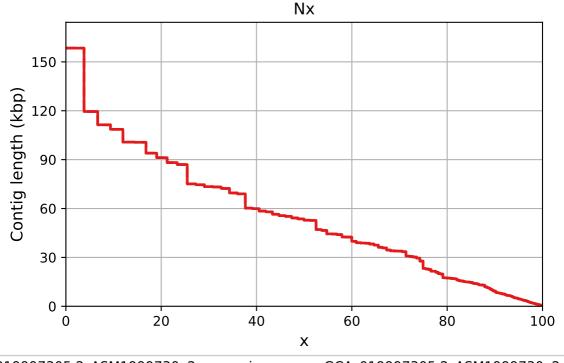
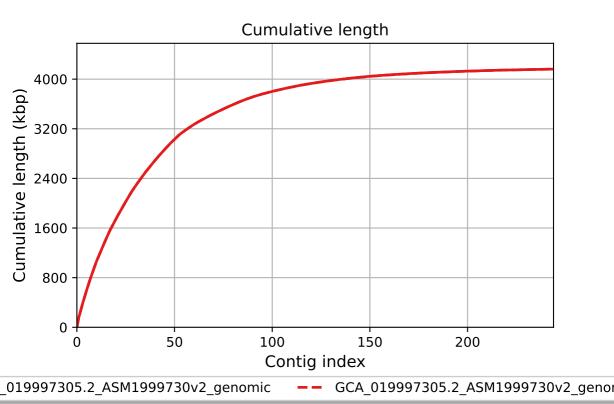
Report

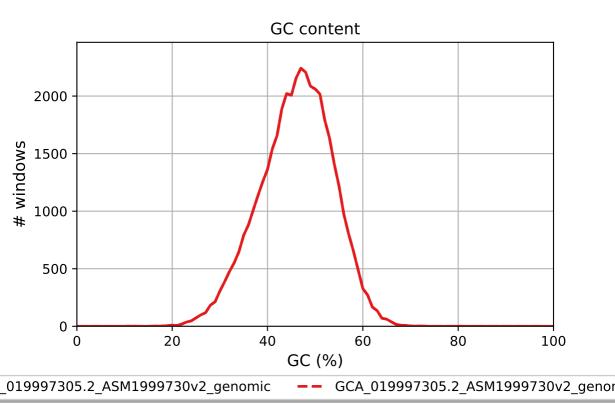
	GCA_019997305.2_ASM1999730v2_genomic	GCA_019997305.2_ASM1999730v2_genomic_broken
# contigs (>= 0 bp)	259	-
# contigs (>= 1000 bp)	205	206
# contigs (>= 5000 bp)	122	122
# contigs (>= 10000 bp)	91	91
# contigs (>= 25000 bp)	53	53
# contigs (>= 50000 bp)	28	28
Total length (>= 0 bp)	4167230	-
Total length (>= 1000 bp)	4134783	4134689
Total length (>= 5000 bp)	3940934	3940934
Total length (>= 10000 bp)	3723926	3723926
Total length (>= 25000 bp)	3118536	3118536
Total length (>= 50000 bp)	2184205	2184205
# contigs	243	244
Largest contig	158427	158427
Total length	4161333	4161239
GC (%)	45.99	45.99
N50	52820	52820
N90	9319	9319
auN	56663.4	56662.3
L50	27	27
L90	94	94
# N's per 100 kbp	2.26	0.00

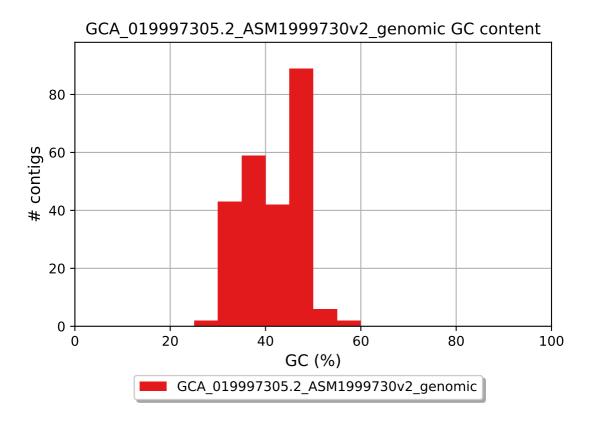
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



____ GCA_019997305.2_ASM1999730v2_genomic ___ GCA_019997305.2_ASM1999730v2_genor







GCA 019997305.2 ASM1999730v2 genomic broken GC content

