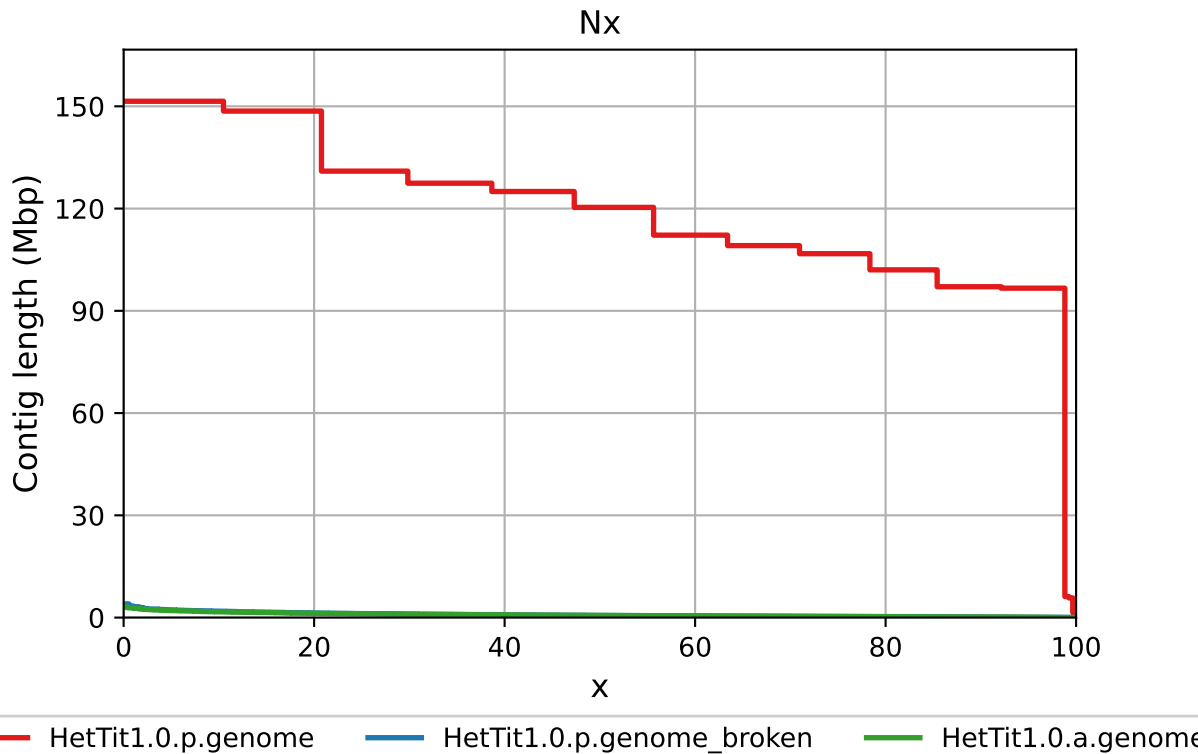


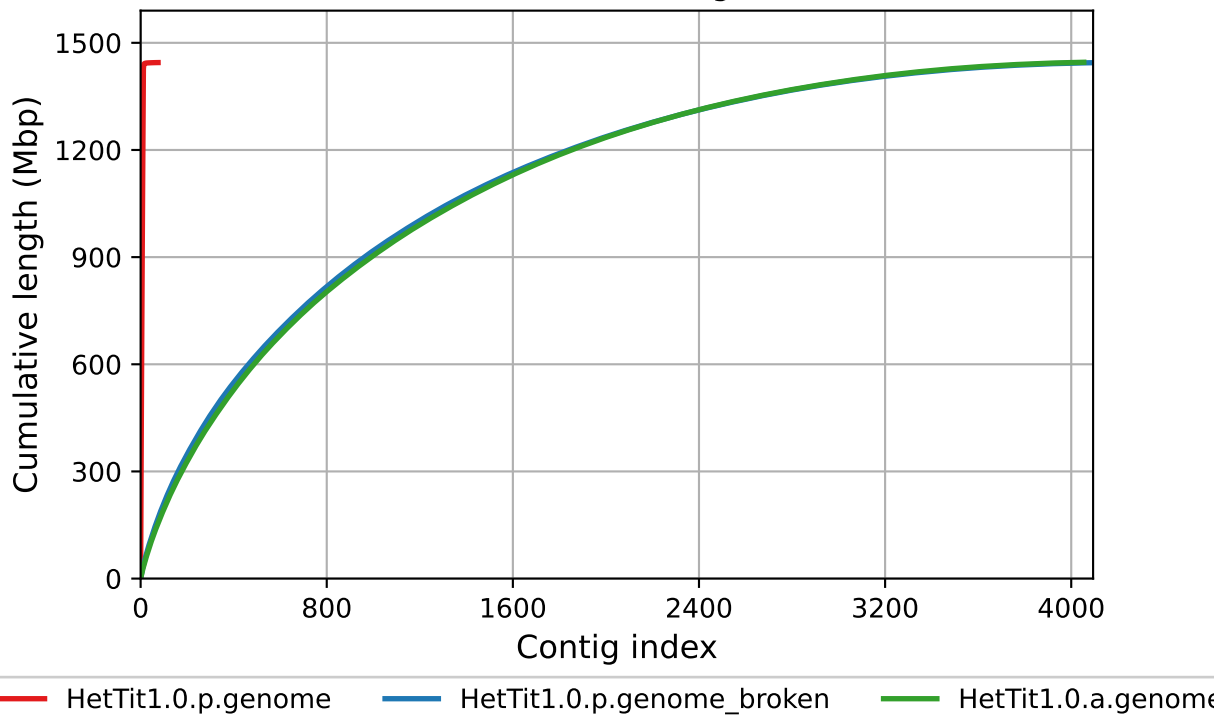
Report

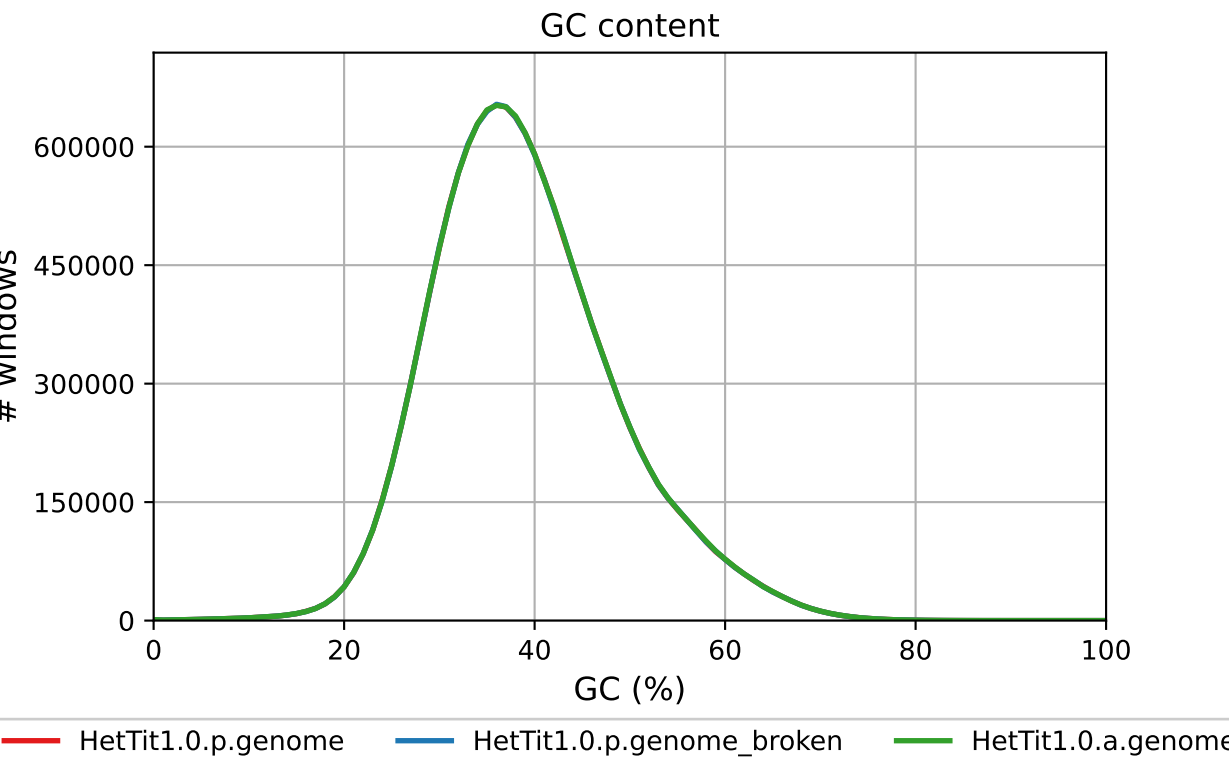
	HetTit1.0.p.genome	HetTit1.0.p.genome_broken	HetTit1.0.a.genome
# contigs (≥ 0 bp)	75	4094	4054
# contigs (≥ 1000 bp)	75	4094	4054
# contigs (≥ 5000 bp)	75	4094	4054
# contigs (≥ 10000 bp)	74	4080	4043
# contigs (≥ 25000 bp)	48	3874	3871
# contigs (≥ 50000 bp)	28	3533	3529
Total length (≥ 0 bp)	1444697970	1444296070	1445542320
Total length (≥ 1000 bp)	1444697970	1444296070	1445542320
Total length (≥ 5000 bp)	1444697970	1444296070	1445542320
Total length (≥ 10000 bp)	1444688748	1444173534	1445445168
Total length (≥ 25000 bp)	1444264645	1440496826	1442235848
Total length (≥ 50000 bp)	1443604017	1427755254	1429703052
# contigs	75	4094	4054
Largest contig	151479759	4037799	3160621
Total length	1444697970	1444296070	1445542320
GC (%)	39.10	39.10	39.11
N50	120343728	638359	631942
N90	97068442	169648	176305
auN	120214200.6	852126.3	803964.5
L50	6	639	664
L90	11	2326	2332
# N's per 100 kbp	27.82	0.00	0.00

All statistics are based on contigs of size ≥ 3000 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

