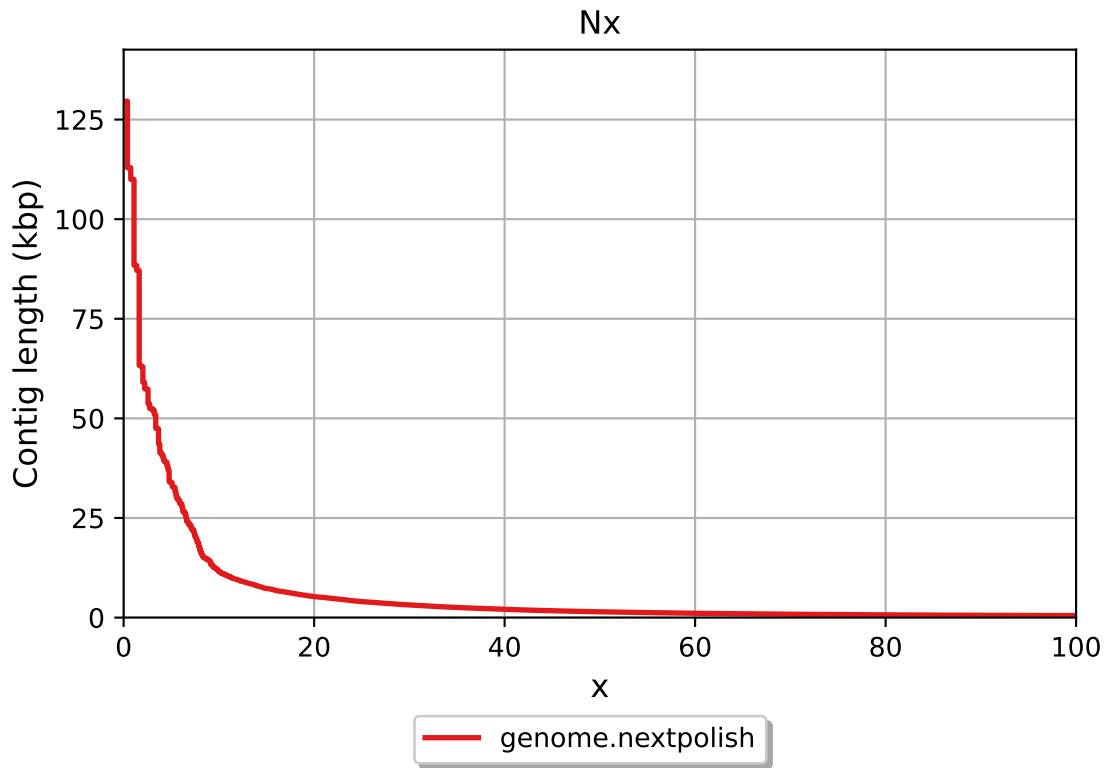


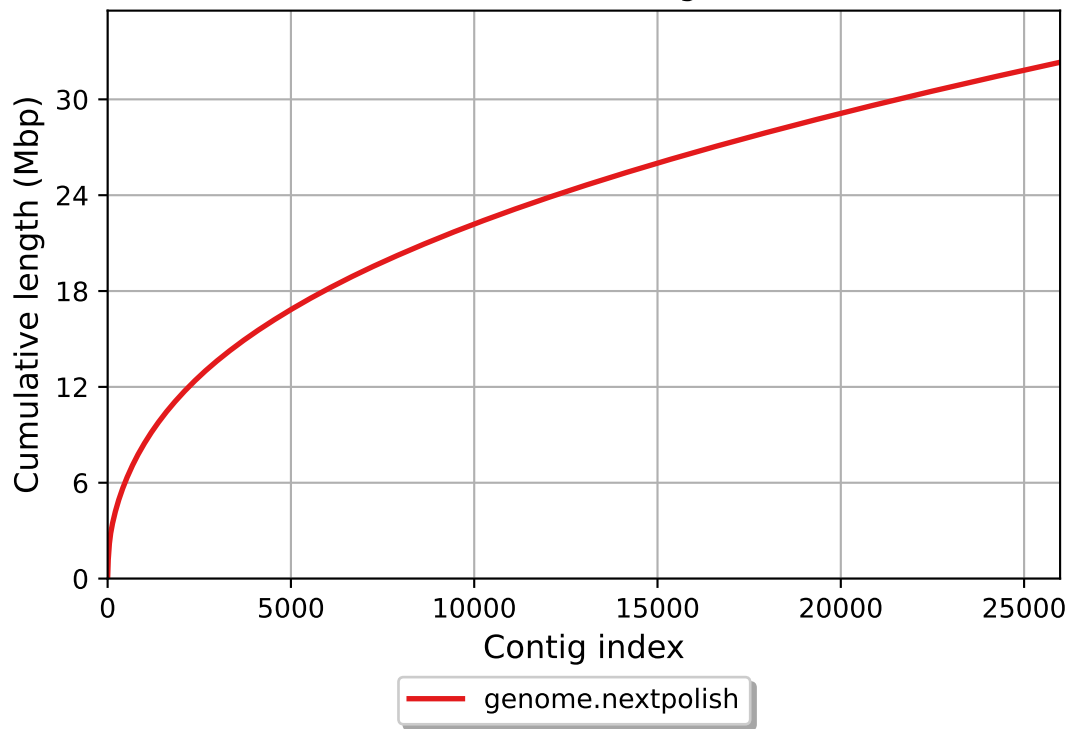
Report

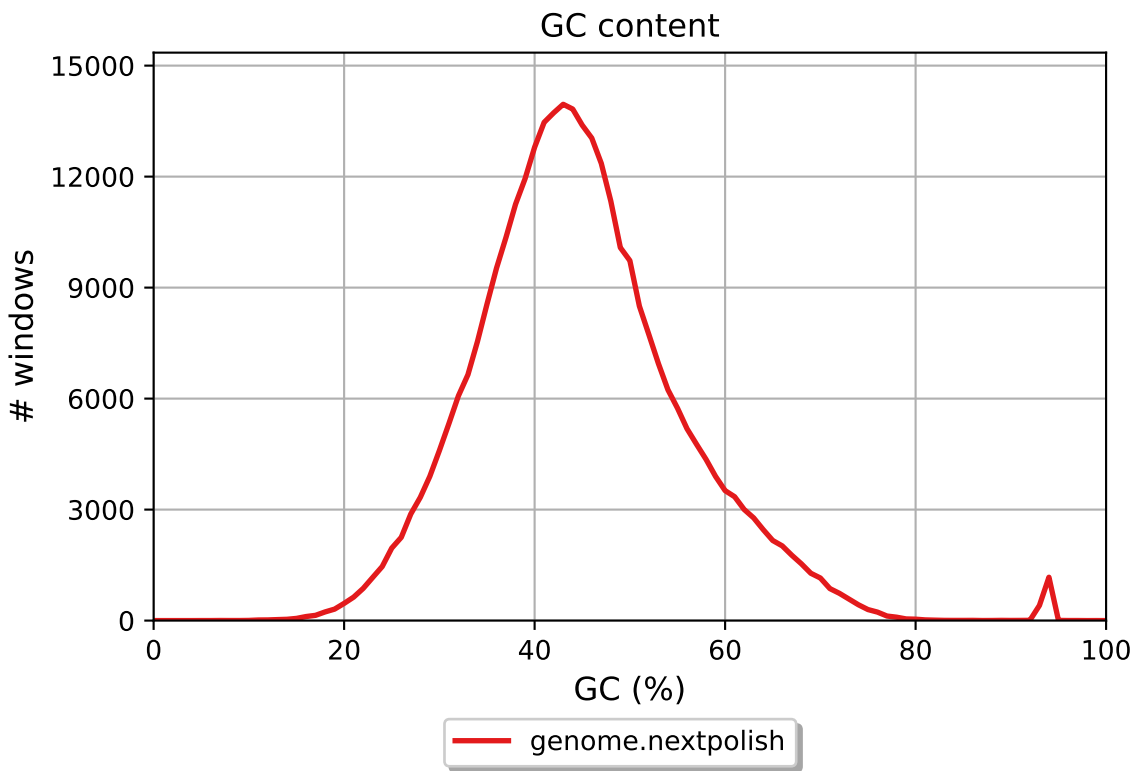
	genome.nextpolish
# contigs (>= 0 bp)	197082
# contigs (>= 1000 bp)	8029
# contigs (>= 5000 bp)	631
# contigs (>= 10000 bp)	154
# contigs (>= 25000 bp)	45
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	79762441
Total length (>= 1000 bp)	20352866
Total length (>= 5000 bp)	6855231
Total length (>= 10000 bp)	3665944
Total length (>= 25000 bp)	2113942
Total length (>= 50000 bp)	1089615
# contigs	25982
Largest contig	129591
Total length	32321200
GC (%)	44.83
N50	1454
N90	576
auN	6486.0
L50	4518
L90	19939
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	38 + 11 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).

