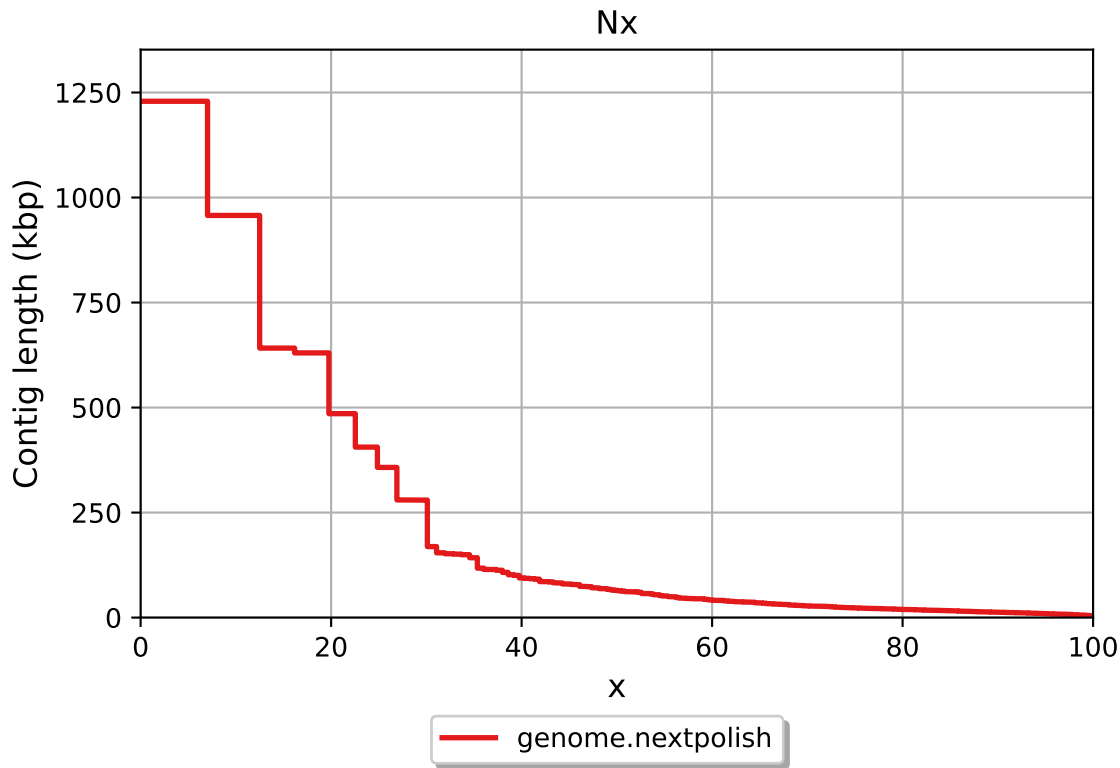


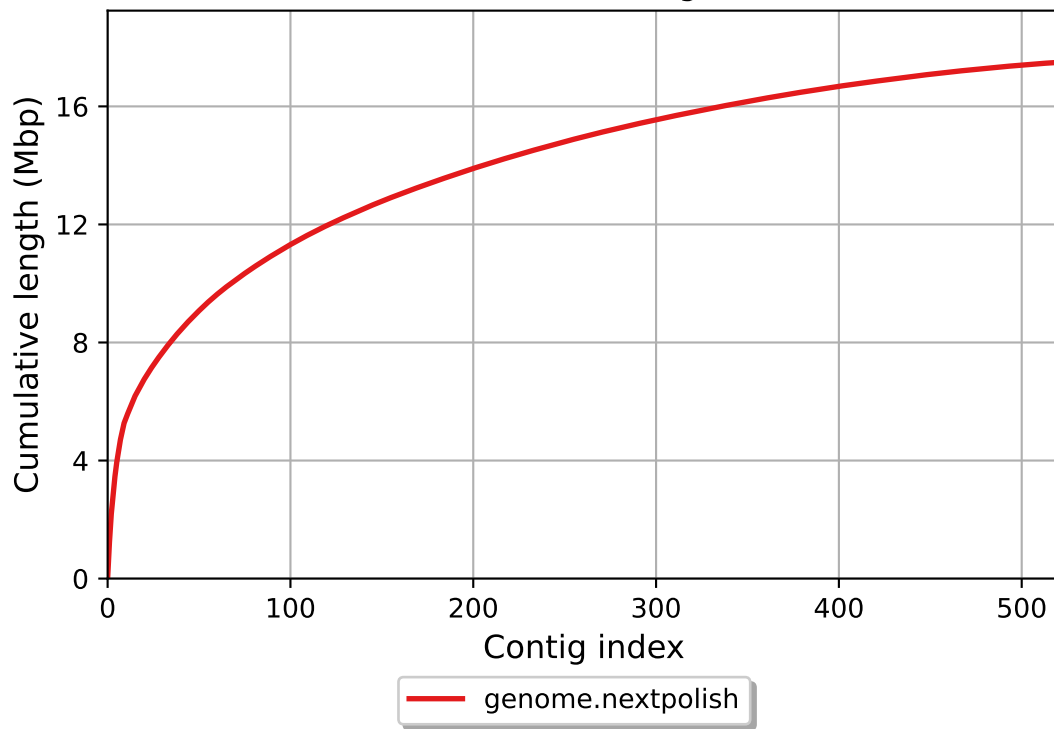
# Report

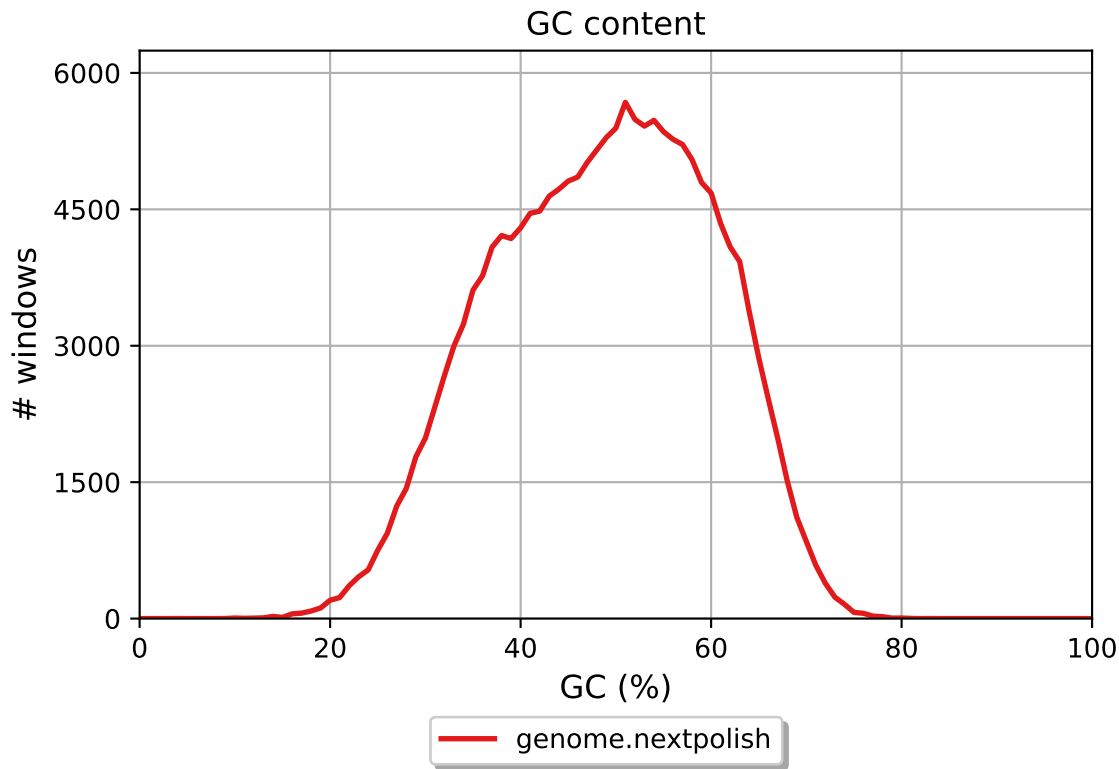
	genome.nextpolish
# contigs (>= 0 bp)	521
# contigs (>= 1000 bp)	521
# contigs (>= 5000 bp)	504
# contigs (>= 10000 bp)	380
# contigs (>= 25000 bp)	149
# contigs (>= 50000 bp)	61
Total length (>= 0 bp)	17494933
Total length (>= 1000 bp)	17494933
Total length (>= 5000 bp)	17420497
Total length (>= 10000 bp)	16487195
Total length (>= 25000 bp)	12762987
Total length (>= 50000 bp)	9689071
# contigs	521
Largest contig	1229155
Total length	17494933
GC (%)	48.54
N50	64814
N90	12692
auN	258813.4
L50	45
L90	316
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	67 + 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

