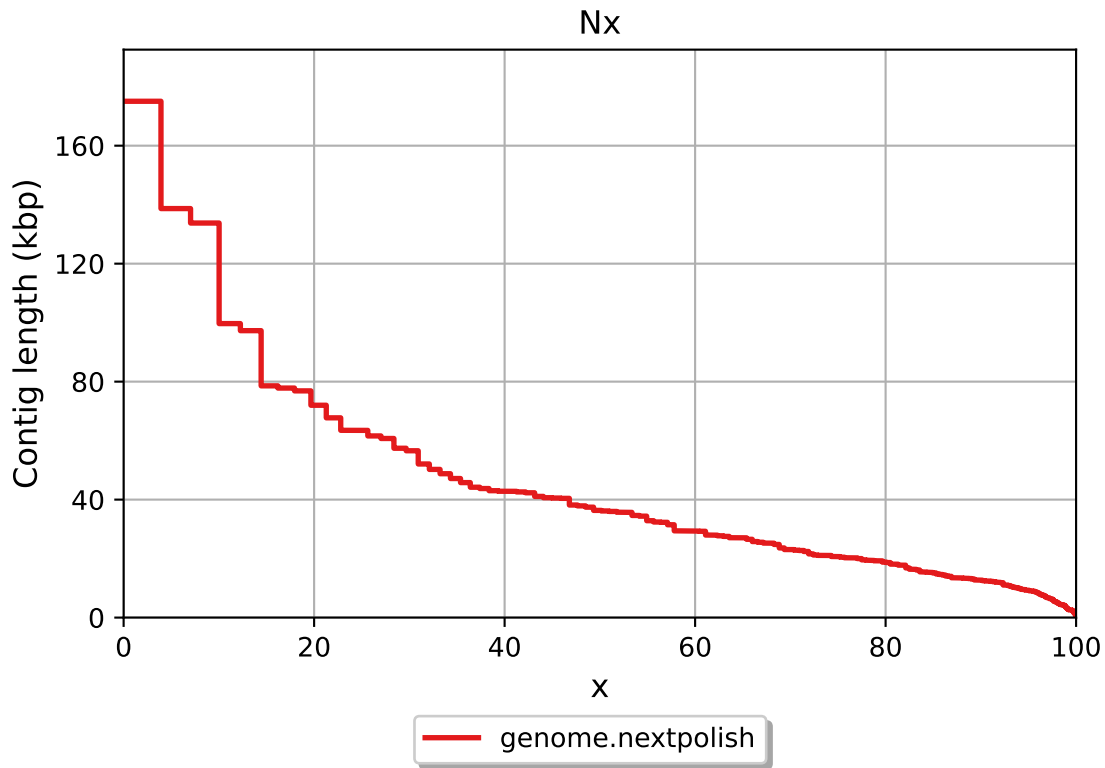


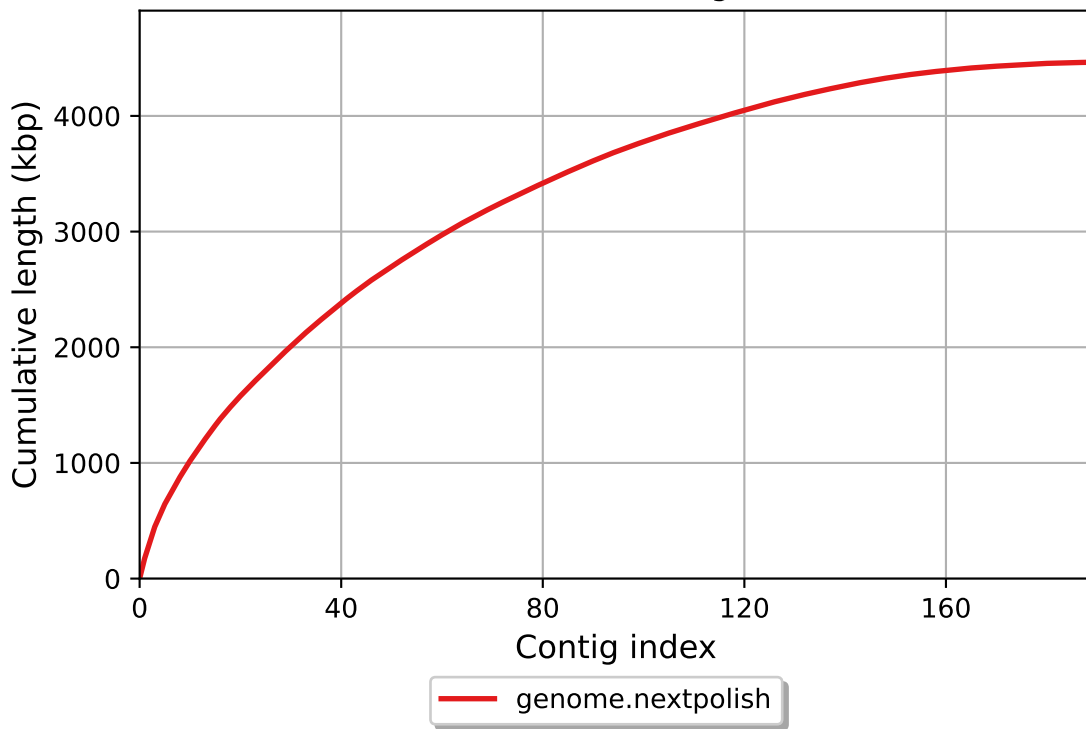
# Report

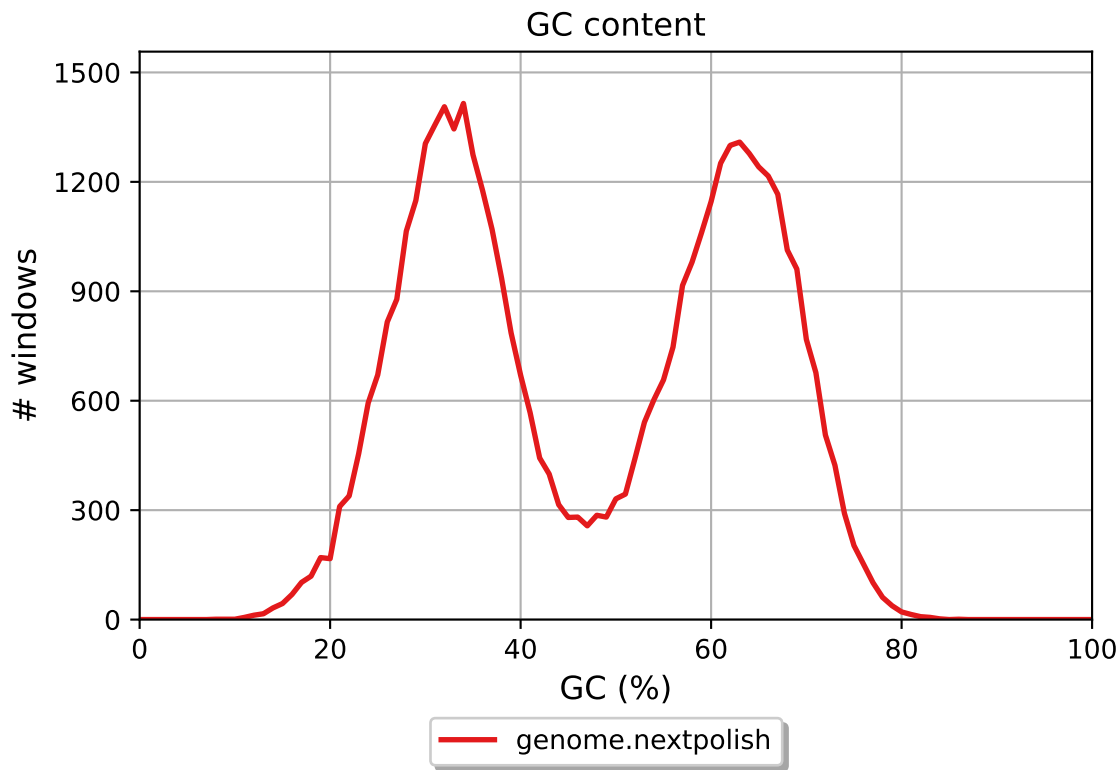
	genome.nextpolish
# contigs (>= 0 bp)	189
# contigs (>= 1000 bp)	185
# contigs (>= 5000 bp)	157
# contigs (>= 10000 bp)	133
# contigs (>= 25000 bp)	63
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	4463797
Total length (>= 1000 bp)	4460917
Total length (>= 5000 bp)	4379372
Total length (>= 10000 bp)	4196082
Total length (>= 25000 bp)	3047793
Total length (>= 50000 bp)	1483027
# contigs	189
Largest contig	175058
Total length	4463797
GC (%)	47.54
N50	36344
N90	12682
auN	49411.6
L50	36
L90	118
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 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

