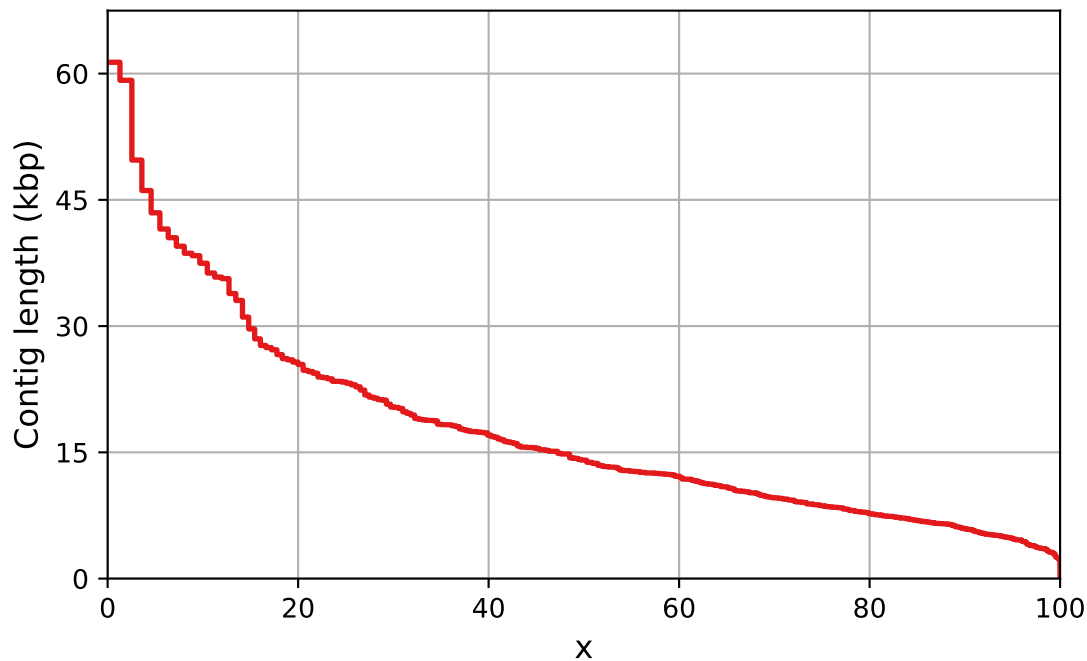


# Report

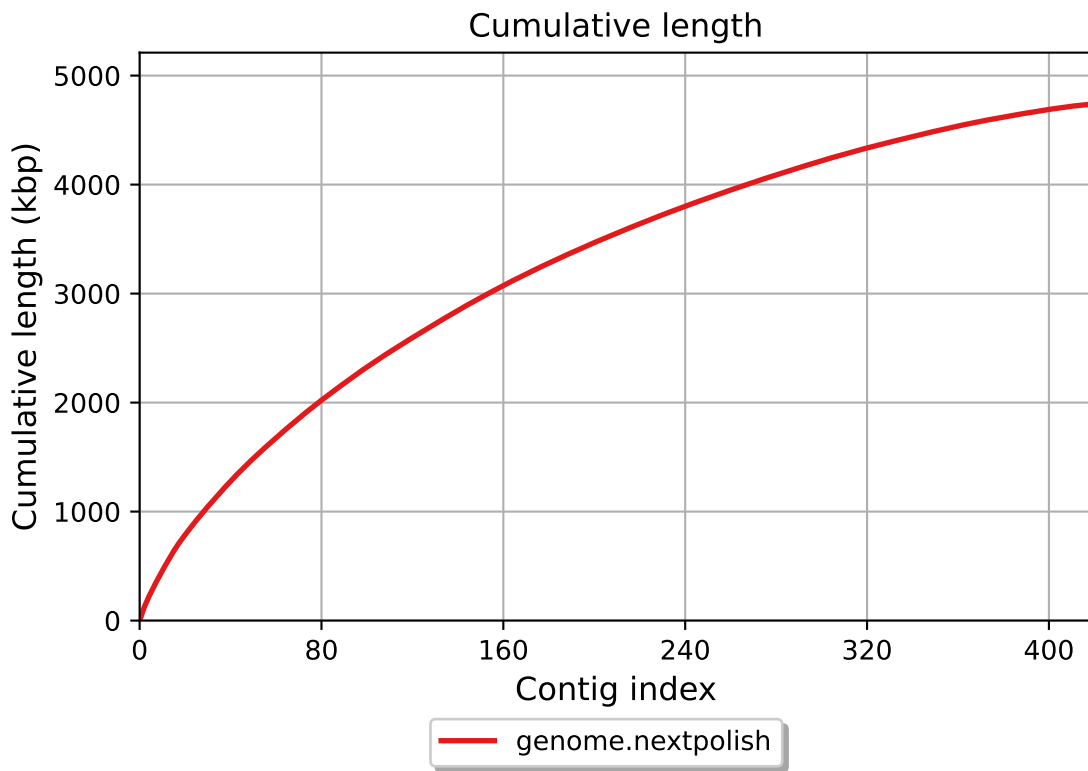
	genome.nextpolish
# contigs (>= 0 bp)	419
# contigs (>= 1000 bp)	417
# contigs (>= 5000 bp)	342
# contigs (>= 10000 bp)	176
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	4736912
Total length (>= 1000 bp)	4735693
Total length (>= 5000 bp)	4450865
Total length (>= 10000 bp)	3240184
Total length (>= 25000 bp)	971981
Total length (>= 50000 bp)	120549
# contigs	419
Largest contig	61346
Total length	4736912
GC (%)	55.92
N50	14088
N90	5915
auN	17862.2
L50	103
L90	308
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