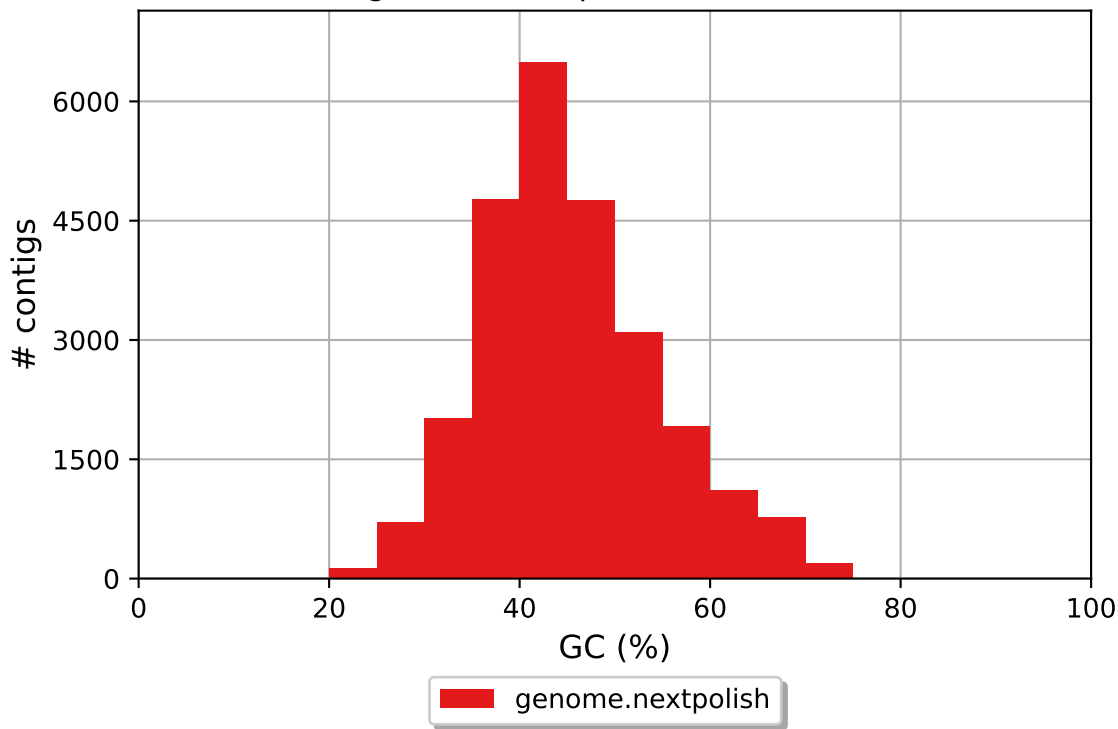


Cumulative length

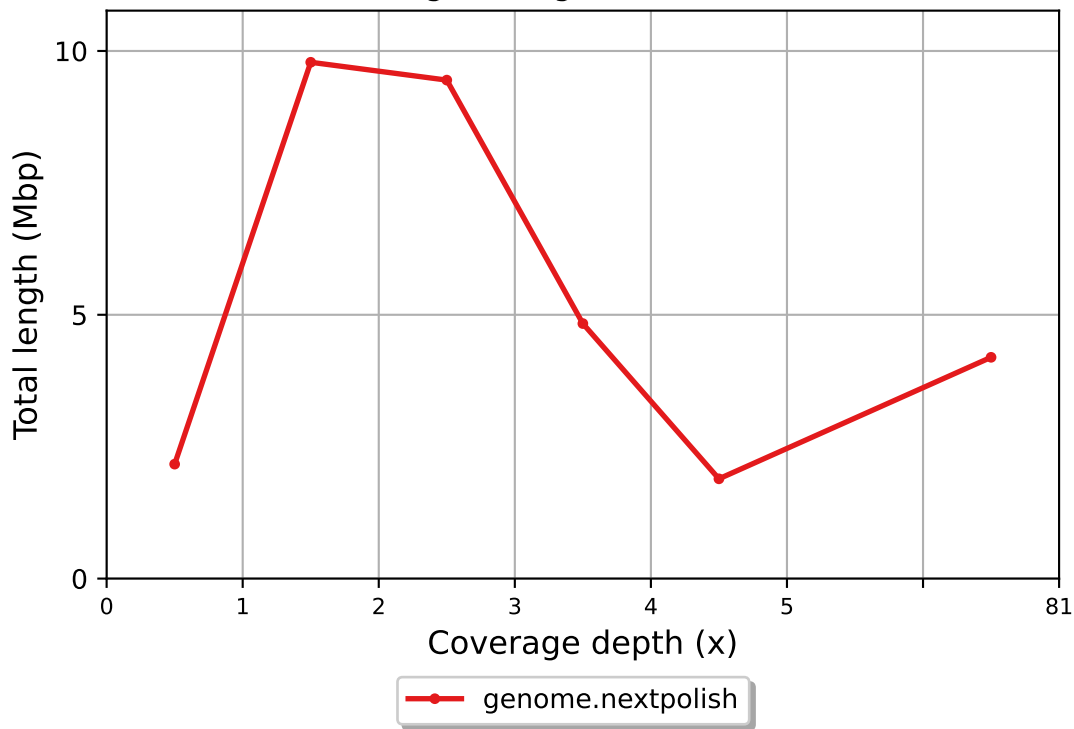




genome.nextpolish GC content



Coverage histogram (bin size: 1x)



genome.nextpolish coverage histogram (bin size: 1x)

