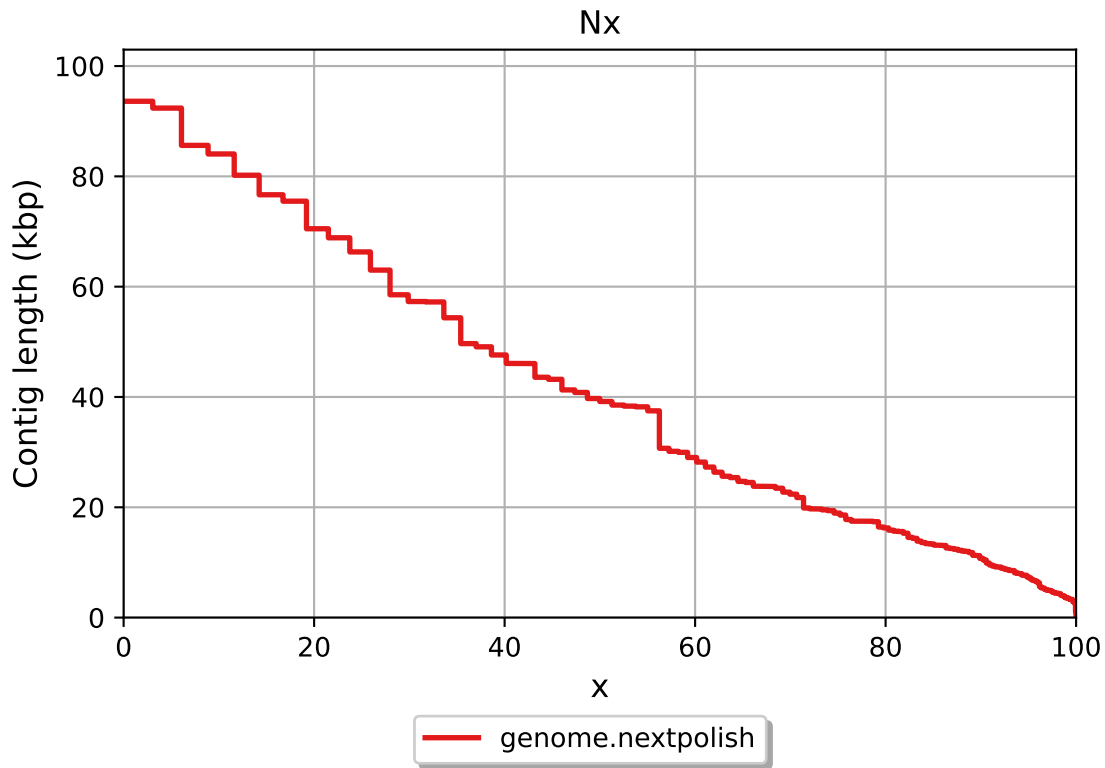
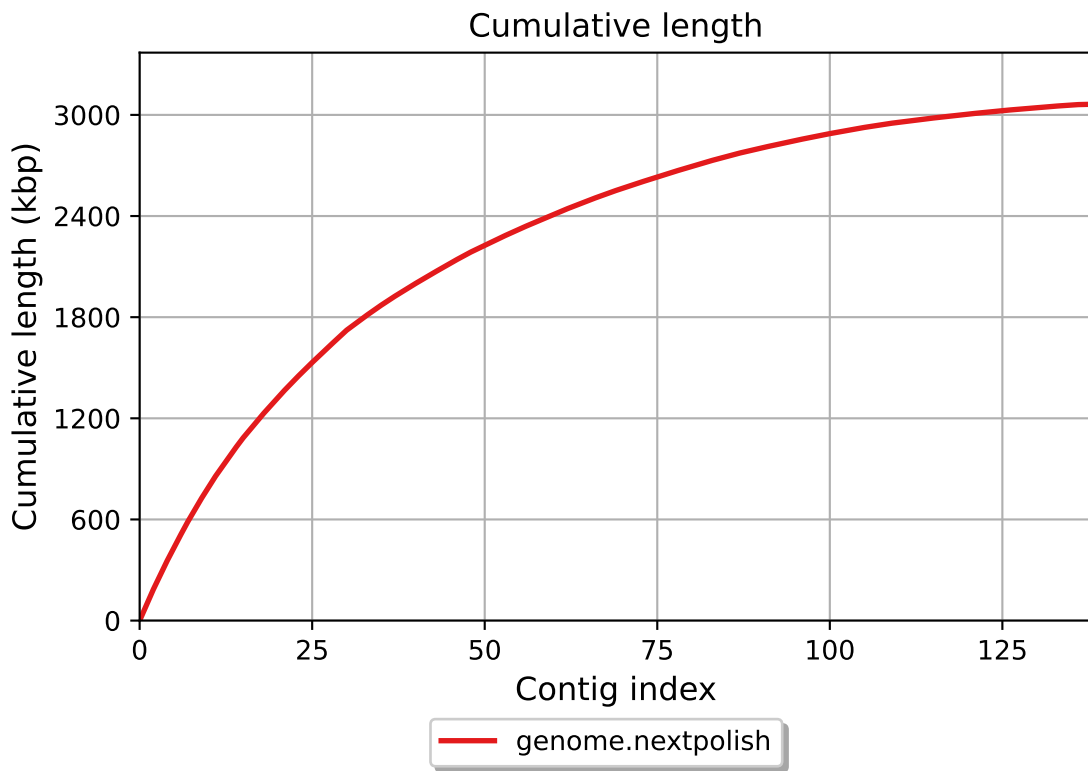


