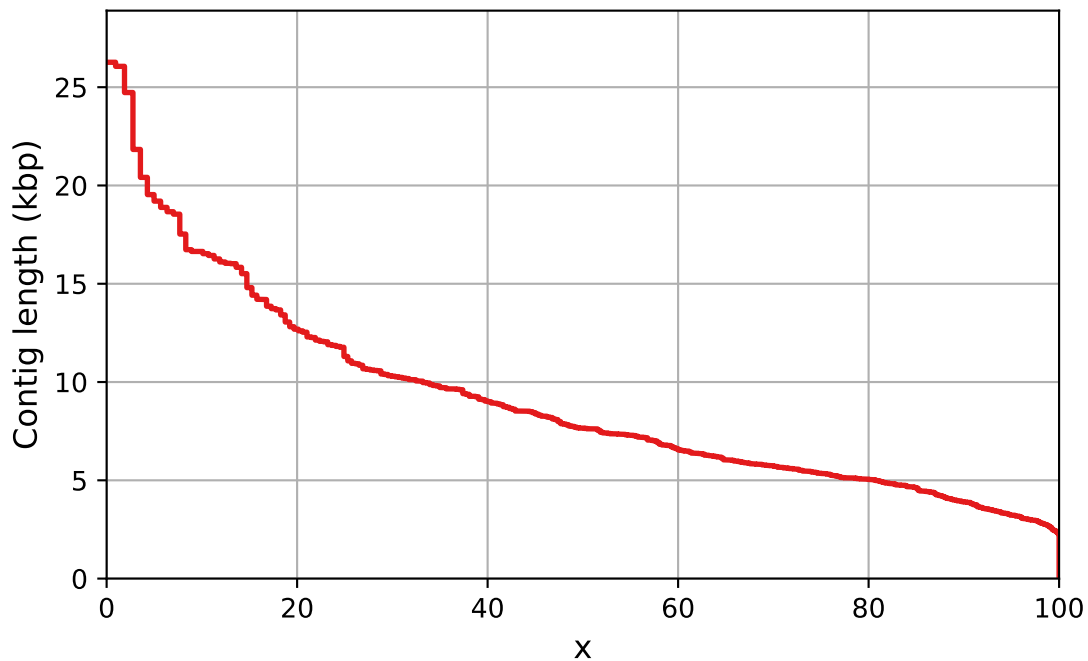


Report

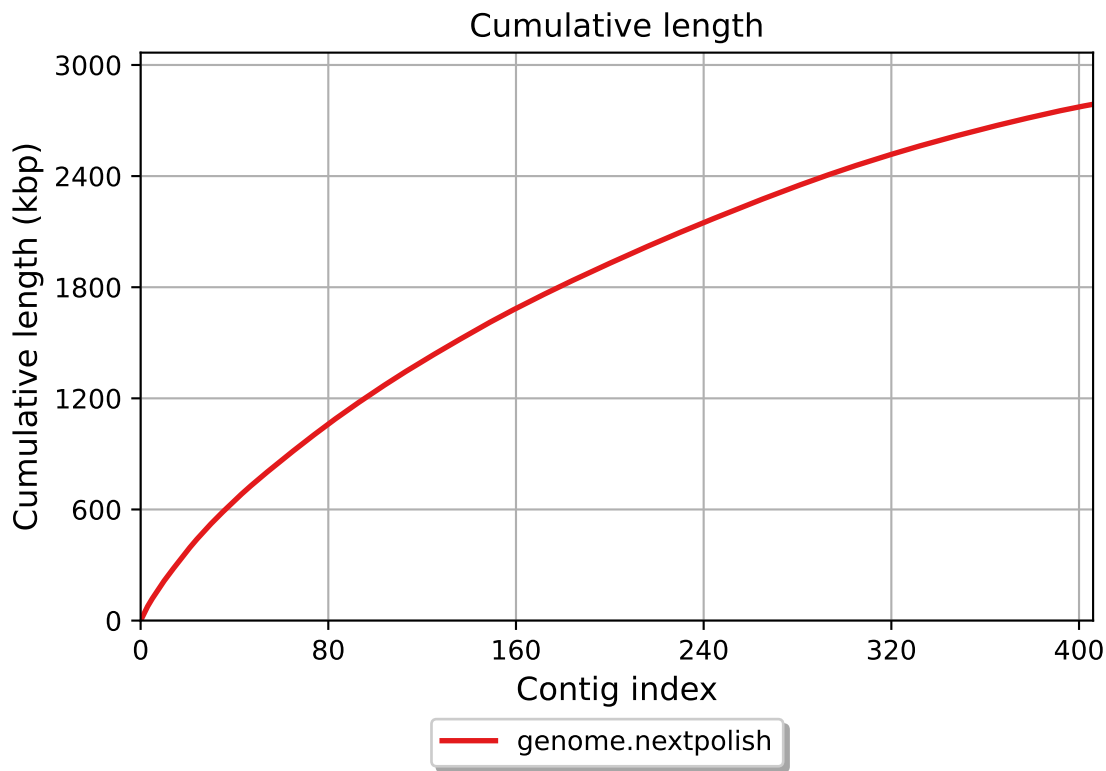
	genome.nextpolish
# contigs (>= 0 bp)	406
# contigs (>= 1000 bp)	406
# contigs (>= 5000 bp)	260
# contigs (>= 10000 bp)	66
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2787845
Total length (>= 1000 bp)	2787845
Total length (>= 5000 bp)	2250702
Total length (>= 10000 bp)	925419
Total length (>= 25000 bp)	52329
Total length (>= 50000 bp)	0
# contigs	406
Largest contig	26268
Total length	2787845
GC (%)	56.19
N50	7655
N90	3904
auN	9149.4
L50	120
L90	318
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

