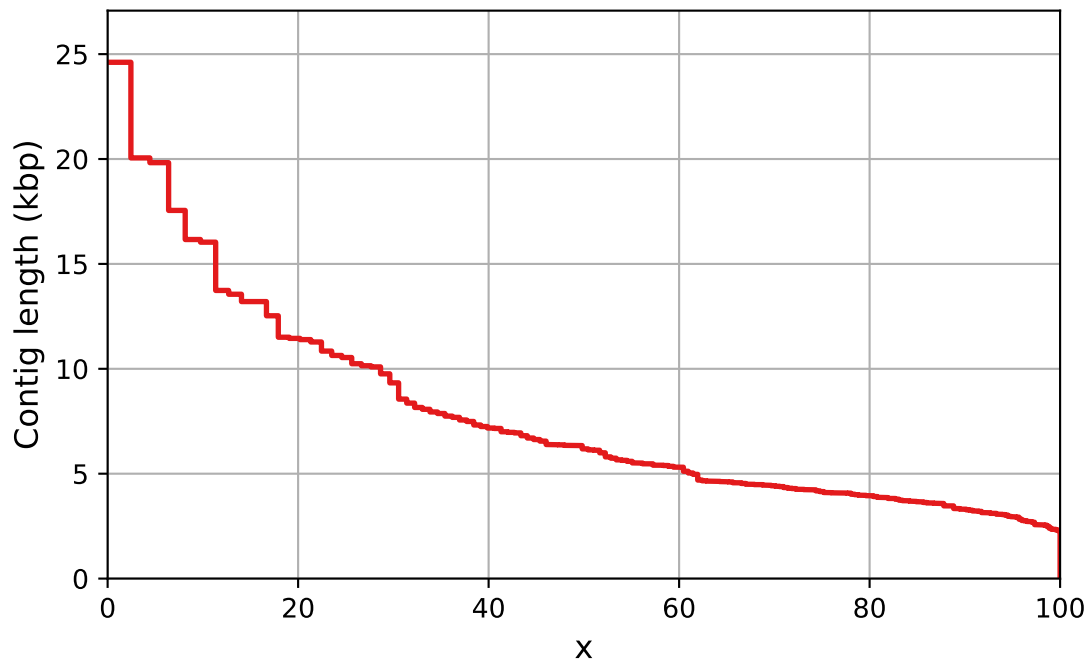


# Report

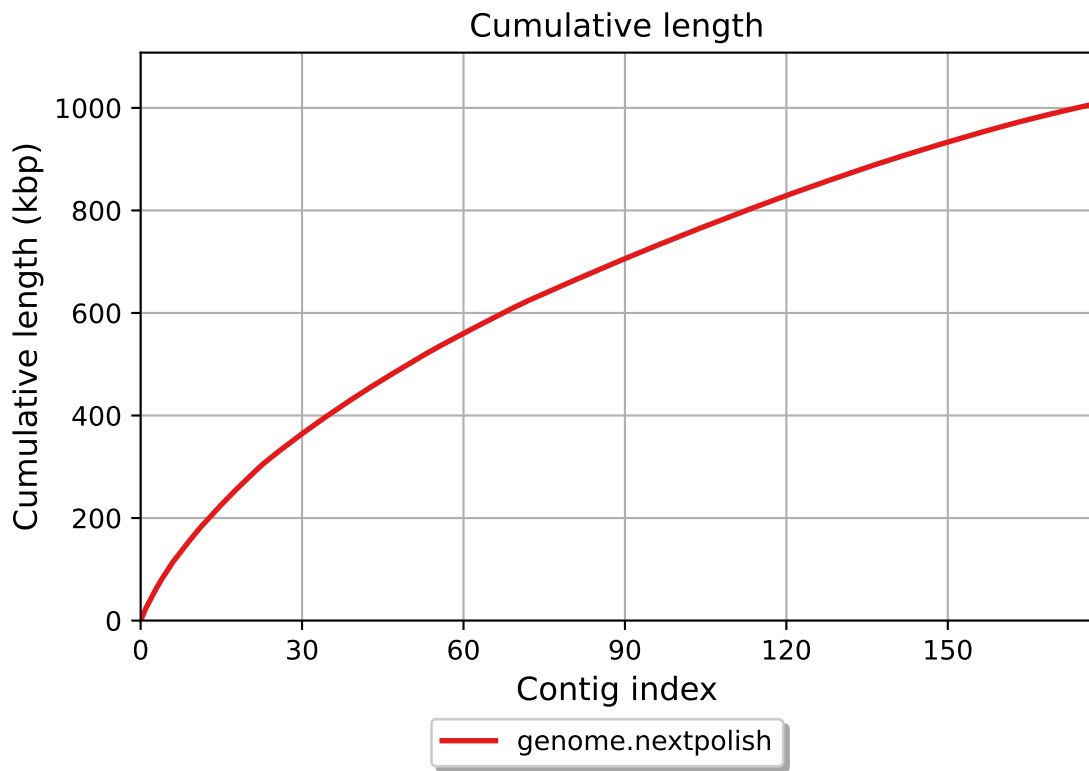
	genome.nextpolish
# contigs (>= 0 bp)	177
# contigs (>= 1000 bp)	177
# contigs (>= 5000 bp)	71
# contigs (>= 10000 bp)	21
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1007080
Total length (>= 1000 bp)	1007080
Total length (>= 5000 bp)	618942
Total length (>= 10000 bp)	288588
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	177
Largest contig	24611
Total length	1007080
GC (%)	61.45
N50	6184
N90	3308
auN	7938.6
L50	51
L90	142
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

