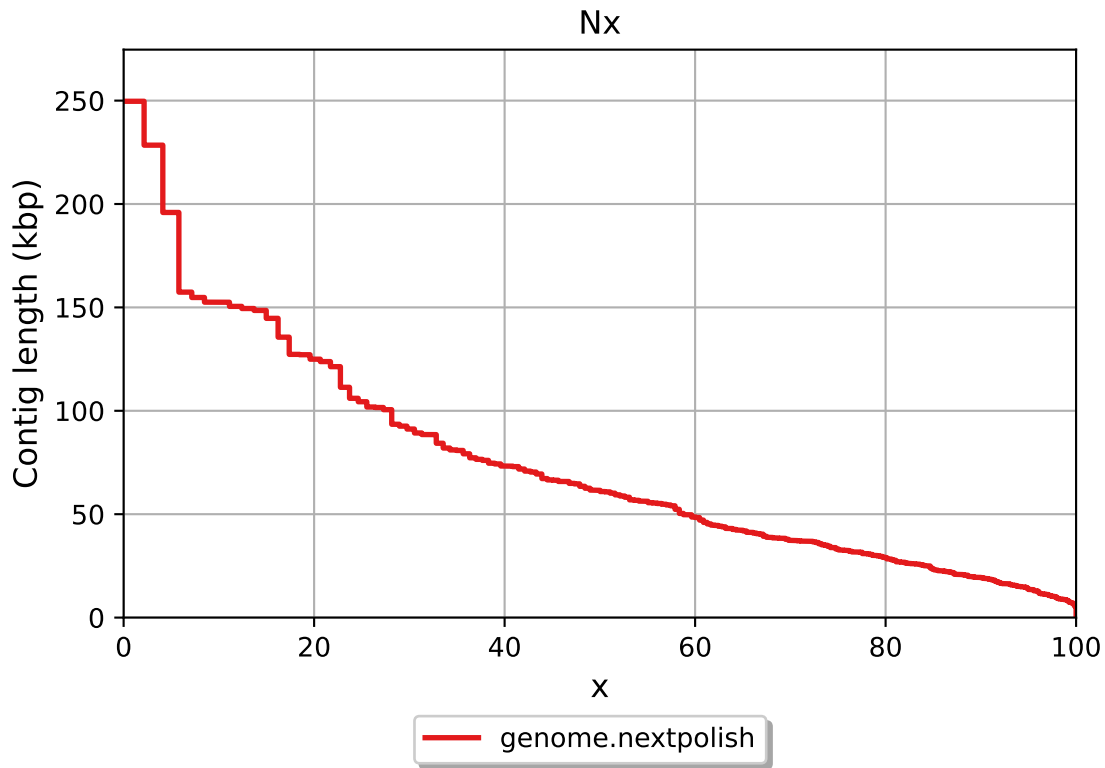


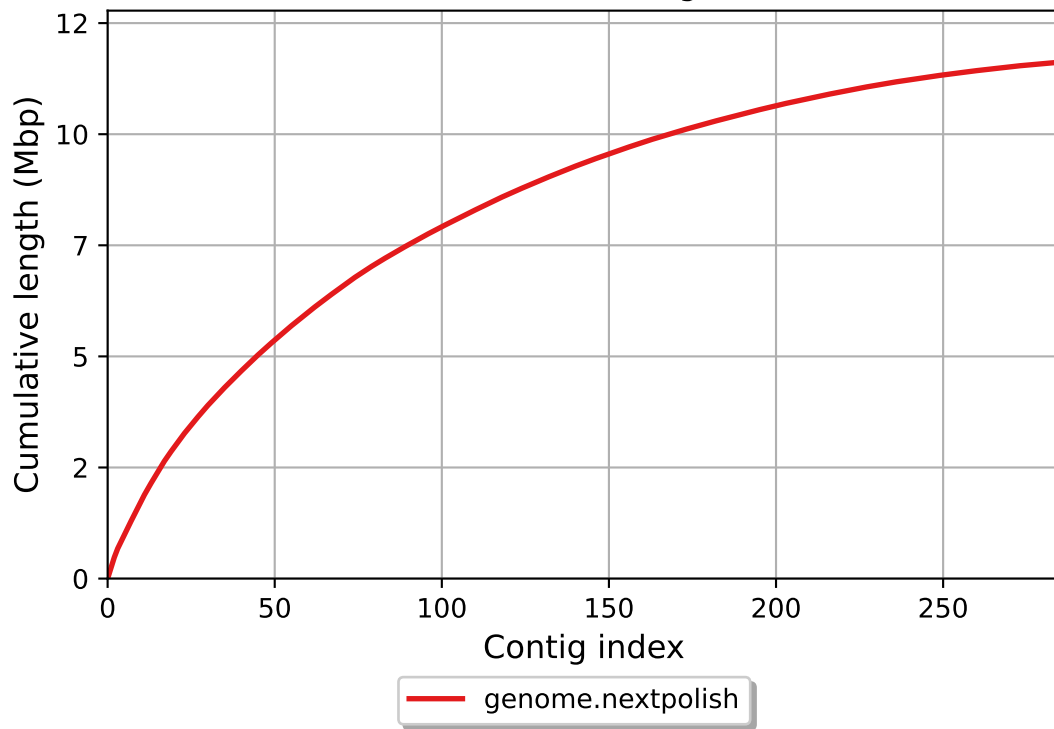
Report

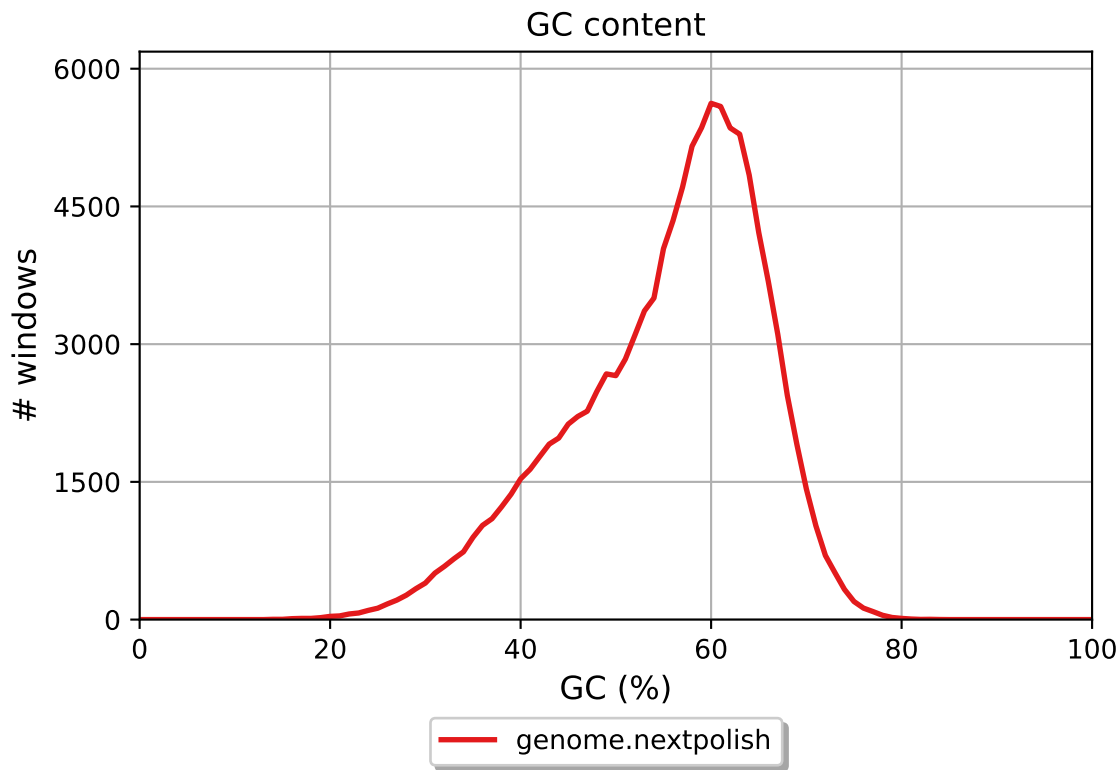
	genome.nextpolish
# contigs (>= 0 bp)	285
# contigs (>= 1000 bp)	285
# contigs (>= 5000 bp)	285
# contigs (>= 10000 bp)	253
# contigs (>= 25000 bp)	159
# contigs (>= 50000 bp)	75
Total length (>= 0 bp)	11618759
Total length (>= 1000 bp)	11618759
Total length (>= 5000 bp)	11618759
Total length (>= 10000 bp)	11364523
Total length (>= 25000 bp)	9788894
Total length (>= 50000 bp)	6829285
# contigs	285
Largest contig	249681
Total length	11618759
GC (%)	55.27
N50	61531
N90	19326
auN	76321.1
L50	57
L90	191
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	42 + 0 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