# Report

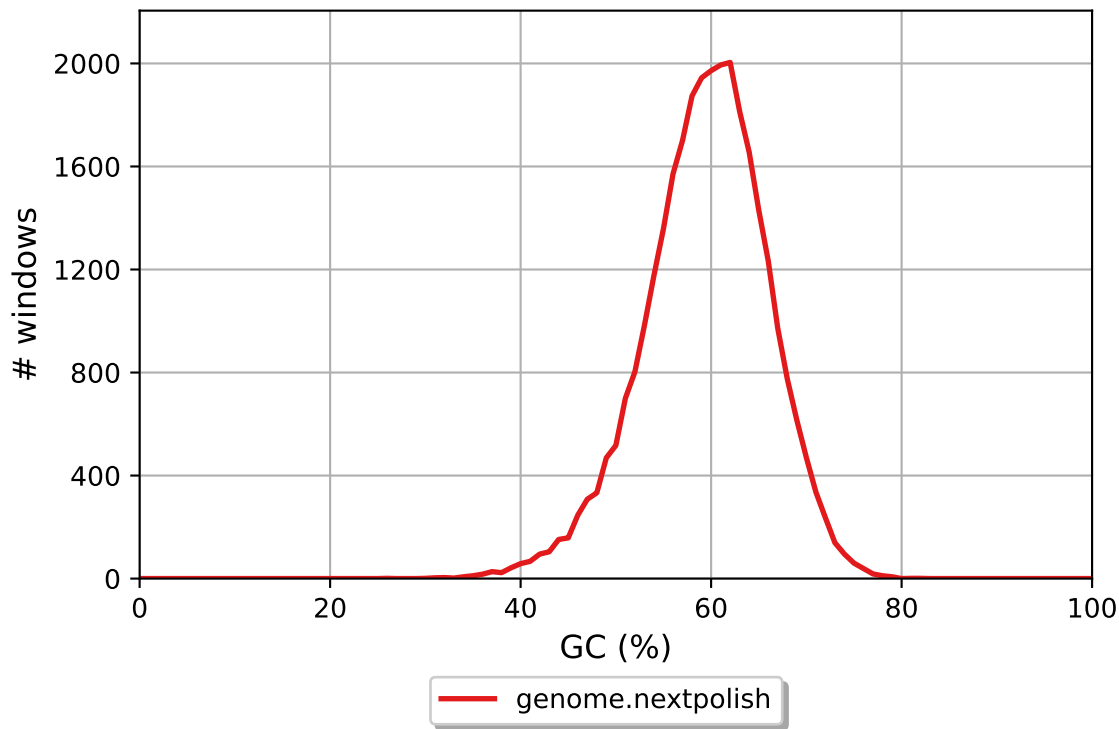
	genome.nextpolish
# contigs (>= 0 bp)	138
# contigs (>= 1000 bp)	137
# contigs (>= 5000 bp)	113
# contigs (>= 10000 bp)	87
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	3063061
Total length (>= 1000 bp)	3062238
Total length (>= 5000 bp)	2971714
Total length (>= 10000 bp)	2774179
Total length (>= 25000 bp)	1975660
Total length (>= 50000 bp)	1084110
# contigs	138
Largest contig	93613
Total length	3063061
GC (%)	59.40
N50	39182
N90	10743
auN	42879.5
L50	26
L90	86
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).





## GC content



genome.nextpolish GC content

