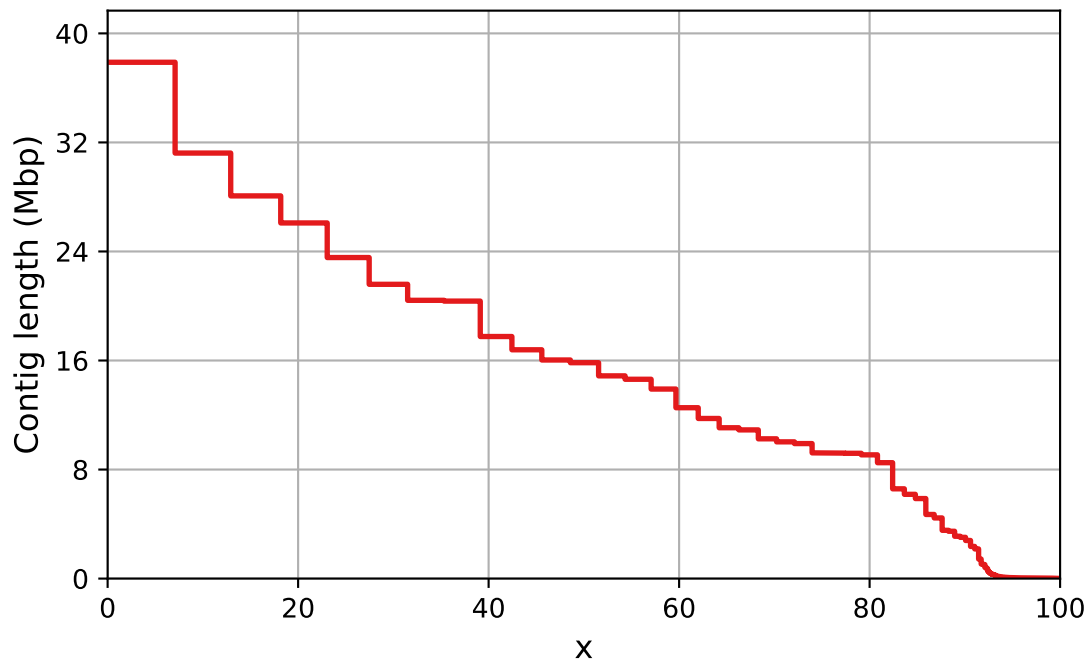


Report

	maternalpaternal.asm.dip.hap1.p_ctg
# contigs (>= 0 bp)	823
# contigs (>= 1000 bp)	823
# contigs (>= 5000 bp)	823
# contigs (>= 10000 bp)	823
# contigs (>= 25000 bp)	787
# contigs (>= 50000 bp)	267
Total length (>= 0 bp)	534683615
Total length (>= 1000 bp)	534683615
Total length (>= 5000 bp)	534683615
Total length (>= 10000 bp)	534683615
Total length (>= 25000 bp)	533888662
Total length (>= 50000 bp)	514753554
# contigs	823
Largest contig	37880845
Total length	534683615
GC (%)	35.47
N50	15838561
N75	9224513
L50	12
L75	23
# N's per 100 kbp	0.00

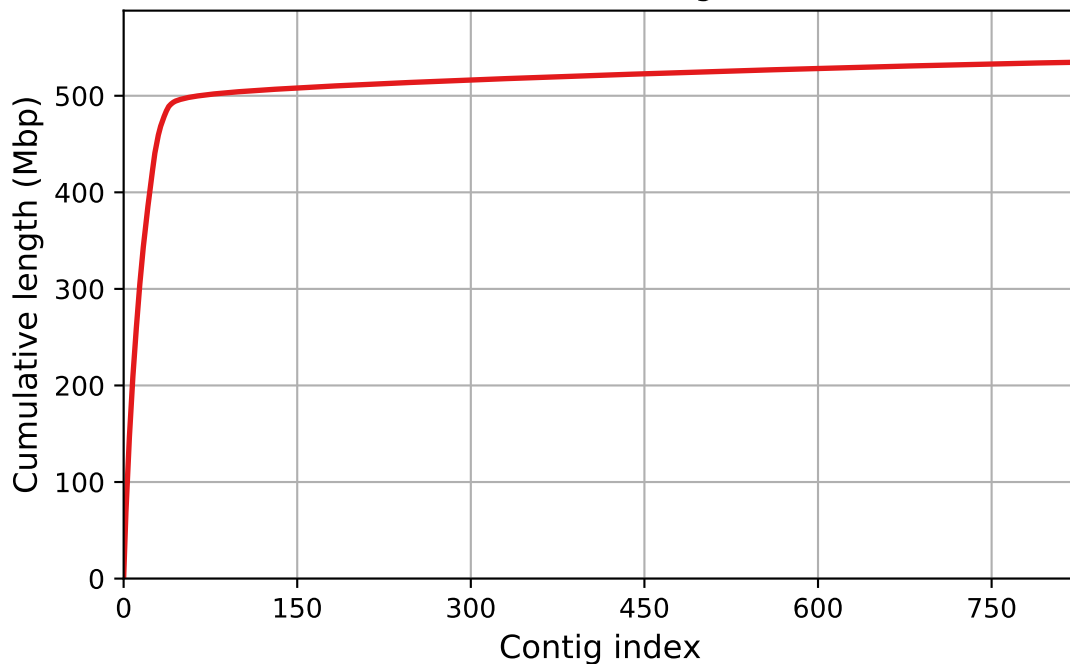
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



