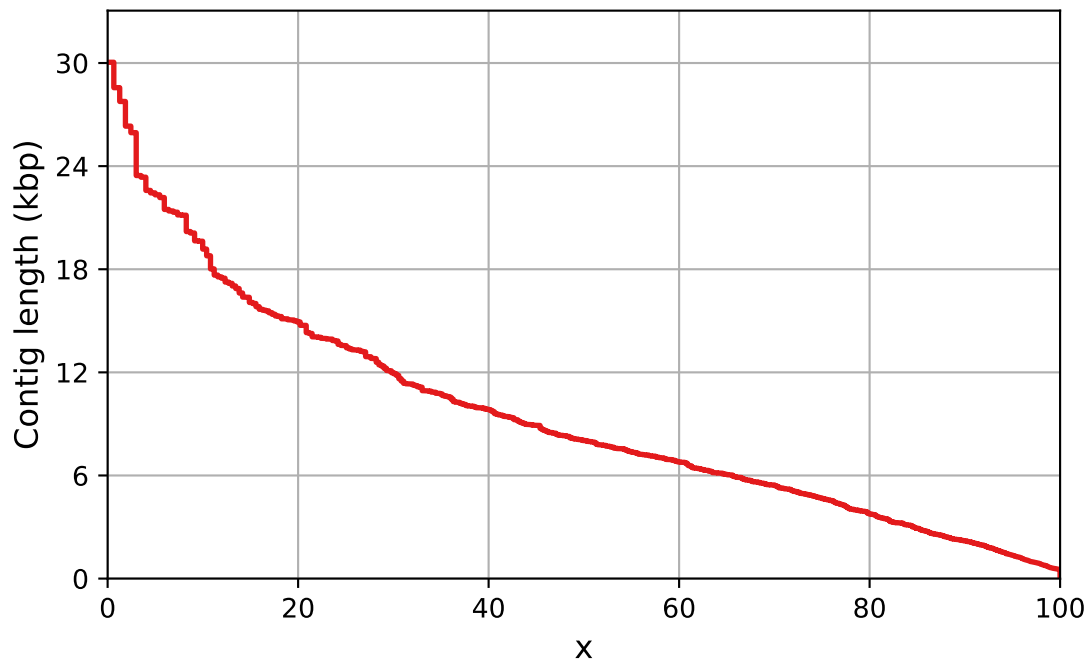


# Report

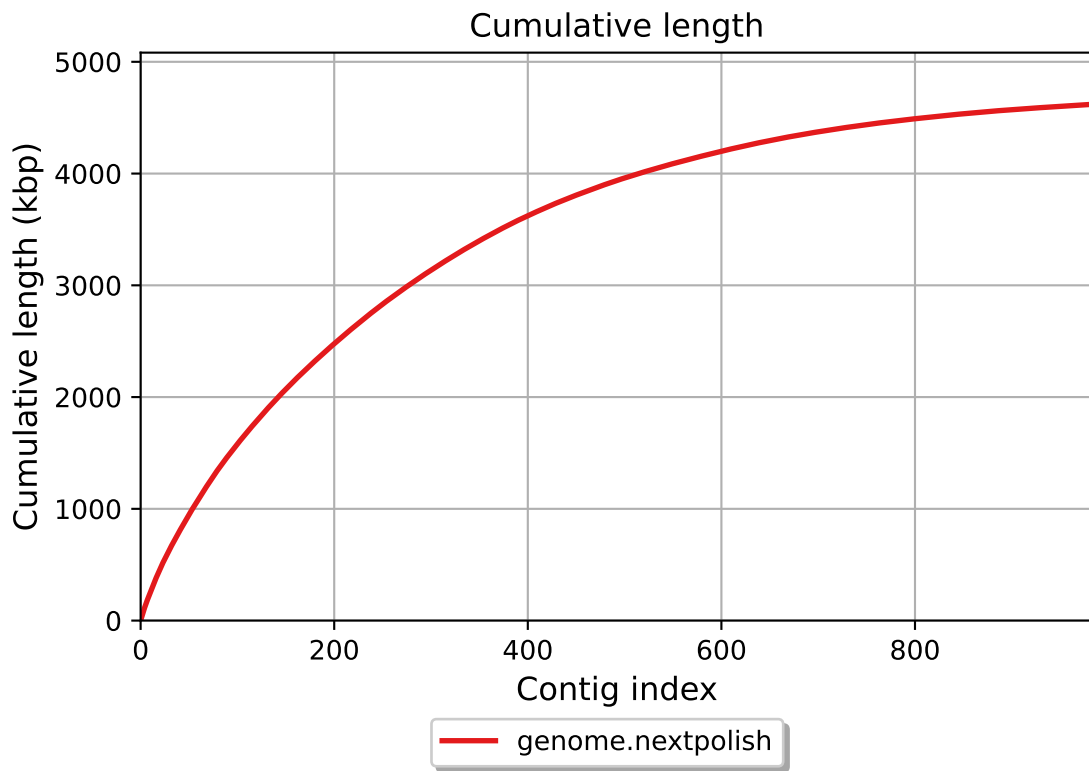
	genome.nextpolish
# contigs (>= 0 bp)	987
# contigs (>= 1000 bp)	785
# contigs (>= 5000 bp)	341
# contigs (>= 10000 bp)	119
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4621552
Total length (>= 1000 bp)	4475921
Total length (>= 5000 bp)	3354788
Total length (>= 10000 bp)	1777682
Total length (>= 25000 bp)	138647
Total length (>= 50000 bp)	0
# contigs	984
Largest contig	30043
Total length	4620095
GC (%)	56.92
N50	8032
N90	2199
auN	9521.5
L50	179
L90	581
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

