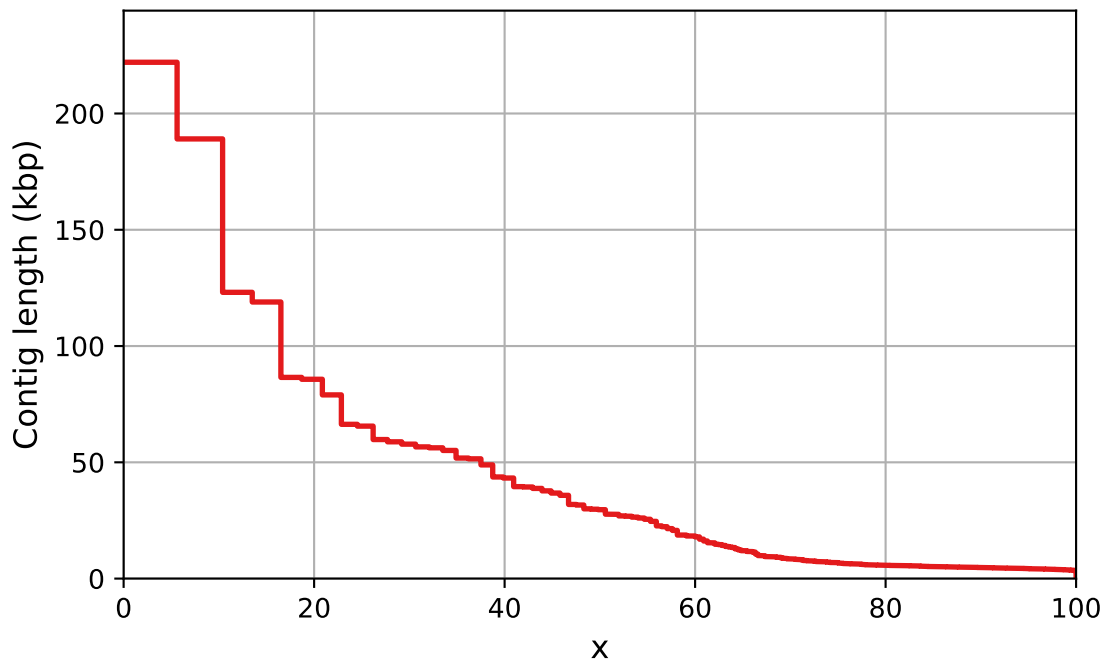


Report

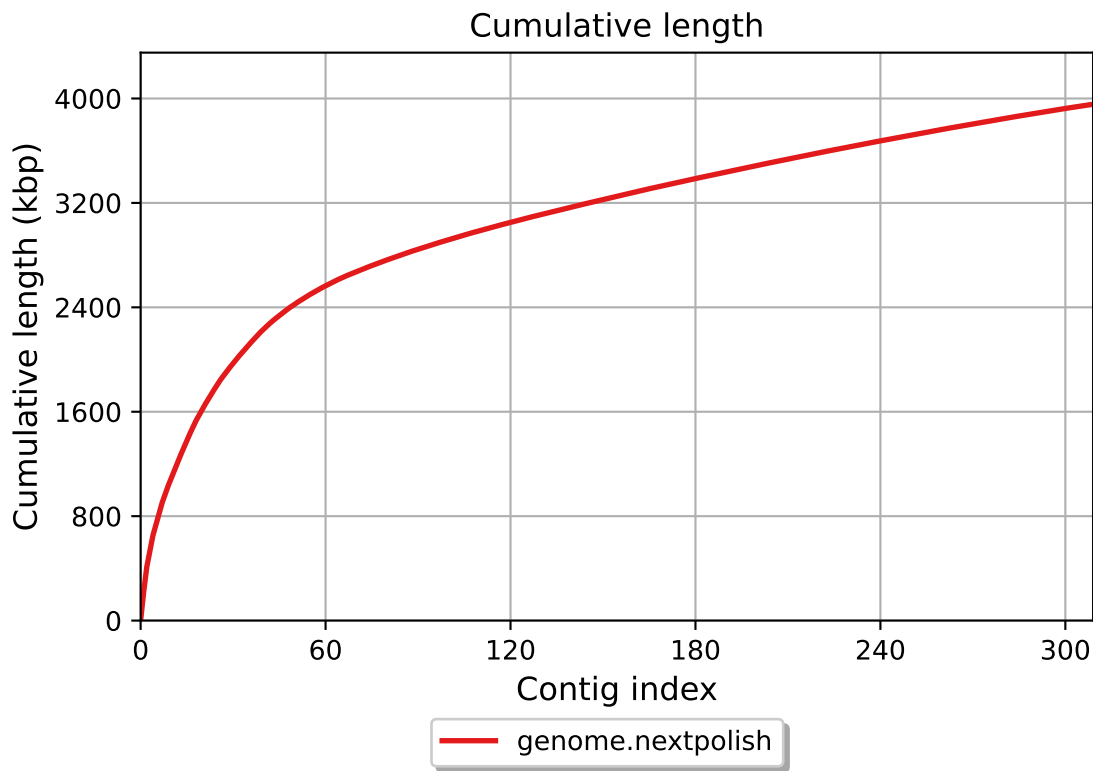
	genome.nextpolish
# contigs (>= 0 bp)	309
# contigs (>= 1000 bp)	309
# contigs (>= 5000 bp)	193
# contigs (>= 10000 bp)	66
# contigs (>= 25000 bp)	38
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	3955737
Total length (>= 1000 bp)	3955737
Total length (>= 5000 bp)	3452250
Total length (>= 10000 bp)	2634185
Total length (>= 25000 bp)	2187842
Total length (>= 50000 bp)	1483981
# contigs	309
Largest contig	222028
Total length	3955737
GC (%)	60.04
N50	29630
N90	4757
auN	52833.6
L50	31
L90	216
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