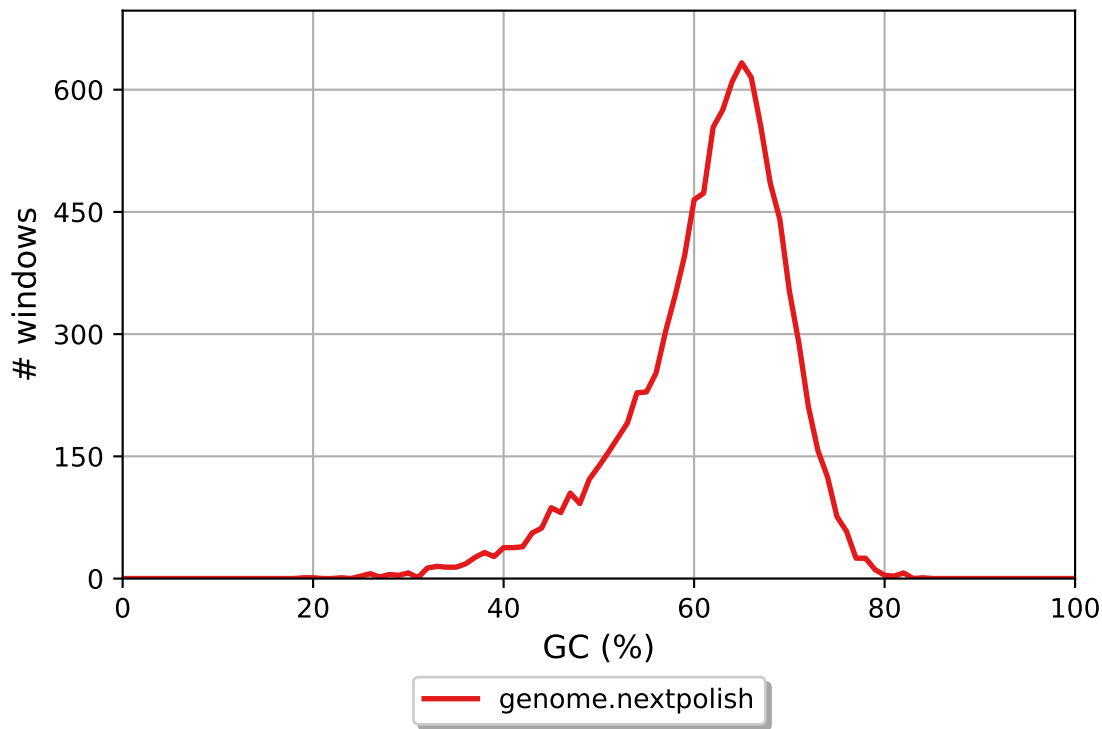
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

