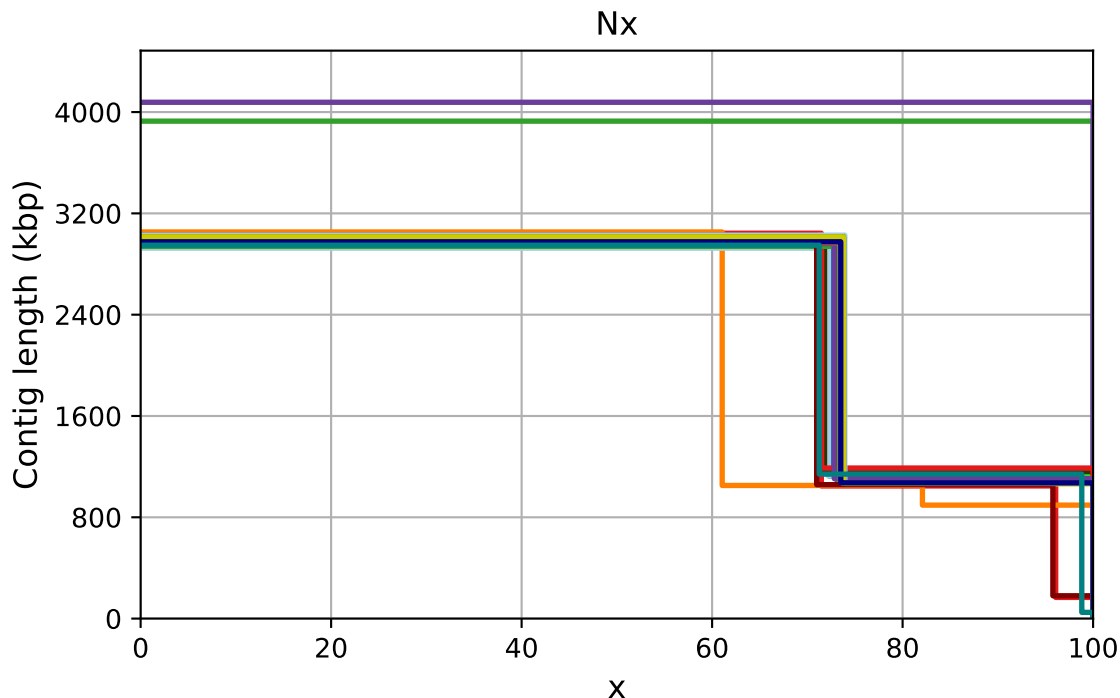


Report

	GCF_000765415.1_ASM76541v1_genomic	GCF_000788715.2_ASM78871v2_genomic	GCF_000969235.1_ASM96923v1_genomic	GCF_000969265.1_ASM96926v1_genomic	GCF_001045415.1_ASM104541v1_genomic	GCF_001250235.2_5174_7_8_v2_genomic	GCF_001854425.1_ASM185442v1_genomic	GCF_001887395.1_ASM188739v1_genomic	GCF_001887415.1_ASM188741v1_genomic	GCF_001887435.1_ASM188743v1_genomic	GCF_001887455.1_ASM188745v1_genomic	GCF_001887475.1_ASM188747v1_genomic	GCF_001887495.1_ASM188749v1_genomic	GCF_001887515.1_ASM188751v1_genomic	GCF_001887615.1_ASM188761v1_genomic	GCF_001887635.1_ASM188763v1_genomic	GCF_001887655.1_ASM188765v1_genomic	GCF_002073335.2_ASM207333v2_genomic	GCF_002313005.1_ASM231300v1_genomic	GCF_002313025.1_ASM231302v1_genomic	GCF_002749635.1_ASM274963v1_genomic	GCF_002892855.1_ASM289285v1_genomic	GCF_002997215.1_ASM299721v1_genomic	GCF_003063785.1_ASM306378v1_genomic	GCF_008369605.1_ASM836960v1_genomic
contigs (>= 0 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
contigs (>= 1000 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
contigs (>= 5000 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
contigs (>= 10000 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
contigs (>= 25000 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
contigs (>= 50000 bp)	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
total length (>= 0 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
total length (>= 1000 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
total length (>= 5000 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
total length (>= 10000 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
total length (>= 25000 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
total length (>= 50000 bp)	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4089299
