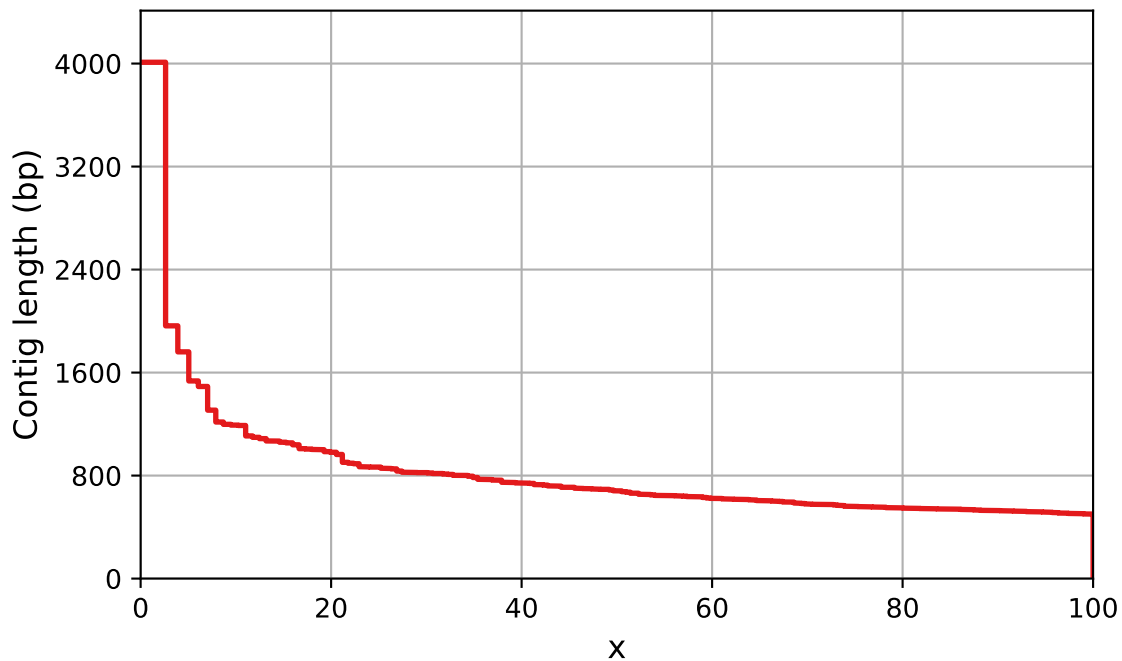


Report

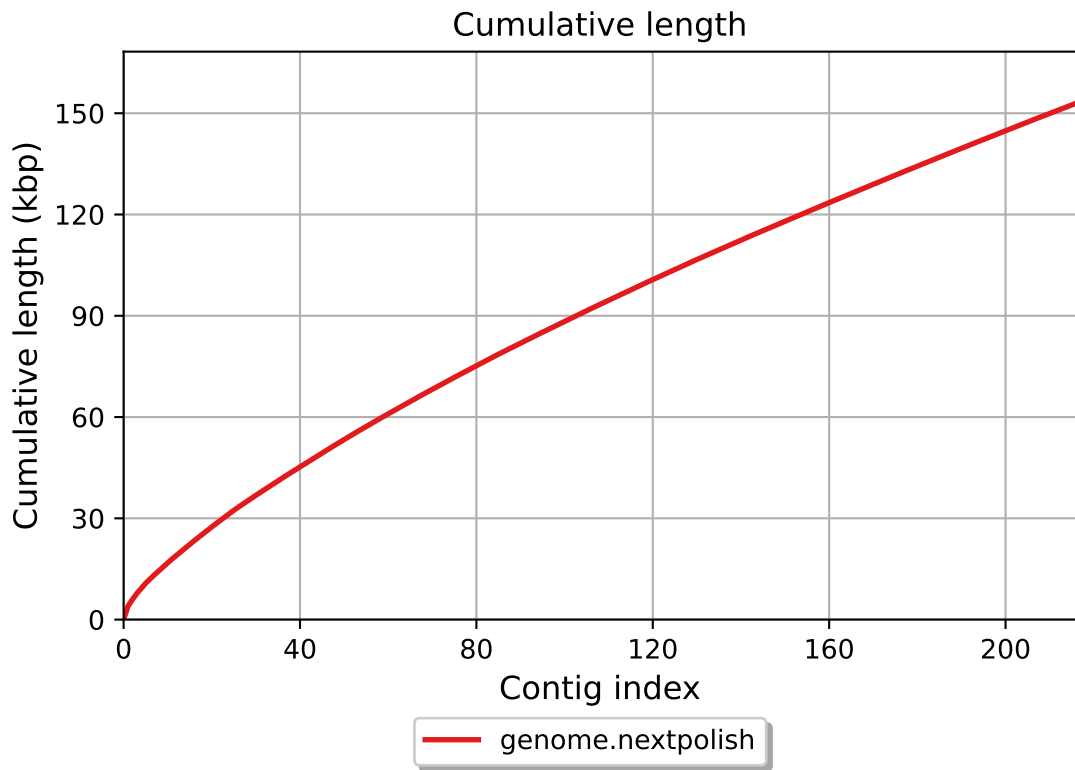
	genome.nextpolish
# contigs (>= 0 bp)	3891
# contigs (>= 1000 bp)	22
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	900406
Total length (>= 1000 bp)	29464
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	216
Largest contig	4010
Total length	152952
GC (%)	54.54
N50	682
N90	528
auN	846.7
L50	82
L90	187
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