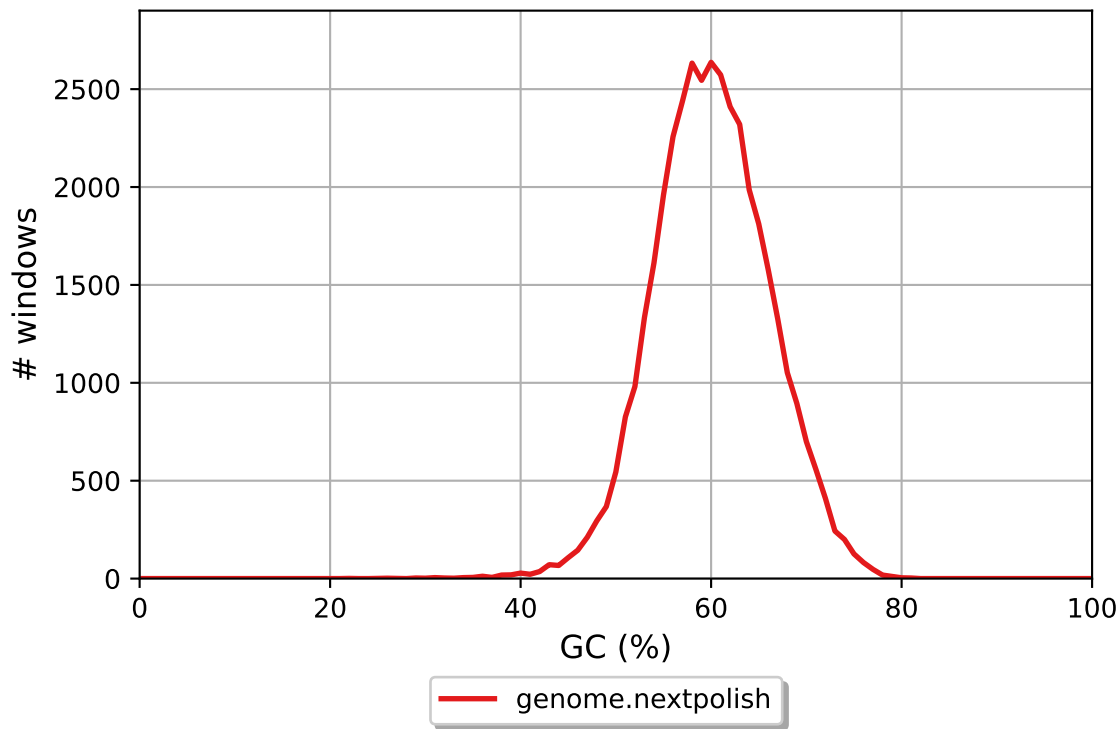
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

