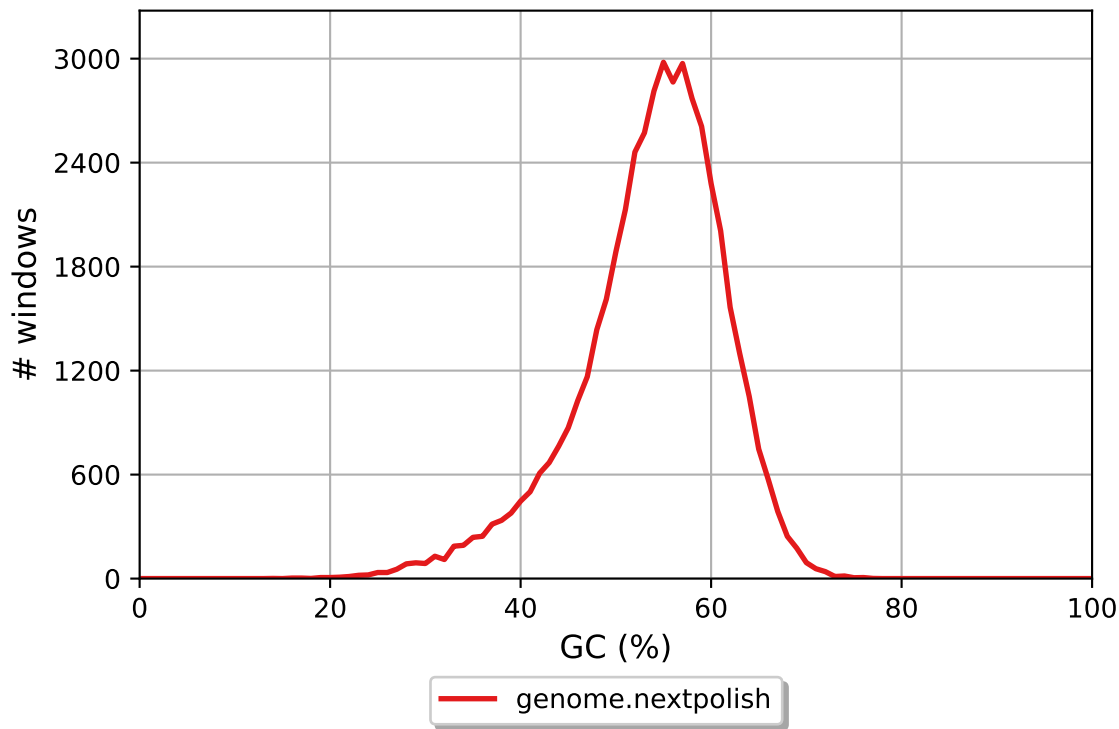
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