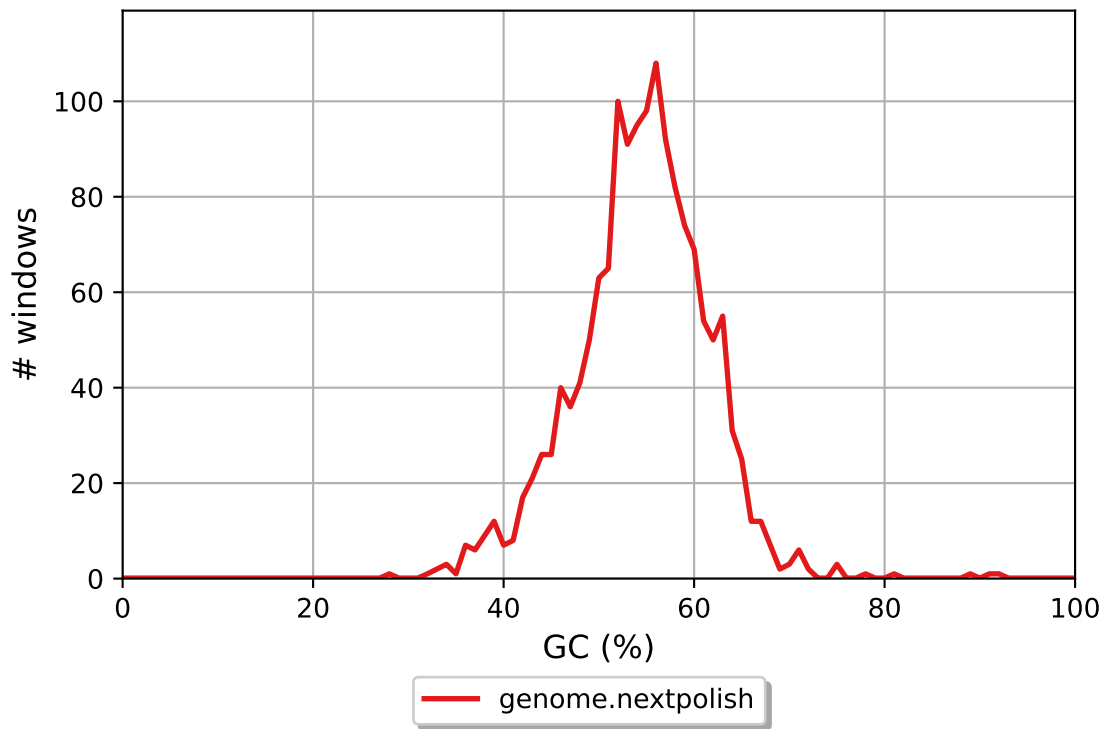
Nx



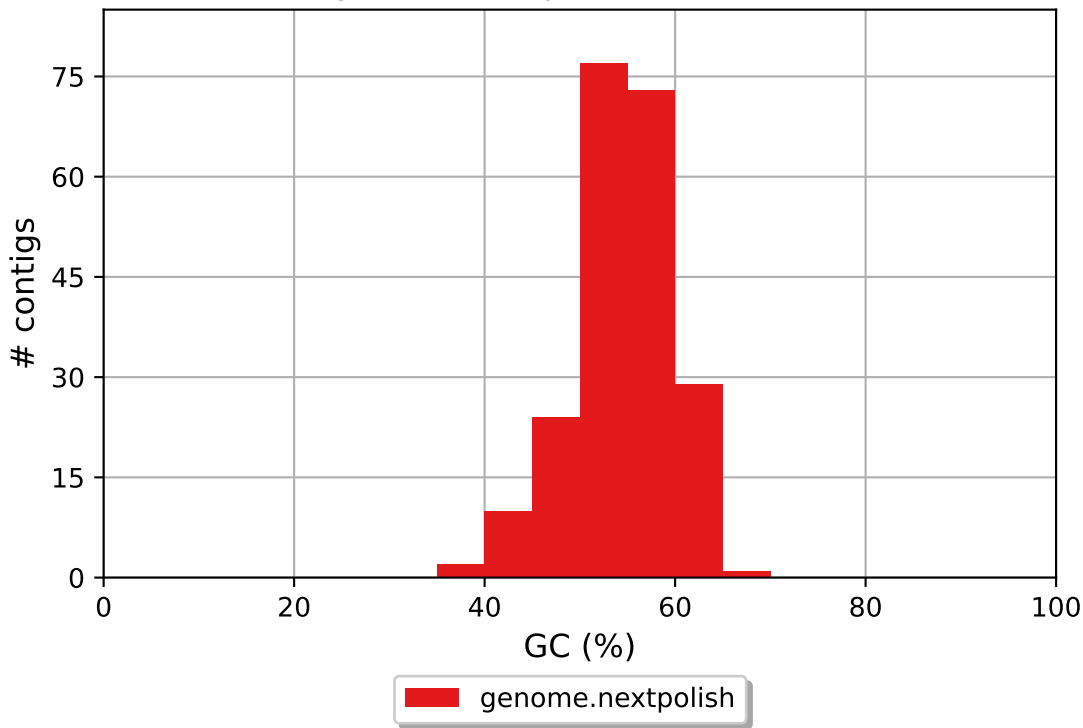
— genome.nextpolish



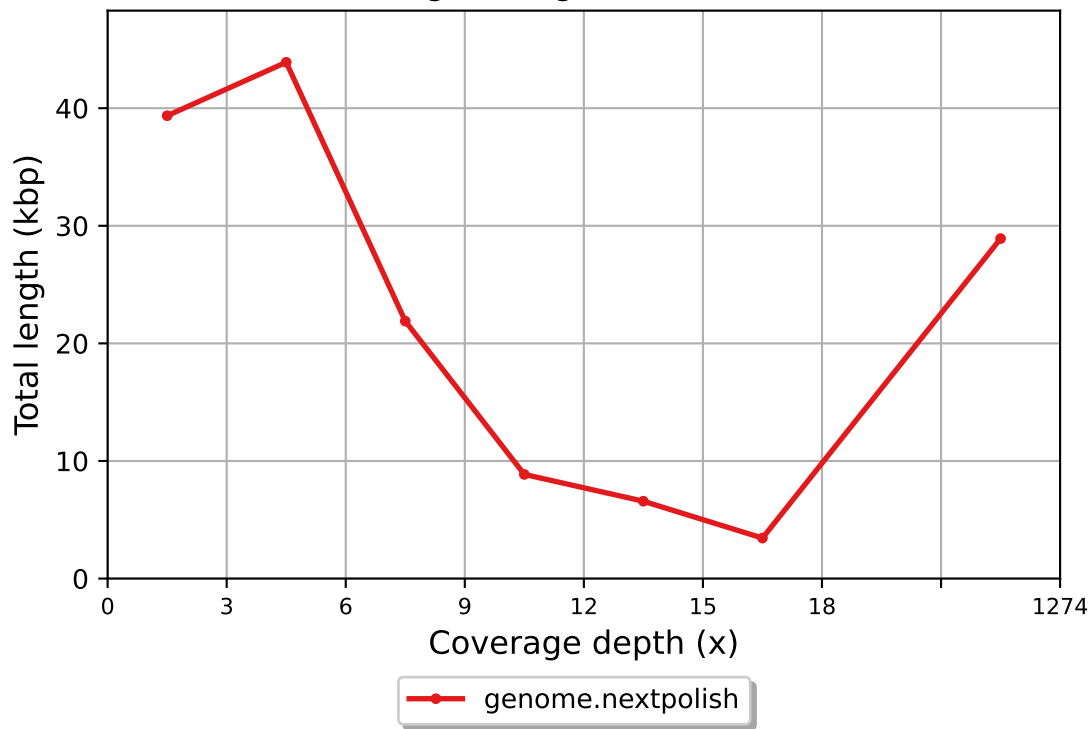
GC content



genome.nextpolish GC content



Coverage histogram (bin size: 3x)



genome.nextpolish coverage histogram (bin size: 3x)

