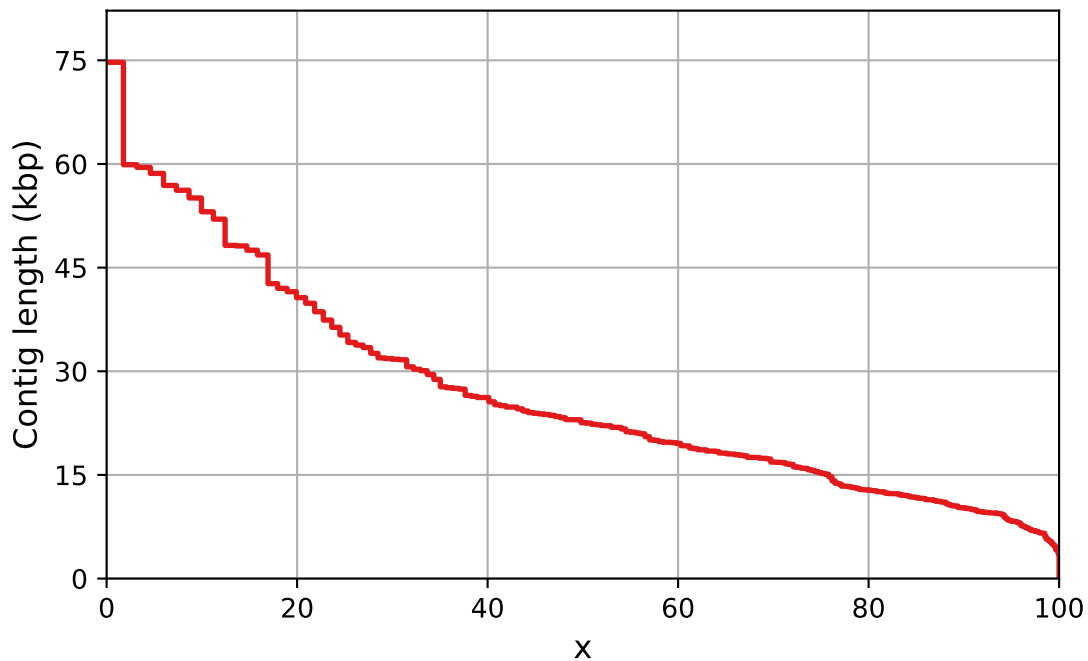


Report

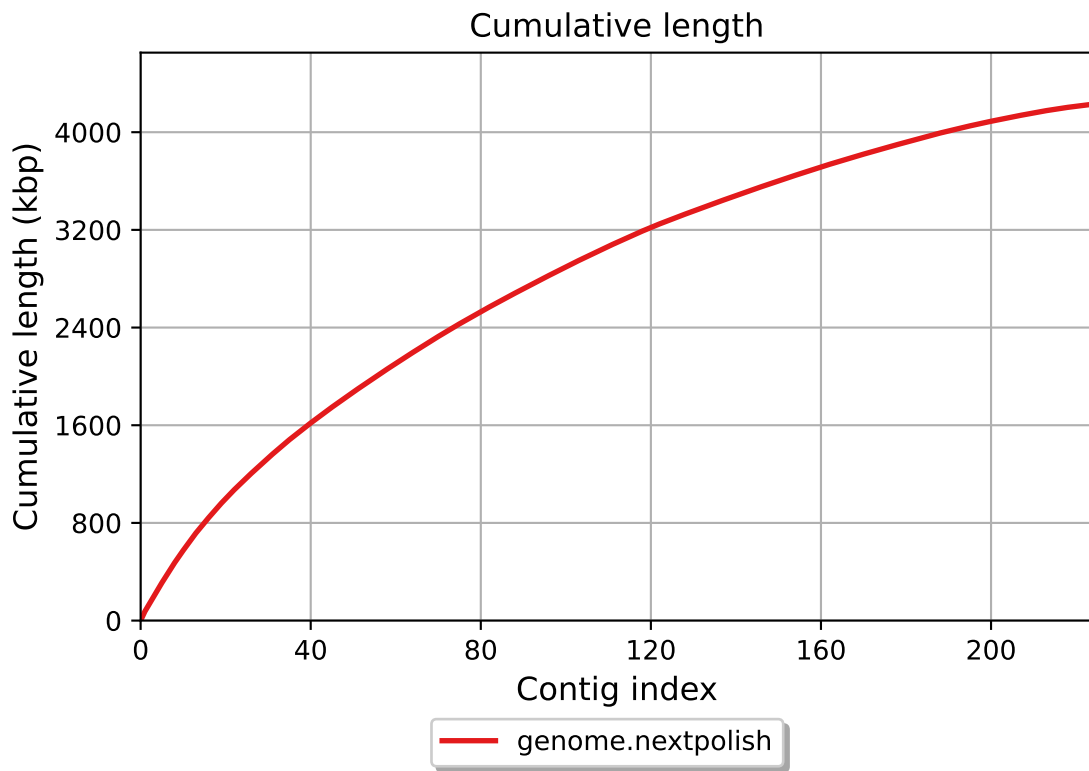
	genome.nextpolish
# contigs (>= 0 bp)	224
# contigs (>= 1000 bp)	224
# contigs (>= 5000 bp)	217
# contigs (>= 10000 bp)	172
# contigs (>= 25000 bp)	46
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	4228773
Total length (>= 1000 bp)	4228773
Total length (>= 5000 bp)	4198358
Total length (>= 10000 bp)	3840285
Total length (>= 25000 bp)	1772927
Total length (>= 50000 bp)	526003
# contigs	224
Largest contig	74716
Total length	4228773
GC (%)	56.60
N50	22555
N90	10249
auN	27043.4
L50	61
L90	169
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

