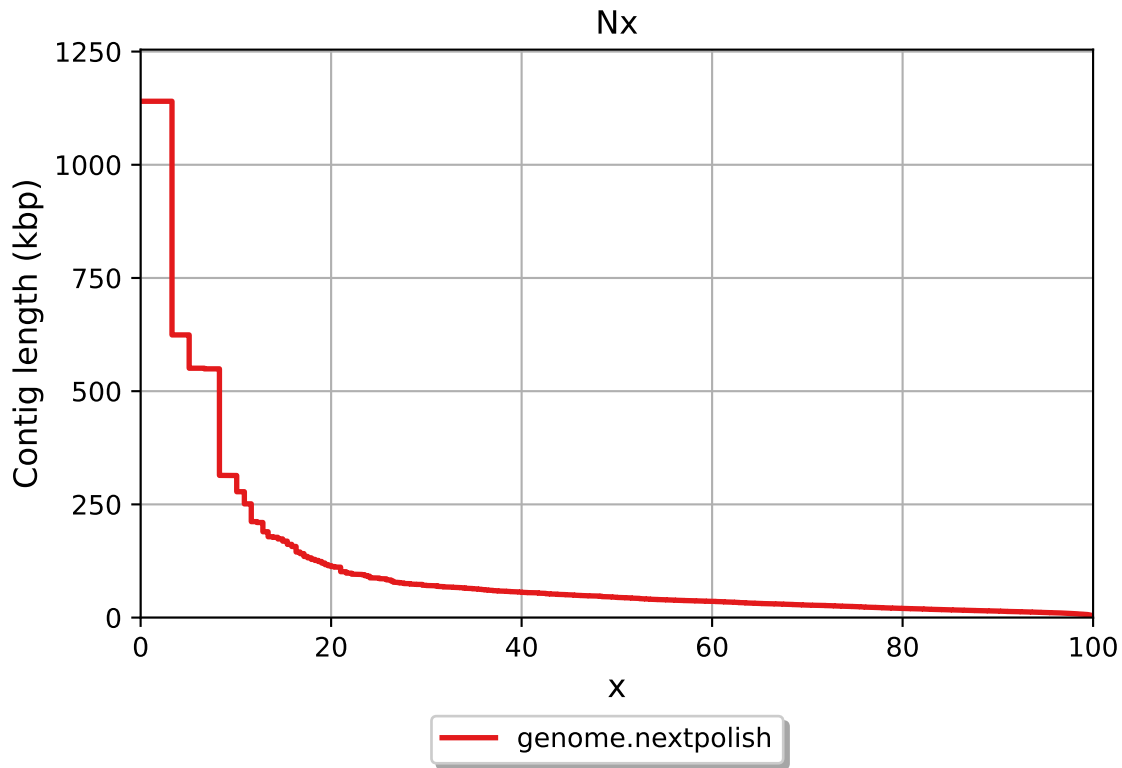


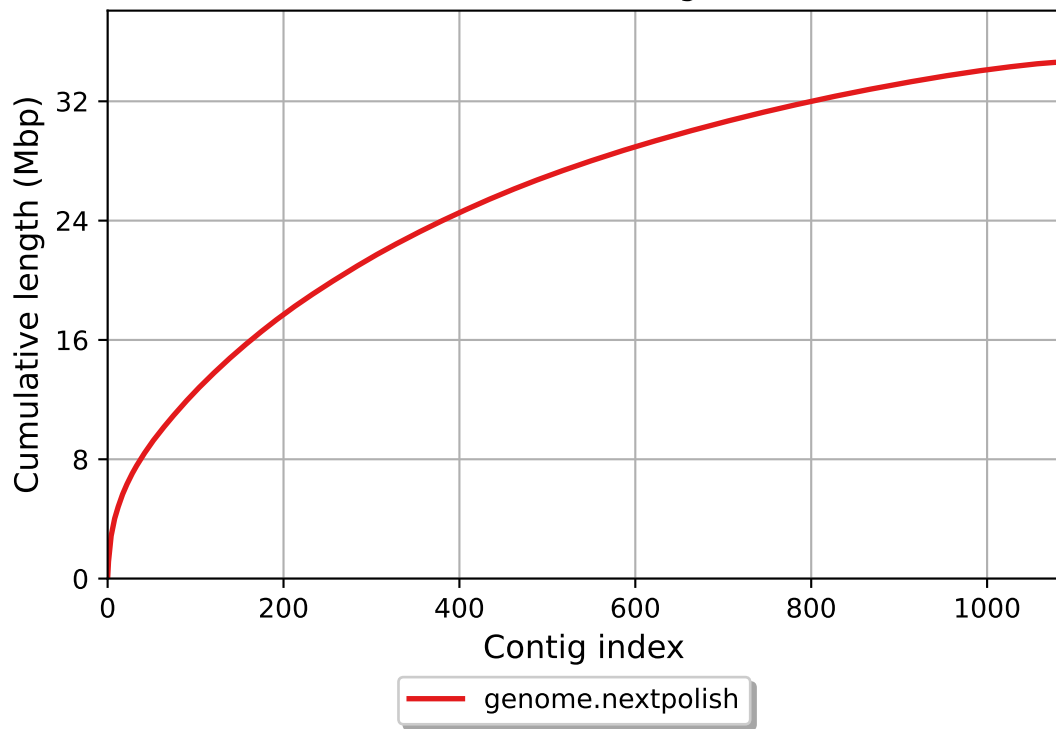
Report

	genome.nextpolish
# contigs (>= 0 bp)	1083
# contigs (>= 1000 bp)	1079
# contigs (>= 5000 bp)	1065
# contigs (>= 10000 bp)	933
# contigs (>= 25000 bp)	439
# contigs (>= 50000 bp)	155
Total length (>= 0 bp)	34617020
Total length (>= 1000 bp)	34614506
Total length (>= 5000 bp)	34558436