maternalpaternal.asm.dip.hap1.p_ctg

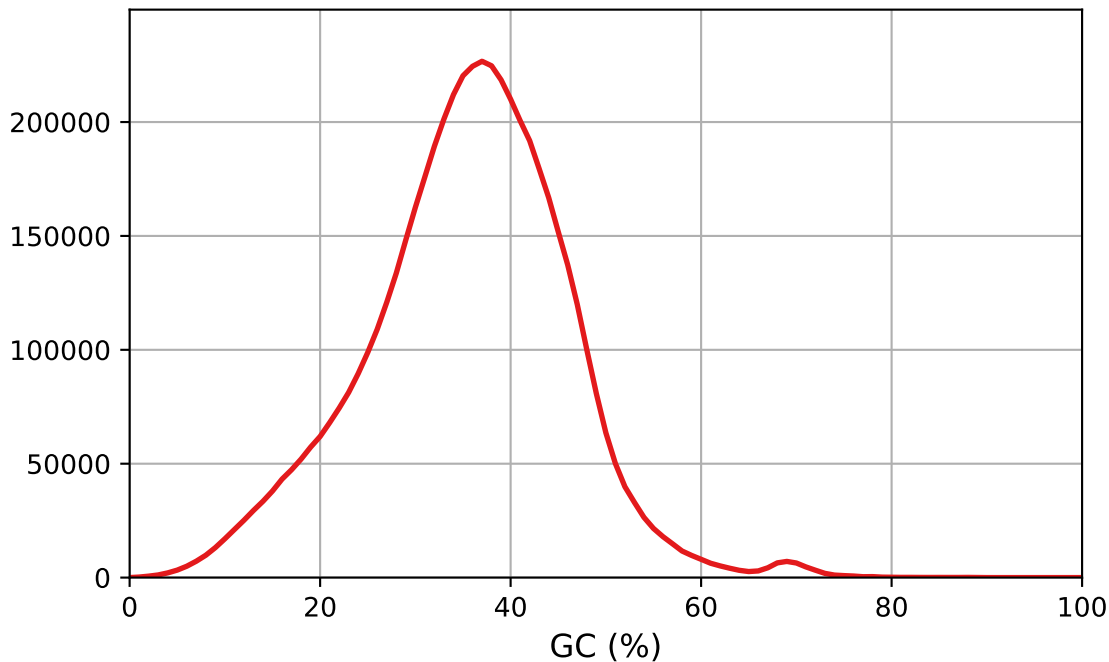
Cumulative length



— maternalpaternal.asm.dip.hap1.p_ctg

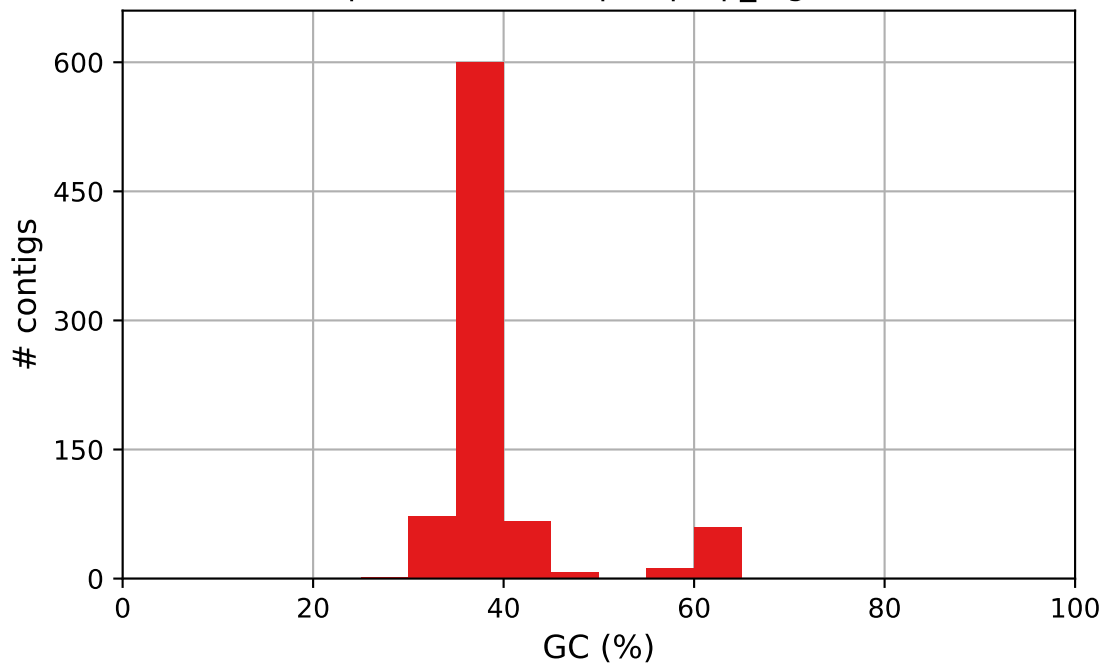
windows

GC content



maternalpaternal.asm.dip.hap1.p_ctg

maternalpaternal.asm.dip.hap1.p_ctg GC content



maternalpaternal.asm.dip.hap1.p_ctg