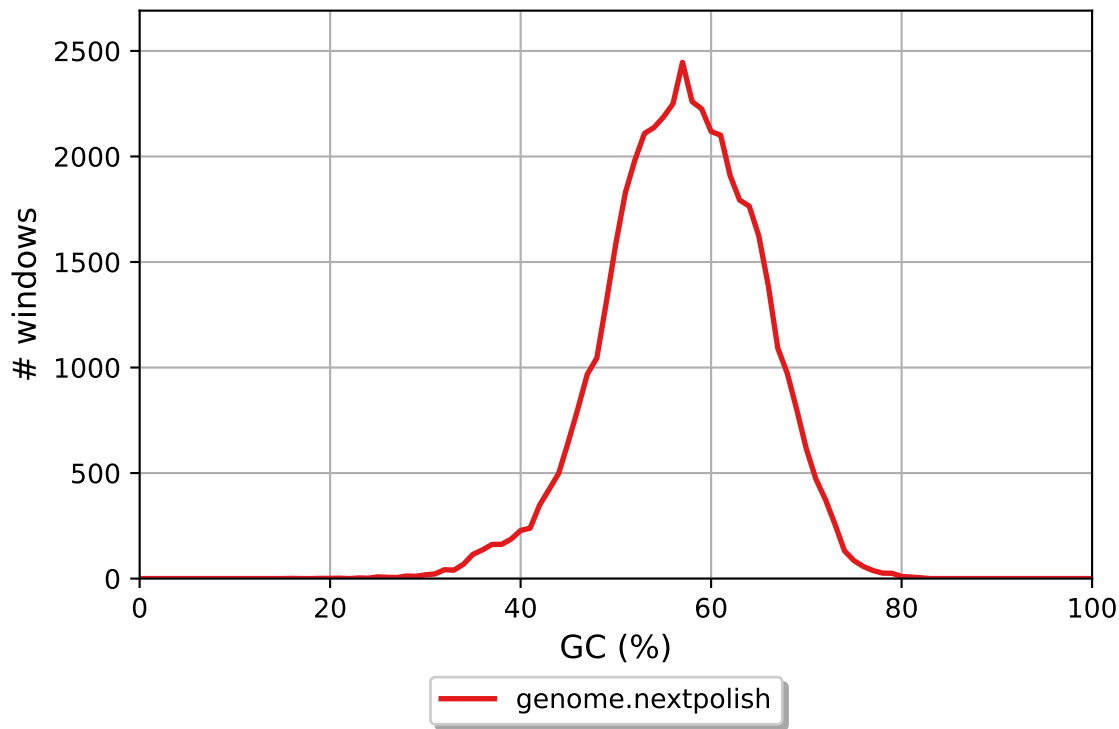
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