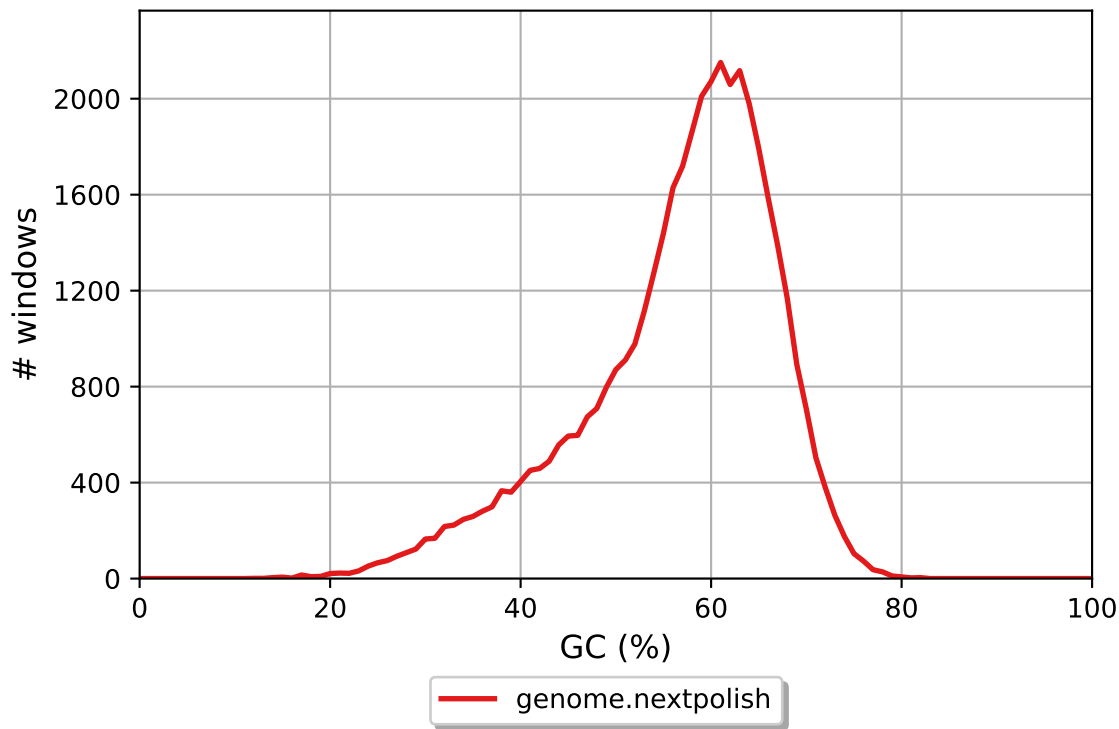
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

