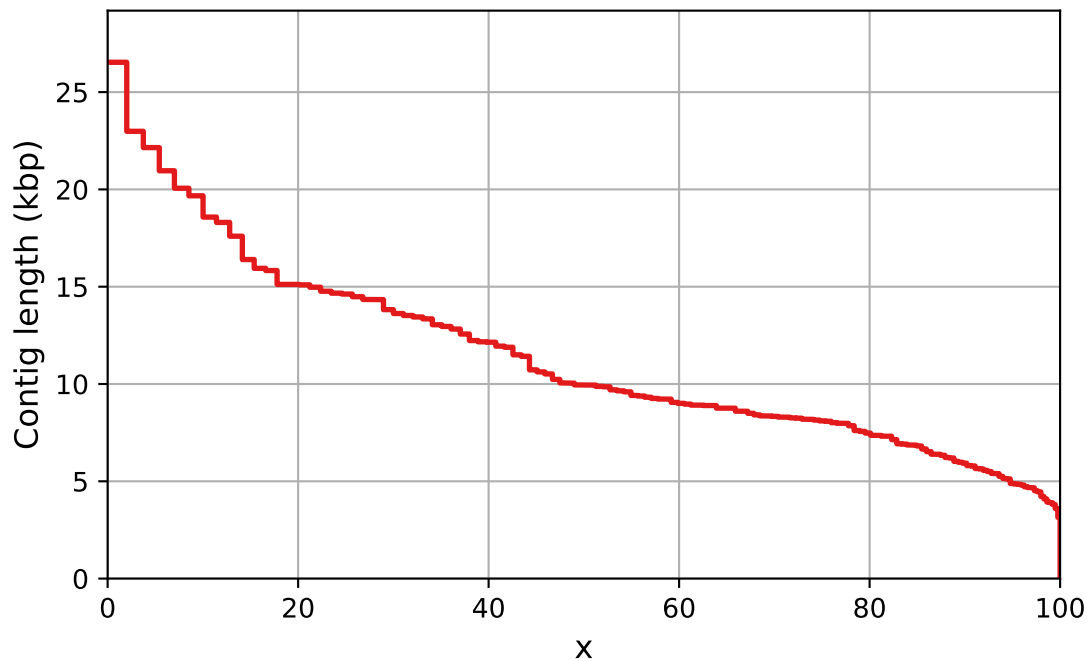


# Report

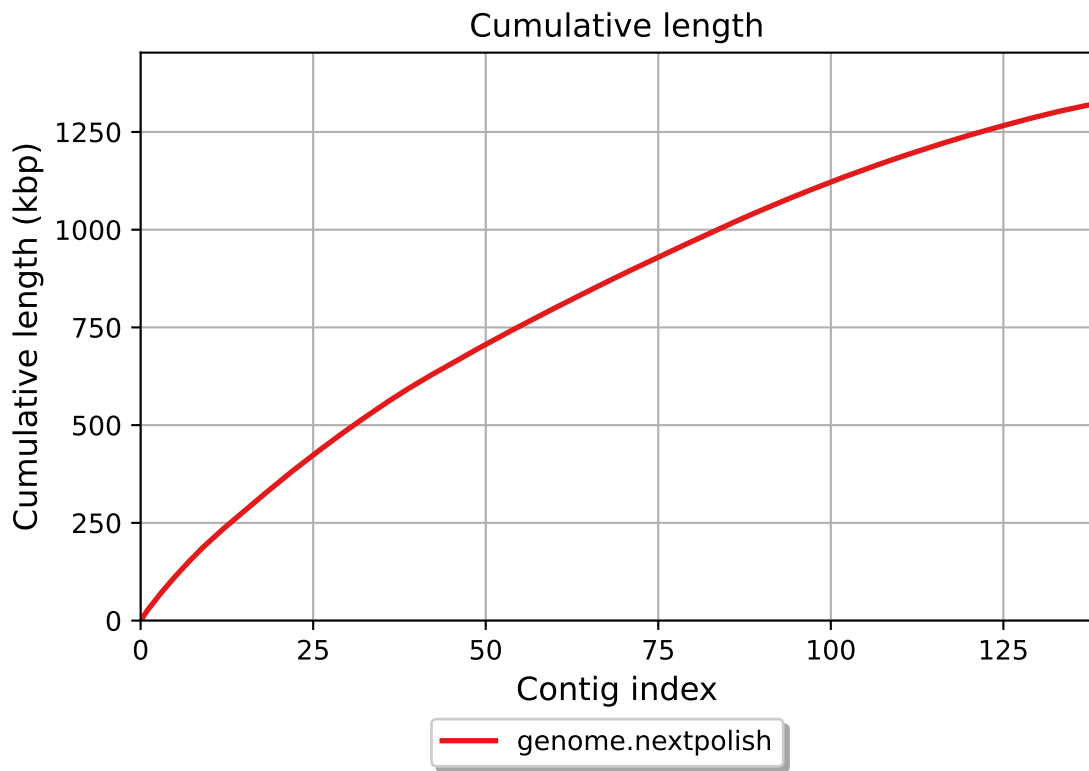
	genome.nextpolish
# contigs (>= 0 bp)	138
# contigs (>= 1000 bp)	138
# contigs (>= 5000 bp)	122
# contigs (>= 10000 bp)	44
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1320829
Total length (>= 1000 bp)	1320829
Total length (>= 5000 bp)	1251680
Total length (>= 10000 bp)	647188
Total length (>= 25000 bp)	26533
Total length (>= 50000 bp)	0
# contigs	138
Largest contig	26533
Total length	1320829
GC (%)	54.21
N50	9954
N90	5928
auN	11544.6
L50	46
L90	111
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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).

