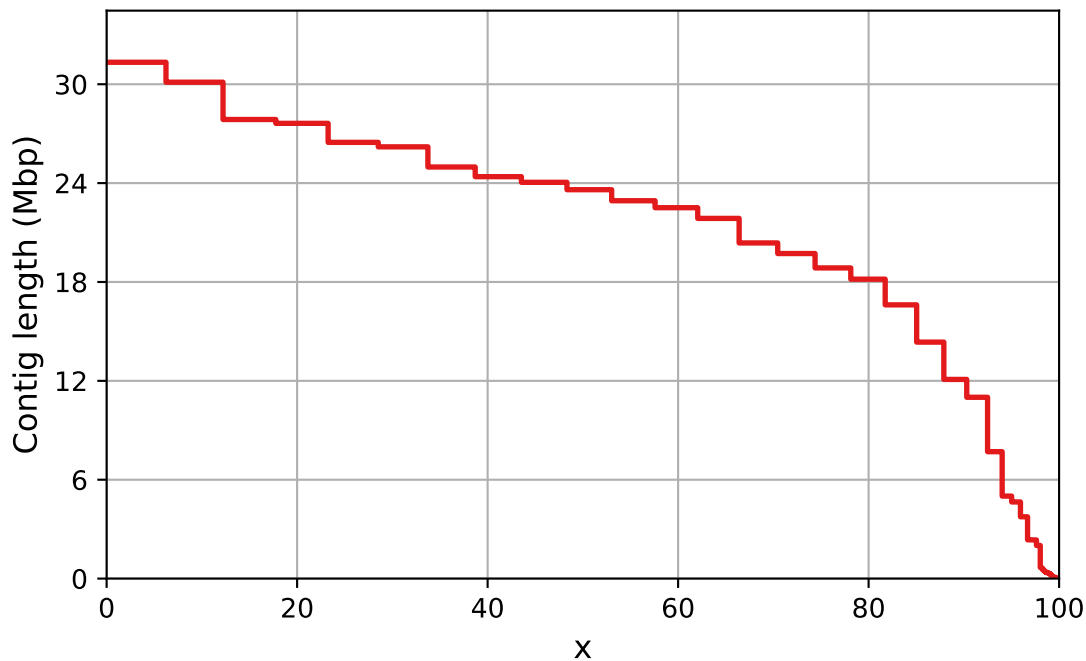


Report

	maternalpaternal.asm.dip.hap2.p_ctg
# contigs (>= 0 bp)	116
# contigs (>= 1000 bp)	116
# contigs (>= 5000 bp)	116
# contigs (>= 10000 bp)	116
# contigs (>= 25000 bp)	103
# contigs (>= 50000 bp)	70
Total length (>= 0 bp)	502852466
Total length (>= 1000 bp)	502852466
Total length (>= 5000 bp)	502852466
Total length (>= 10000 bp)	502852466
Total length (>= 25000 bp)	502591852
Total length (>= 50000 bp)	501424838
# contigs	116
Largest contig	31333120
Total length	502852466
GC (%)	35.28
N50	23599708
N75	18852907
L50	10
L75	16
# N's per 100 kbp	0.00

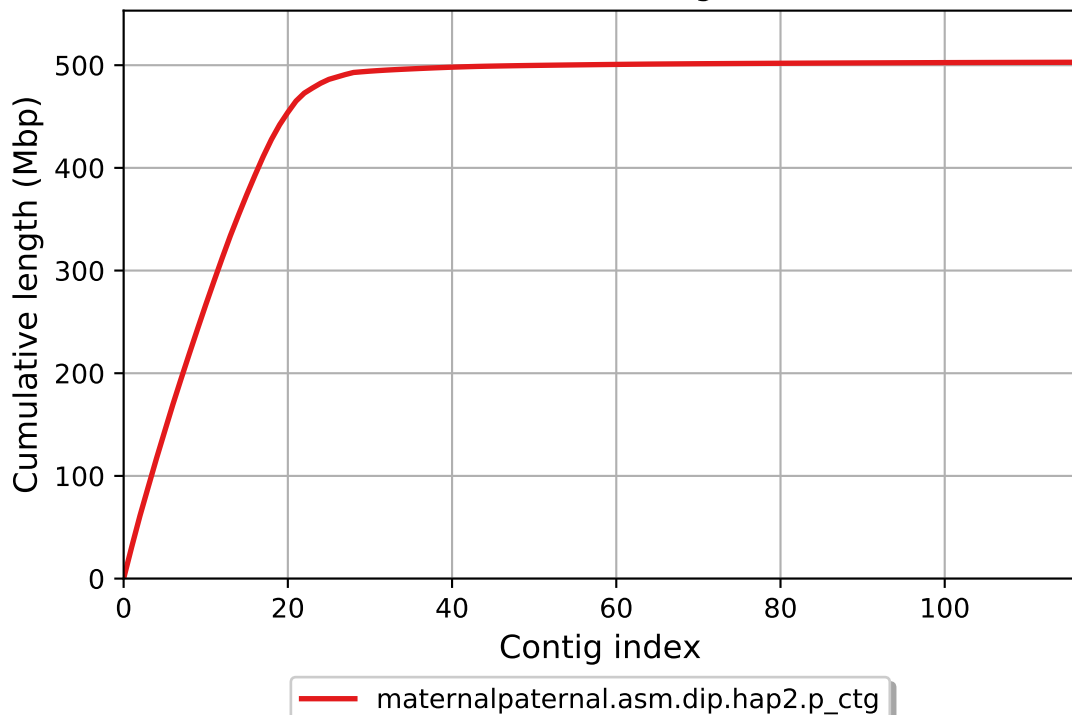
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