Total length (>= 10000 bp)	33512887
Total length (>= 25000 bp)	25556681
Total length (>= 50000 bp)	15599095
# contigs	1083
Largest contig	1140243
Total length	34617020
GC (%)	51.99
N50	44792
N90	14291
auN	121878.9
L50	191
L90	738
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	116 + 9 part

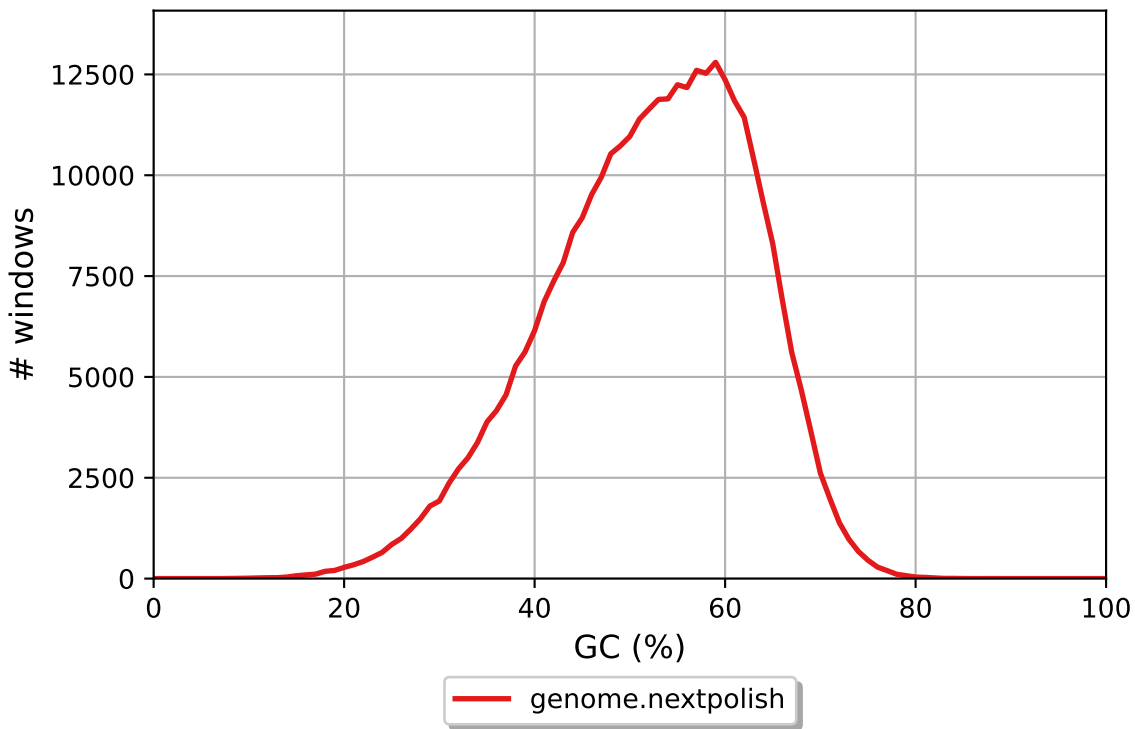
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