contigs	3	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	3	2	2	2	2	2	3
largest contig	3043139	2937102	3928357	4077462	3053204	2961182	2925021	2996195	2951445	2993255	2999012	2969901	2996535	2981414	3008486	2939488	2961701	2969957	3026998	3028272	2992571	3015092	2975504	2948589	
total length	4258023	4061813	3928357	4077462	5000494	4033501	4050927	4102038	4110440	4057520	4099119	4062508	4110872	4170245	4076583	4014863	4066727	4042269	4264715	4095733	4063073	4085463	4085460	4047835	4138412
GC (%)	47.75	47.62	47.77	47.69	45.67	47.49	47.61	47.47	47.53	47.51	47.47	47.56	47.48	47.58	47.47	47.58	47.60	47.49	47.66	47.74	47.50	47.46	47.46	47.50	47.42
50	3043139	2937102	3928357	4077462	3053204	2961182	2925021	2996195	2951445	2993255	2999012	2969901	2996535	2981414	3008486	2939488	2961701	2969957	3026998	3028272	2992571	3015092	2975504	2948589	
90	1047617	1124711	3928357	4077462	896006	1072319	1125906	1105843	1158995	1064265	1100107	1092607	1114337	1188831	1068097	1075375	1105026	1072312	1057601	1067461	1070502	1070371	1072331	1140710	
auN	2439201.2	2435253.2	3928357.0	4077462.0	2245794.0	2459021.8	2424978.8	2486586.7	2446039.1	2487291.6	2489390.6	2465004.9	2486326.2	2470394.1	2500088.7	2440188.1	2457198.4	2466559.6	2418374.6	2517230.6	2486161.5	2505584.8	2505586.0	2471325.5	2415856.2
50	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	
90	2	2	1	1	3	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	
N's per 100 kbp	0.00	0.00	0.00	0.00	0.00	0.30	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.30	0.00	0.00	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).



5541v1_genomic

GCF_001887435.1_ASM188743v1_genomic

GCF_00207

3871v2_genomic

GCF_001887455.1_ASM188745v1_genomic

GCF_00231

5923v1_genomic

GCF_001887475.1_ASM188747v1_genomic

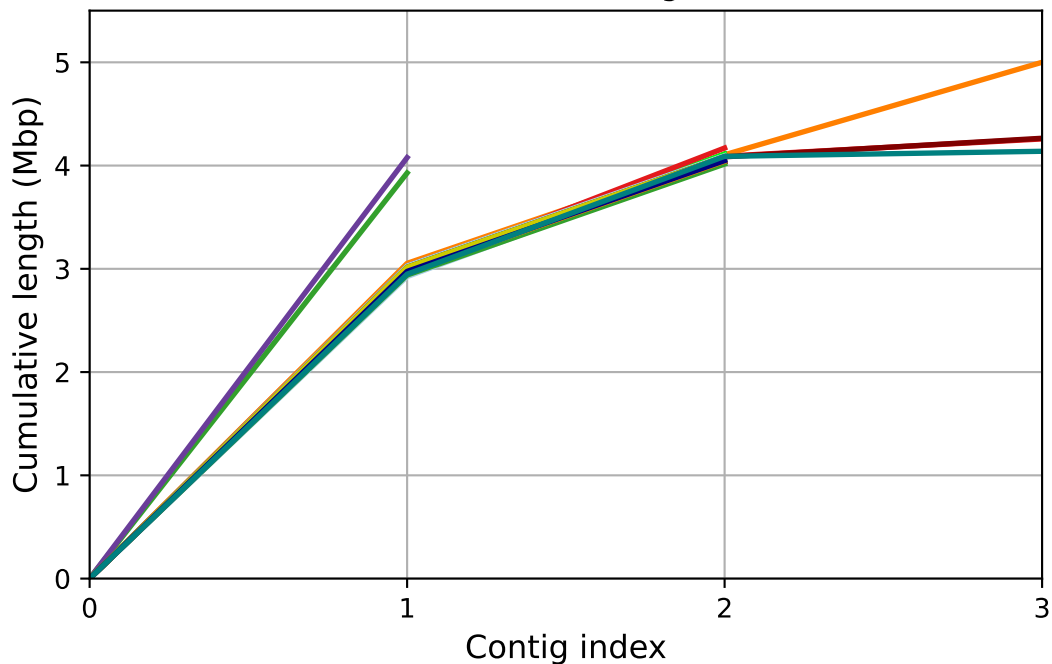
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5926v1_genomic

