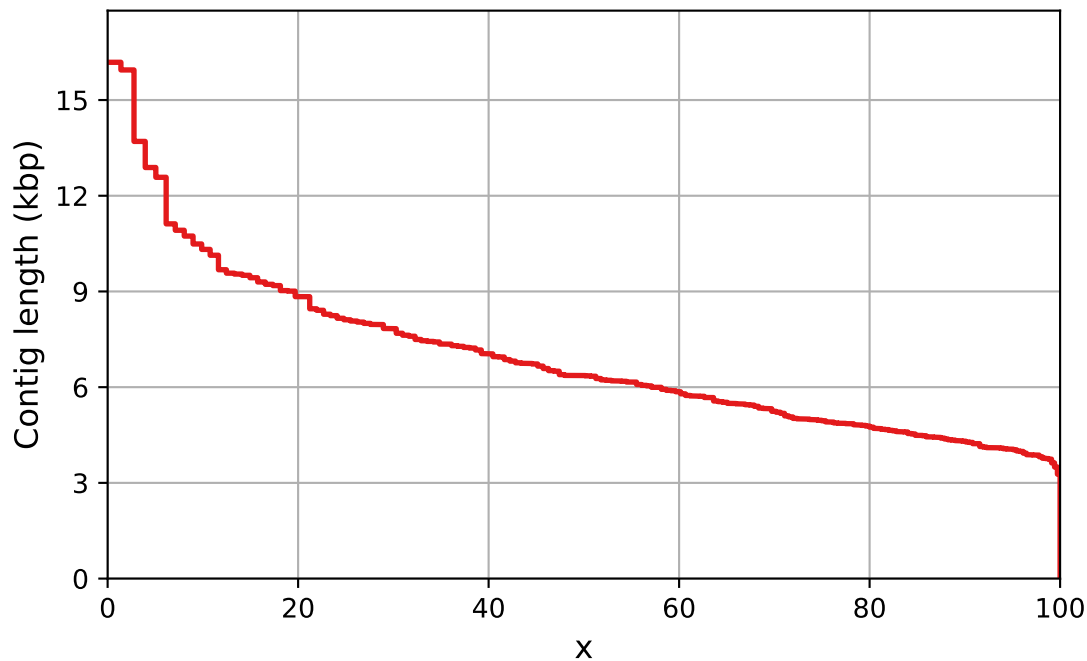


Report

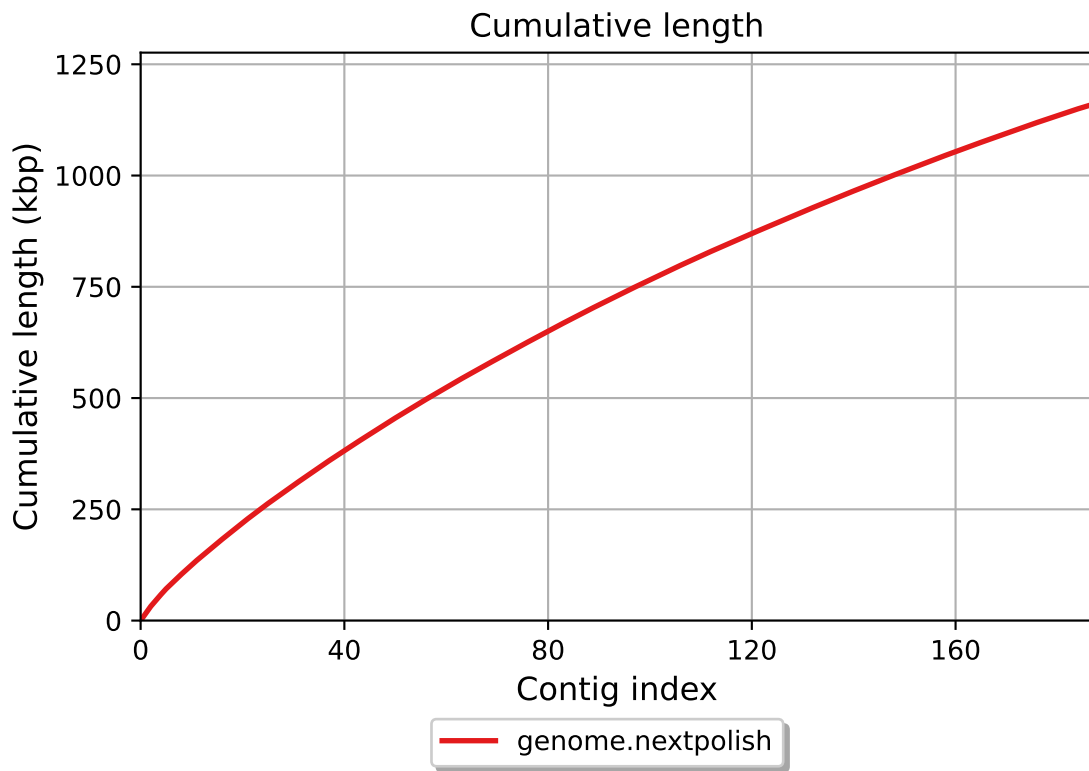
	genome.nextpolish
# contigs (>= 0 bp)	187
# contigs (>= 1000 bp)	187
# contigs (>= 5000 bp)	116
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1160138
Total length (>= 1000 bp)	1160138
Total length (>= 5000 bp)	849727
Total length (>= 10000 bp)	135006
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	187
Largest contig	16183
Total length	1160138
GC (%)	58.40
N50	6364
N90	4306
auN	6987.8
L50	69
L90	158
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