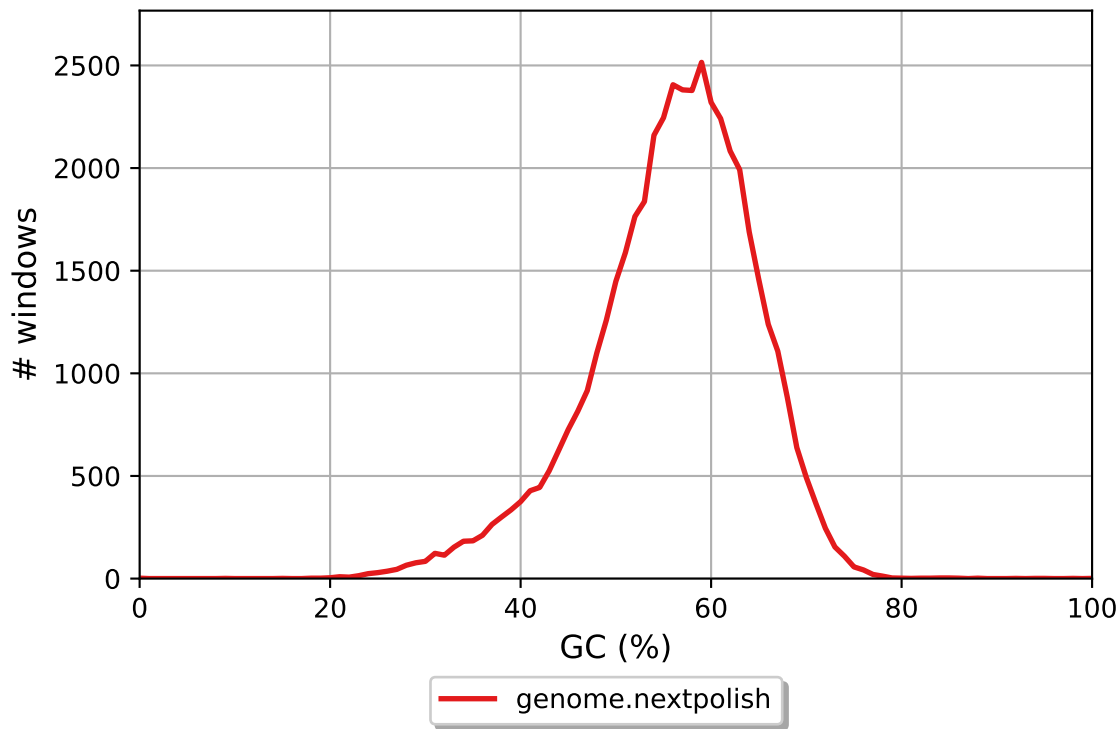
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

