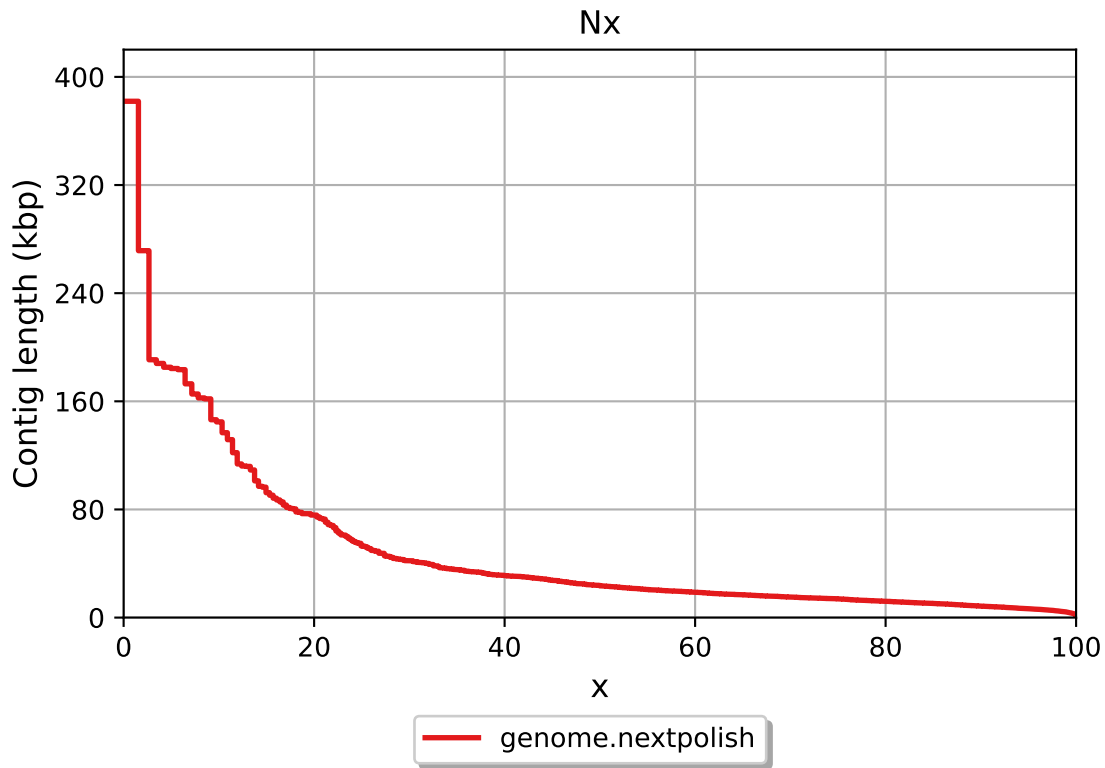


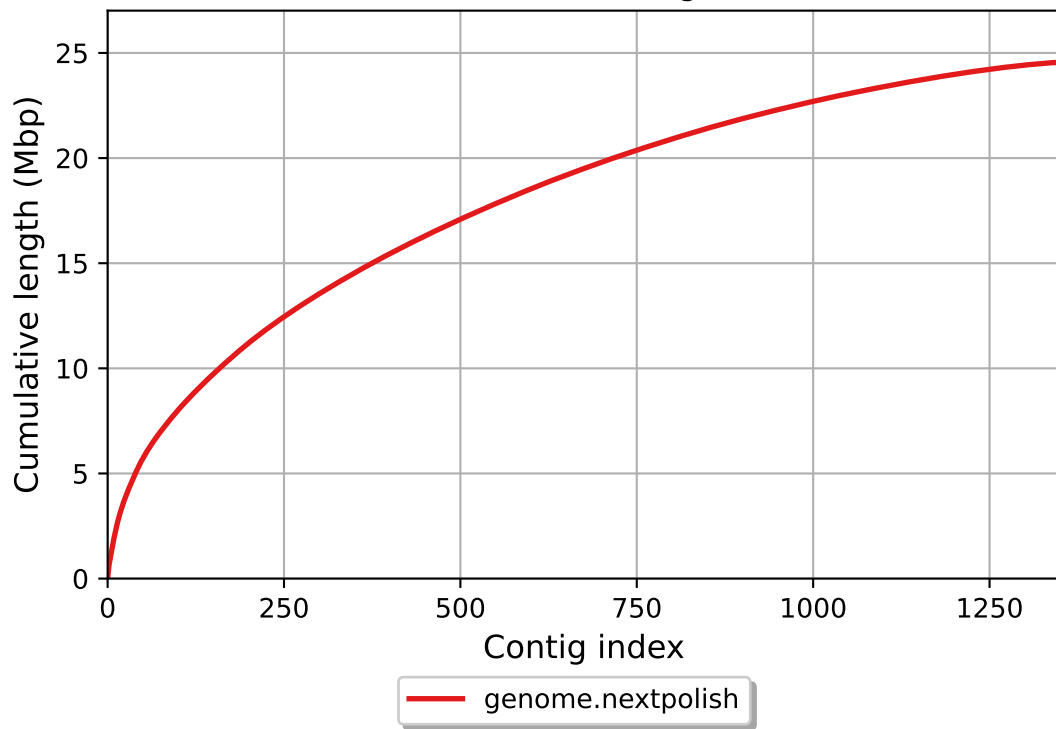
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1350
# contigs (>= 1000 bp)	1347
# contigs (>= 5000 bp)	1217
# contigs (>= 10000 bp)	831
# contigs (>= 25000 bp)	222
# contigs (>= 50000 bp)	62
Total length (>= 0 bp)	24557211
Total length (>= 1000 bp)	24554981
Total length (>= 5000 bp)	24069078