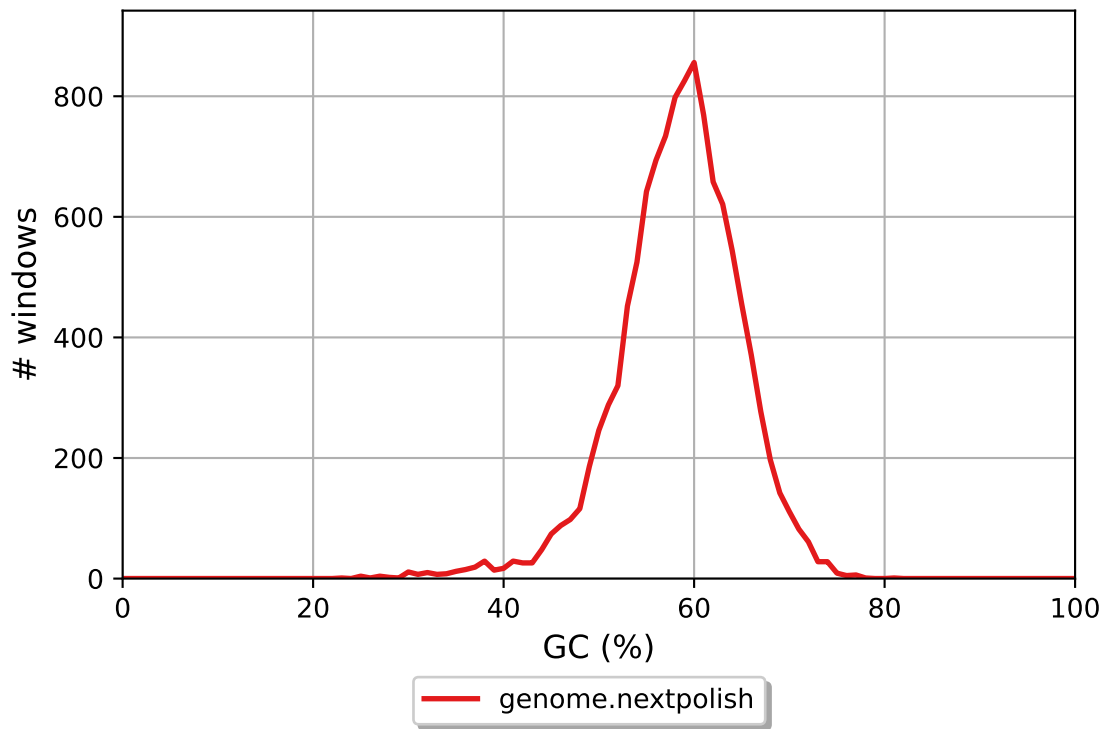
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

