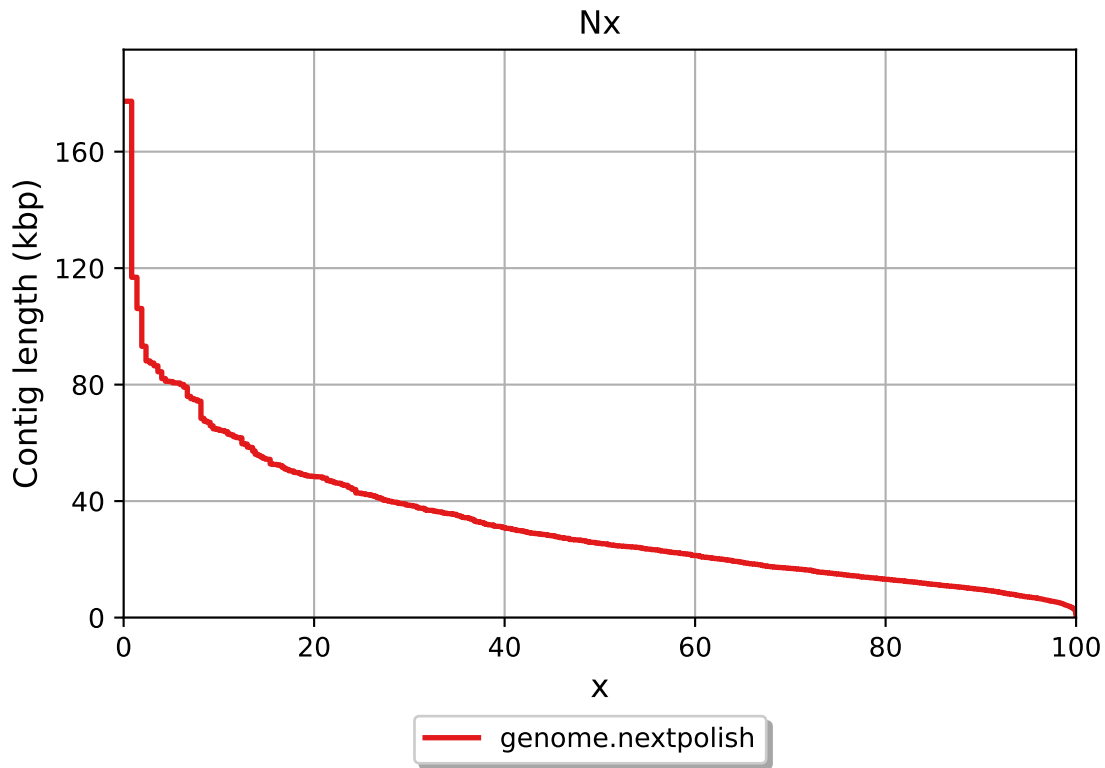


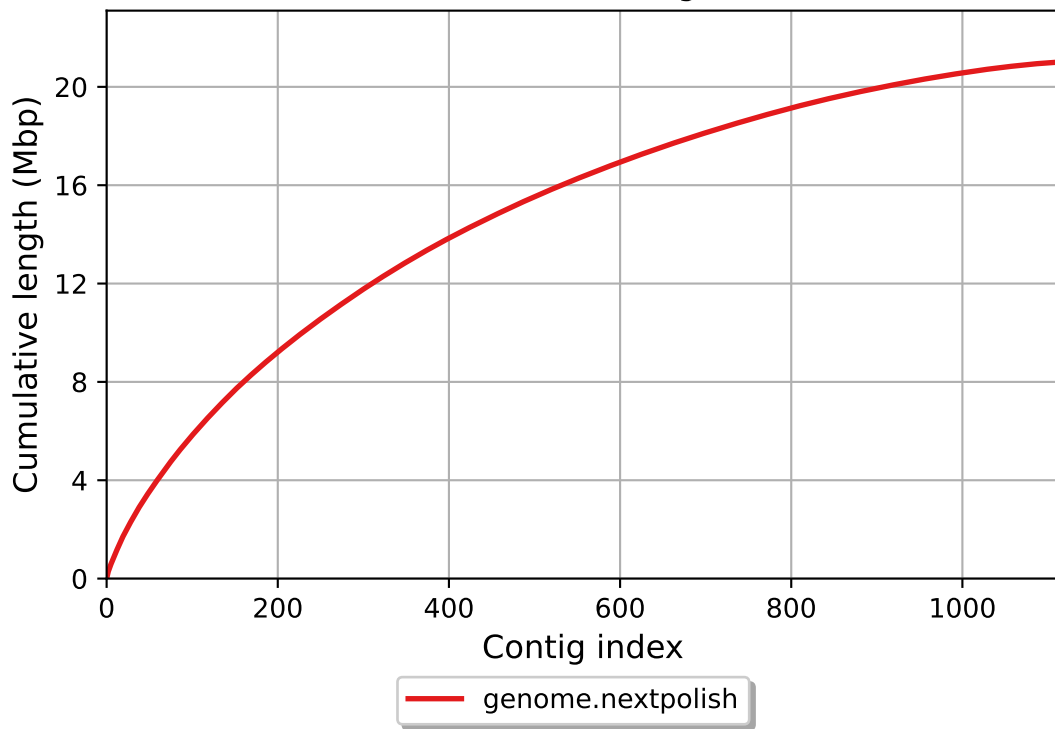
# Report

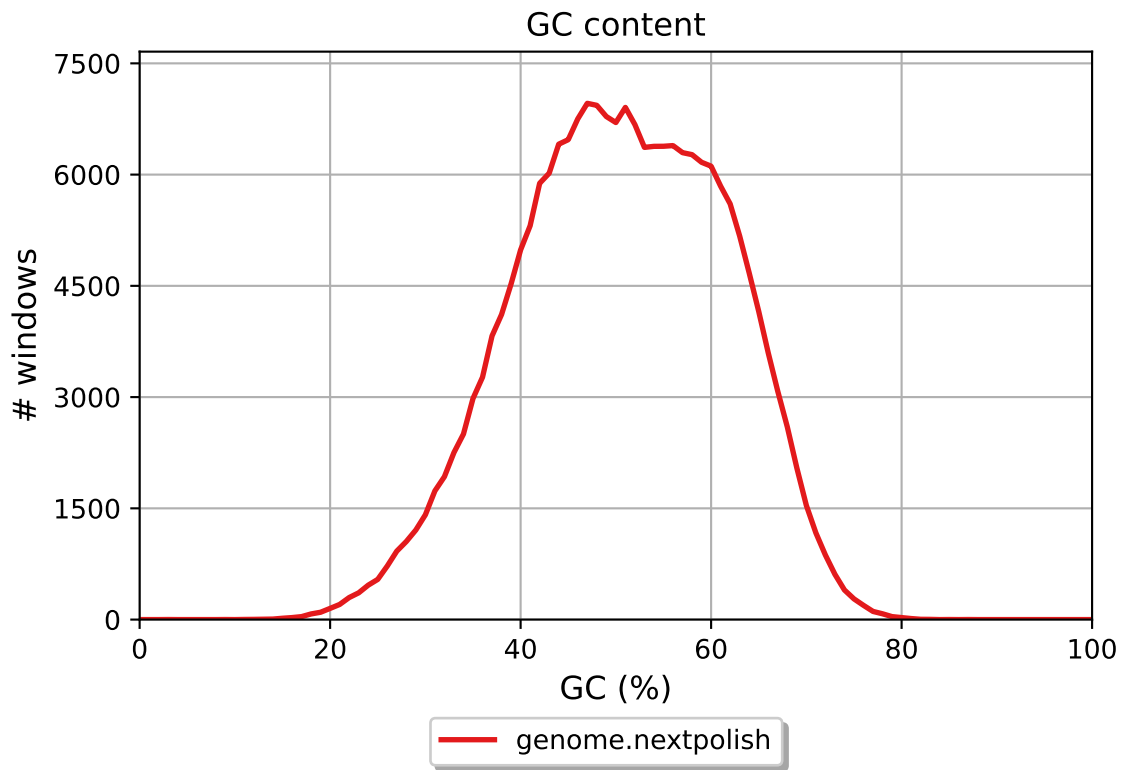
	genome.nextpolish
# contigs (>= 0 bp)	1114
# contigs (>= 1000 bp)	1109
# contigs (>= 5000 bp)	1017
# contigs (>= 10000 bp)	756
# contigs (>= 25000 bp)	256
# contigs (>= 50000 bp)	54
Total length (>= 0 bp)	21000531
Total length (>= 1000 bp)	20997121
Total length (>= 5000 bp)	20656904
Total length (>= 10000 bp)	18720033
Total length (>= 25000 bp)	10709475
Total length (>= 50000 bp)	3747547
# contigs	1113
Largest contig	177301
Total length	21000135
GC (%)	50.46
N50	25483
N90	9623
auN	32748.7
L50	248
L90	775
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	53 + 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





genome.nextpolish GC content