Total length (>= 10000 bp)	21227612
Total length (>= 25000 bp)	11780289
Total length (>= 50000 bp)	6381232
# contigs	1350
Largest contig	381966
Total length	24557211
GC (%)	50.93
N50	23725
N90	8495
auN	50390.1
L50	243
L90	926
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	55 + 4 part

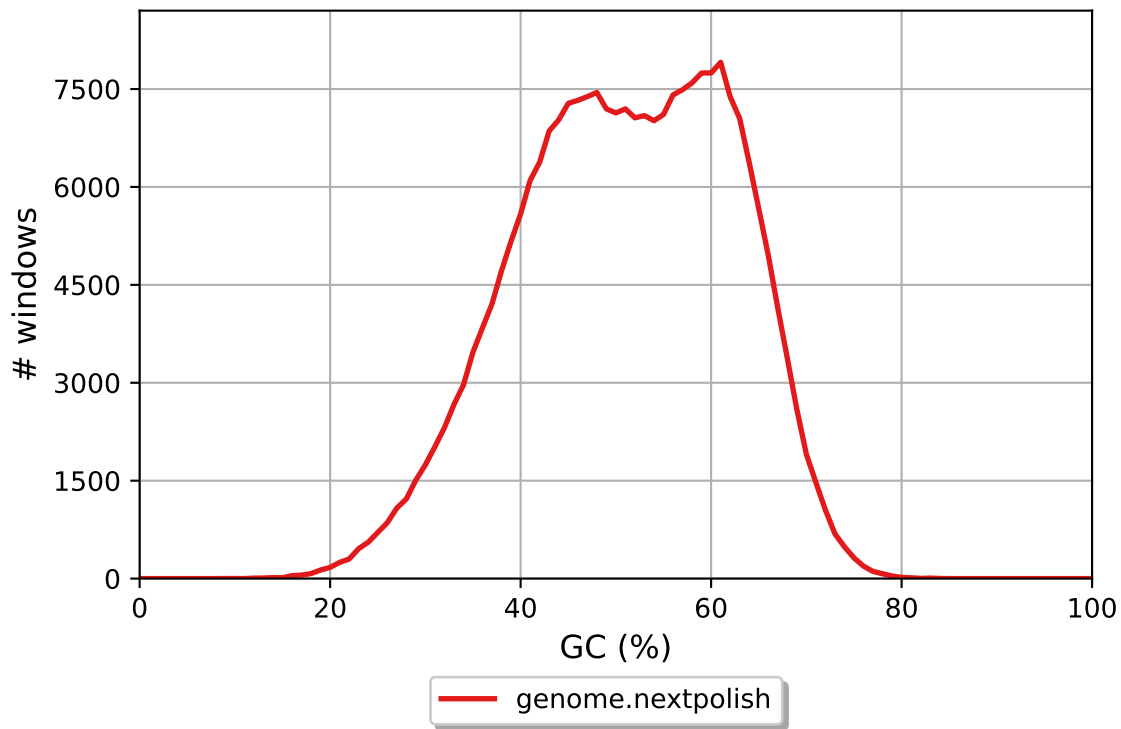
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