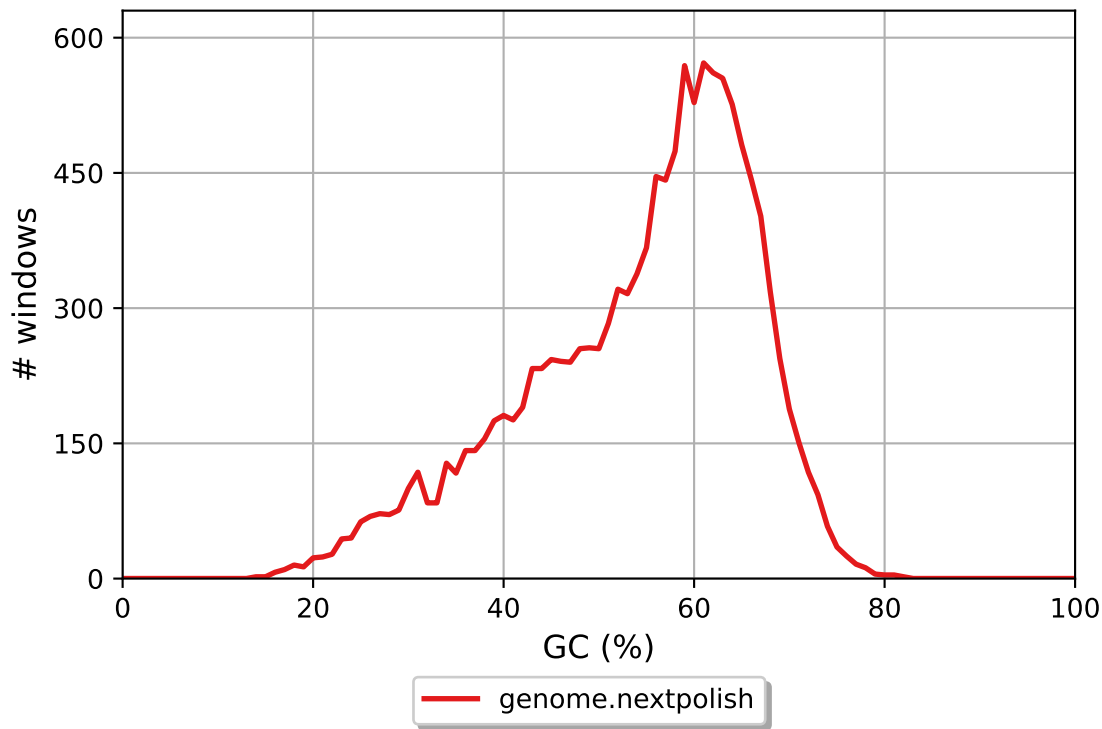
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

