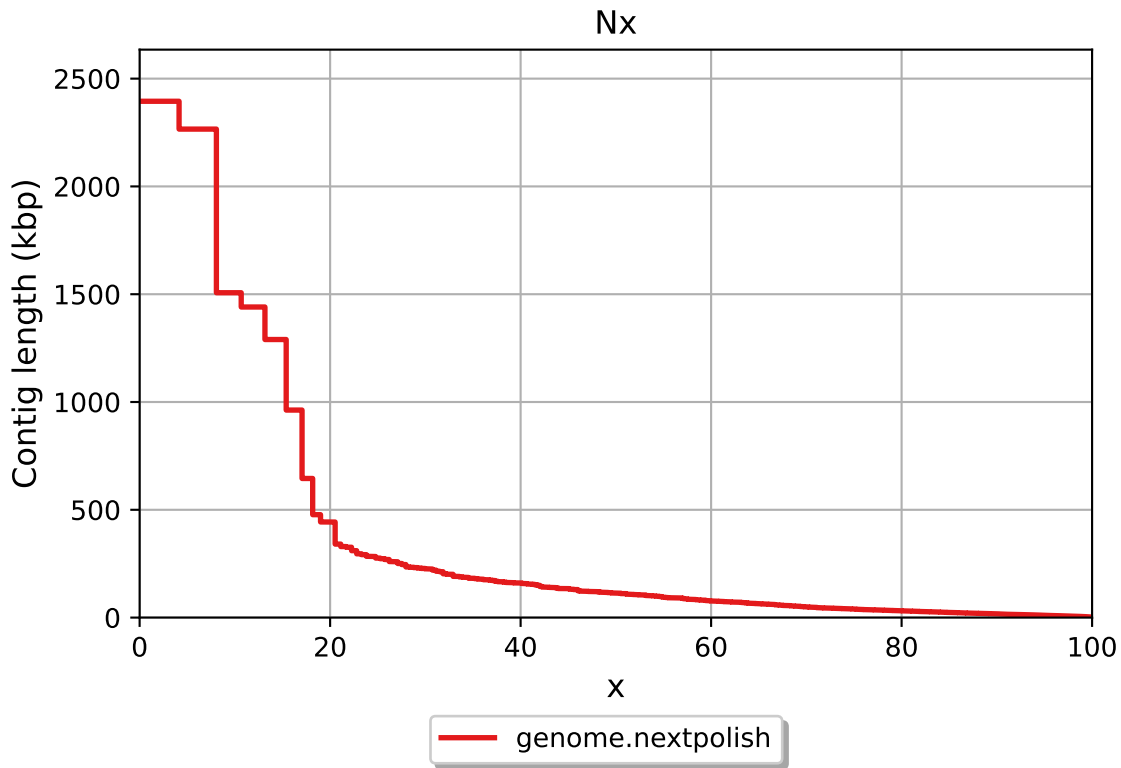


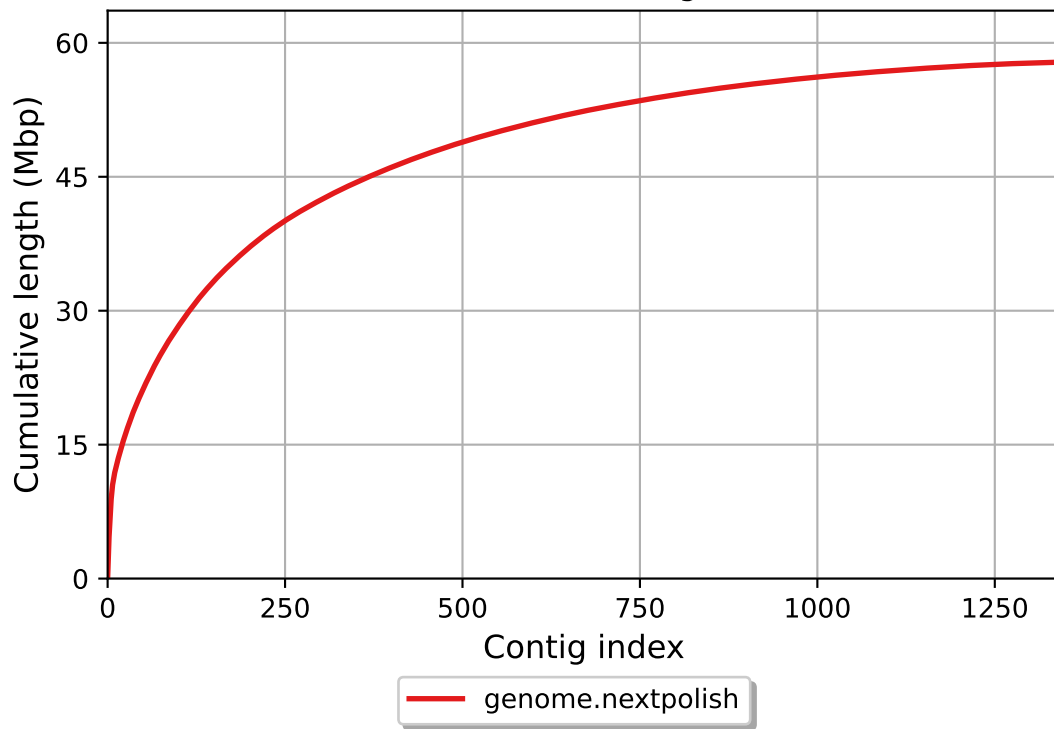
# Report

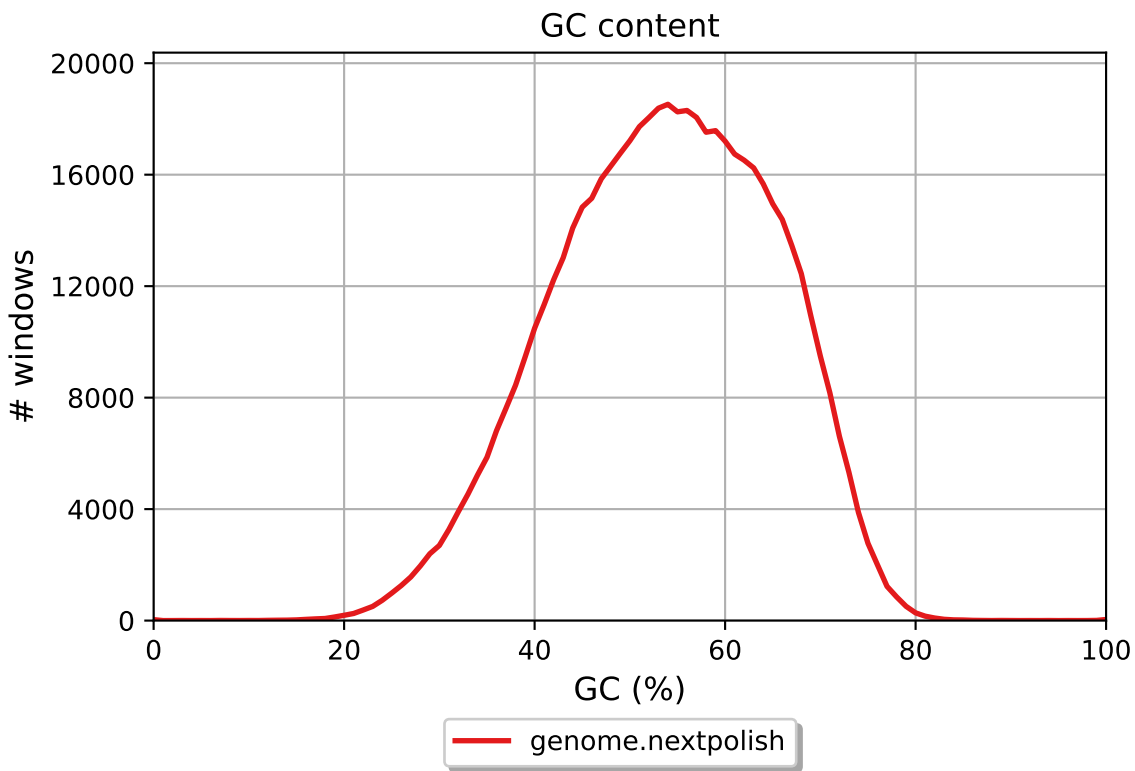
	genome.nextpolish
# contigs (>= 0 bp)	1342
# contigs (>= 1000 bp)	1337
# contigs (>= 5000 bp)	1176
# contigs (>= 10000 bp)	890
# contigs (>= 25000 bp)	496
# contigs (>= 50000 bp)	257
Total length (>= 0 bp)	57824862
Total length (>= 1000 bp)	57821487
Total length (>= 5000 bp)	57276936
Total length (>= 10000 bp)	55205206
Total length (>= 25000 bp)	48789961
Total length (>= 50000 bp)	40438313
# contigs	1342
Largest contig	2394988
Total length	57824862
GC (%)	53.54
N50	113149
N90	17543
auN	407492.4
L50	106
L90	653
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	126 + 3 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