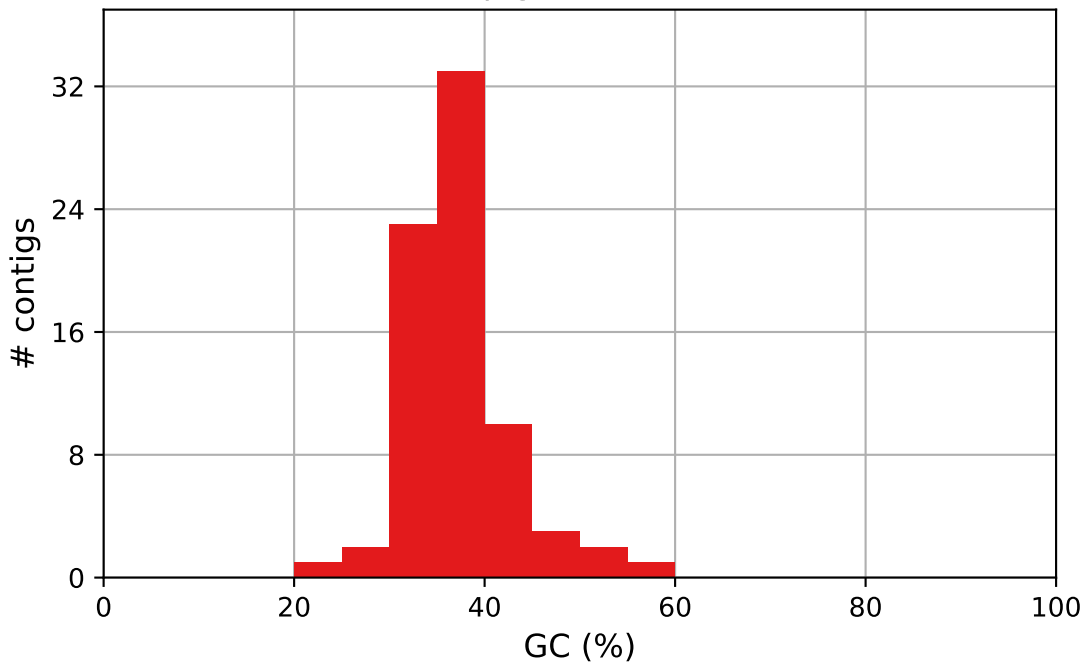


Cumulative length



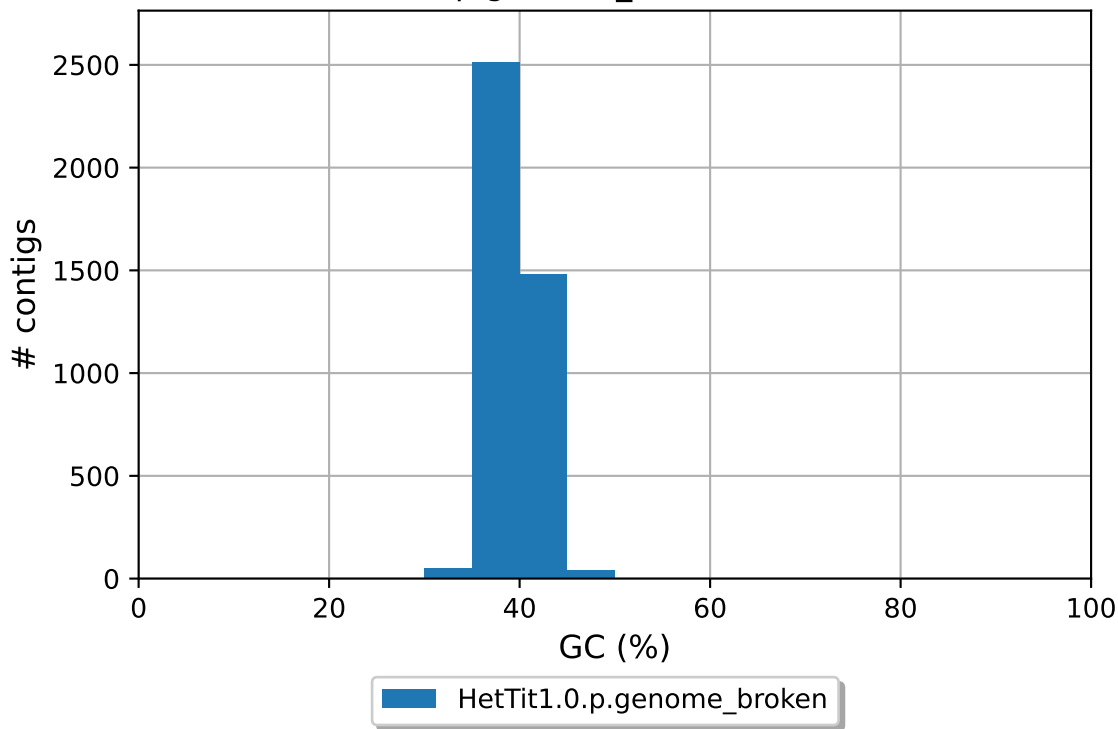


HetTit1.0.p.genome GC content



HetTit1.0.p.genome

HetTit1.0.p.genome_broken GC content



HetTit1.0.a.genome GC content