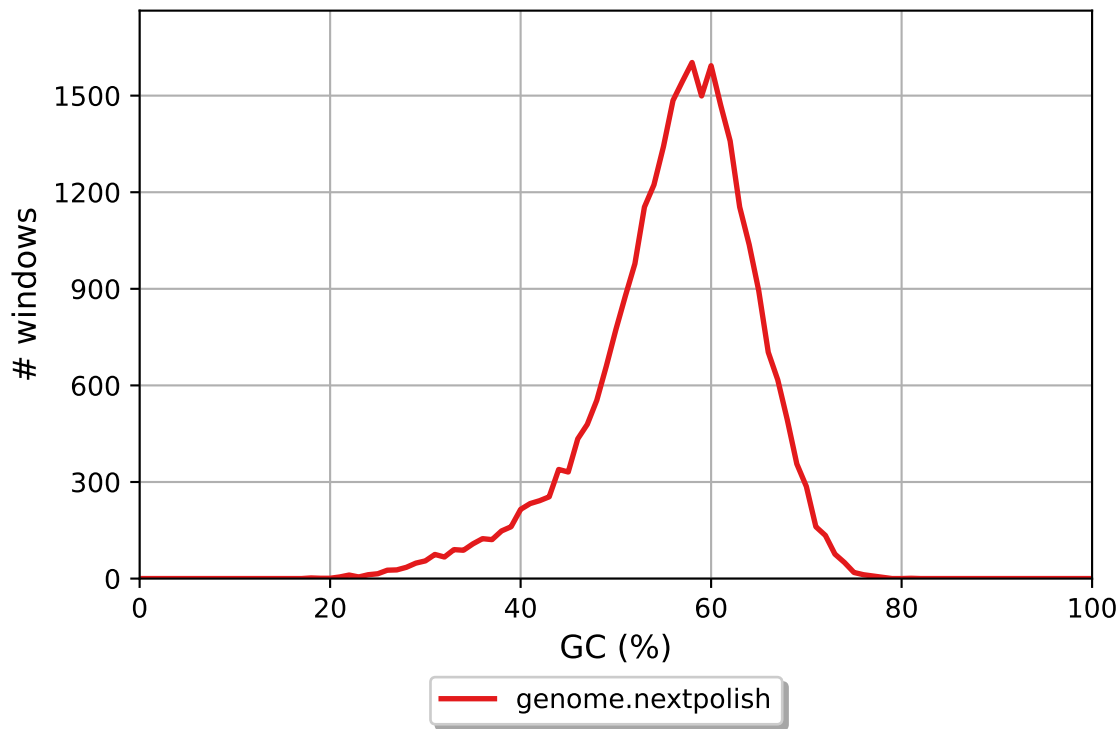
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