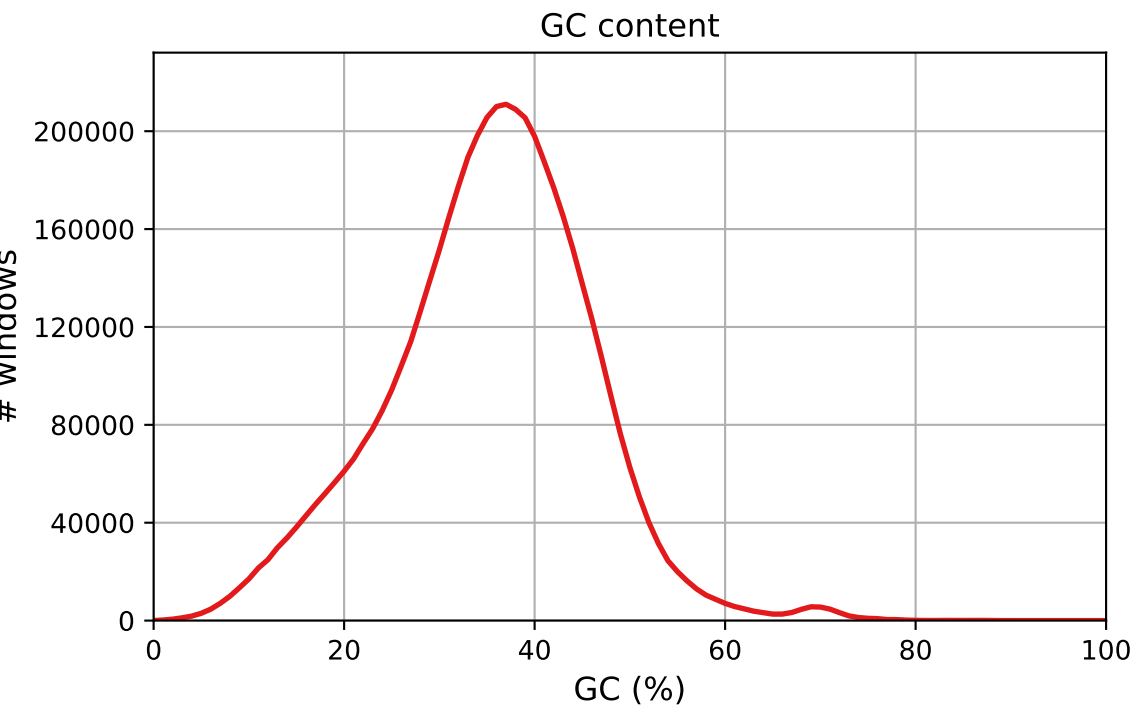
Nx



maternalpaternal.asm.dip.hap2.p_ctg

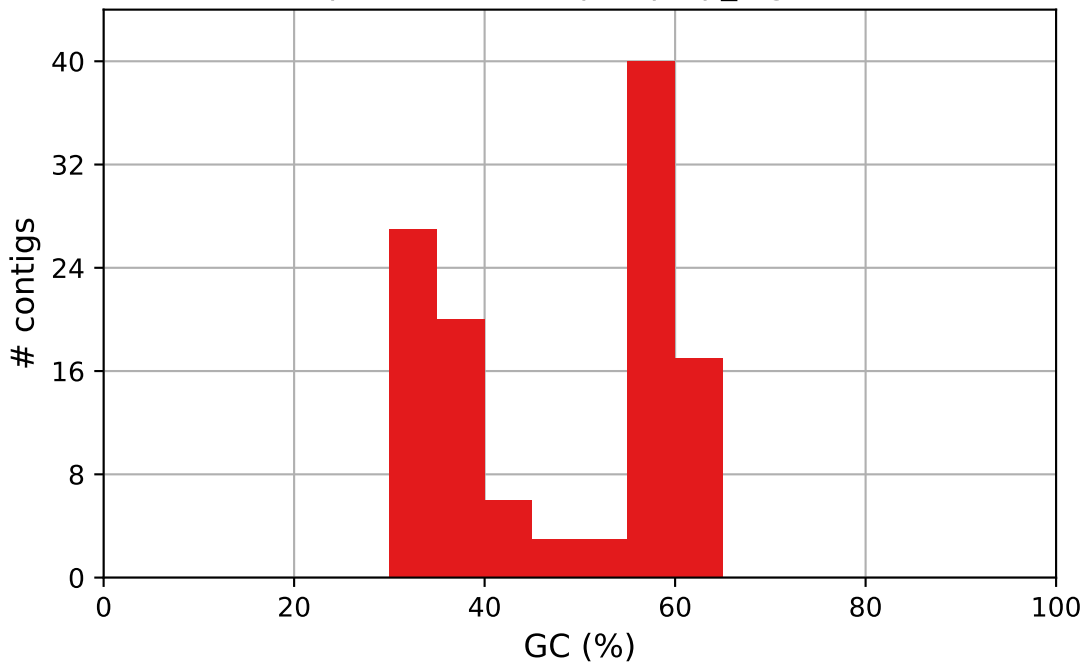
Cumulative length





— maternalpaternal.asm.dip.hap2.p_ctg

maternalpaternal.asm.dip.hap2.p_ctg GC content



maternalpaternal.asm.dip.hap2.p_ctg