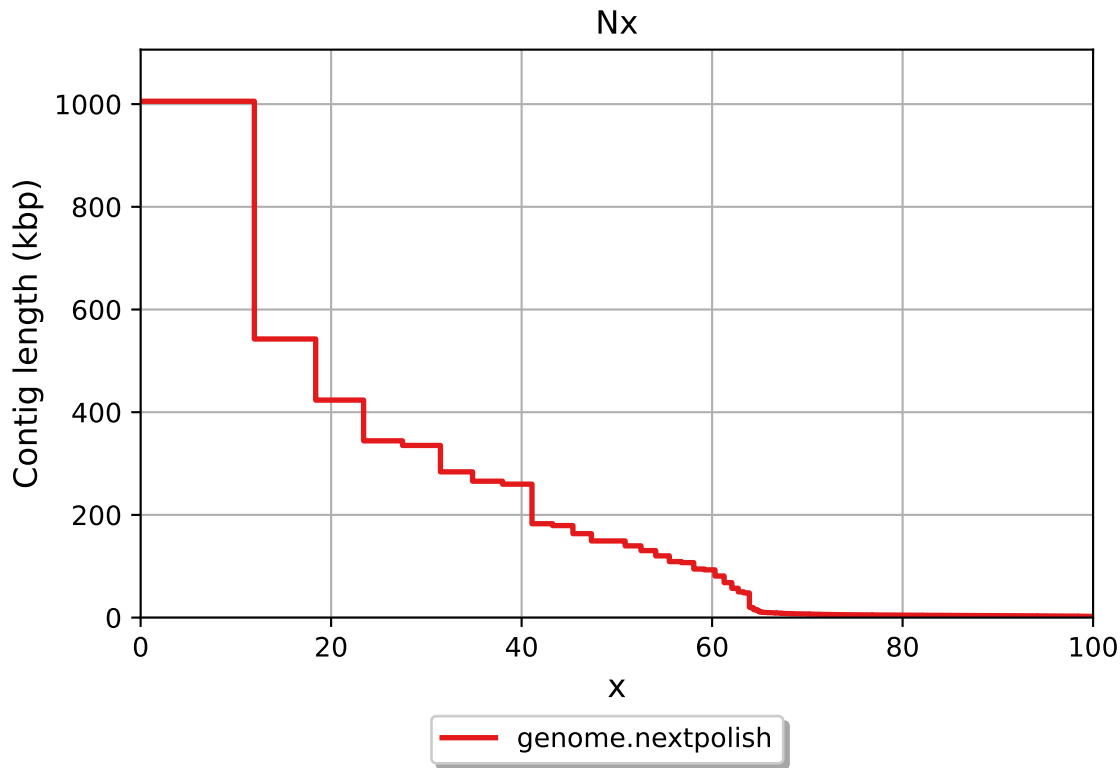


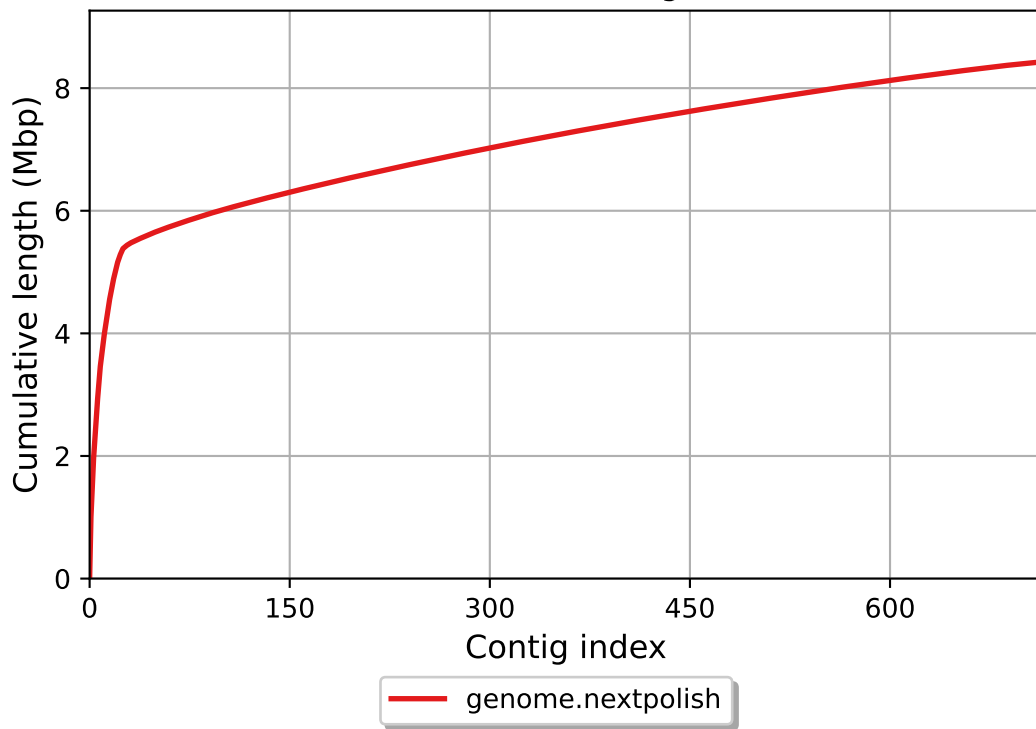
Report

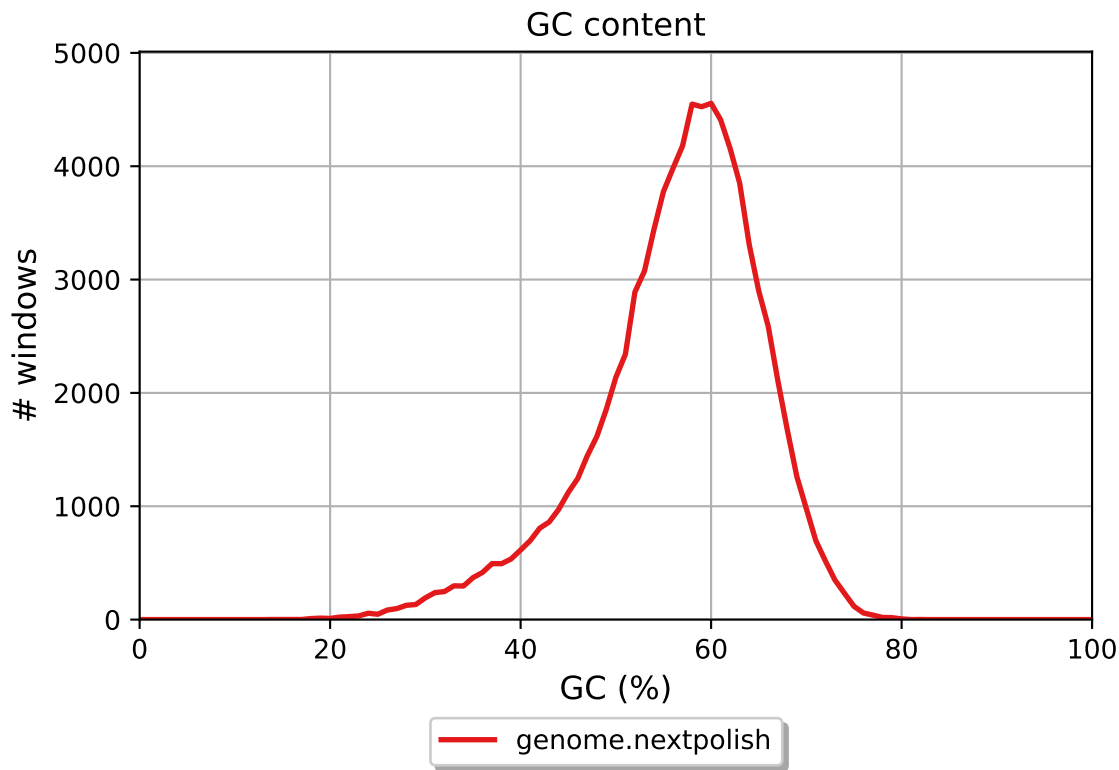
	genome.nextpolish
# contigs (>= 0 bp)	716
# contigs (>= 1000 bp)	708
# contigs (>= 5000 bp)	191
# contigs (>= 10000 bp)	36
# contigs (>= 25000 bp)	25
# contigs (>= 50000 bp)	24
Total length (>= 0 bp)	8422964
Total length (>= 1000 bp)	8417735
Total length (>= 5000 bp)	6515565
Total length (>= 10000 bp)	5529788
Total length (>= 25000 bp)	5382650
Total length (>= 50000 bp)	5334643
# contigs	714
Largest contig	1005542
Total length	8422033
GC (%)	56.44
N50	149226
N90	3685
auN	261023.1
L50	13
L90	439
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	27 + 7 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