GCF_001887495.1_ASM188749v1_genomic

GCF_00274

Cumulative length



5541v1_genomic

GCF_001887435.1_ASM188743v1_genomic

GCF_00207

8871v2_genomic

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GCF_00231

5923v1_genomic

GCF_001887475.1_ASM188747v1_genomic

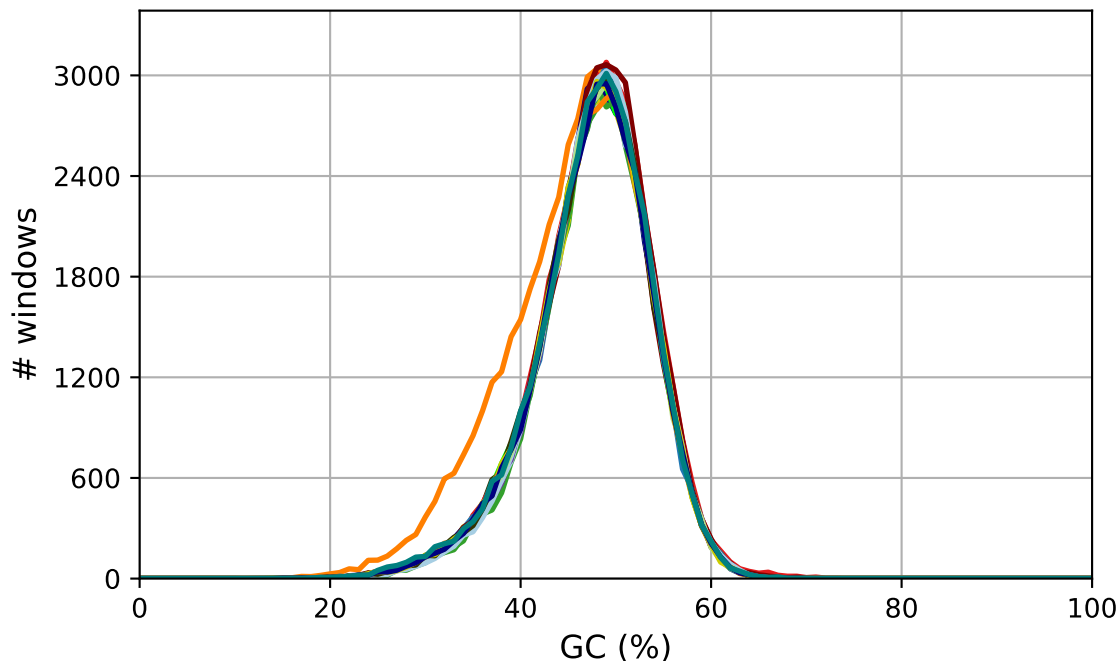
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5926v1_genomic

GCF_001887495.1_ASM188749v1_genomic

GCF_00274

GC content



5541v1_genomic

8871v2_genomic

5923v1_genomic

5926v1_genomic

GCF_001887435.1_ASM188743v1_genomic

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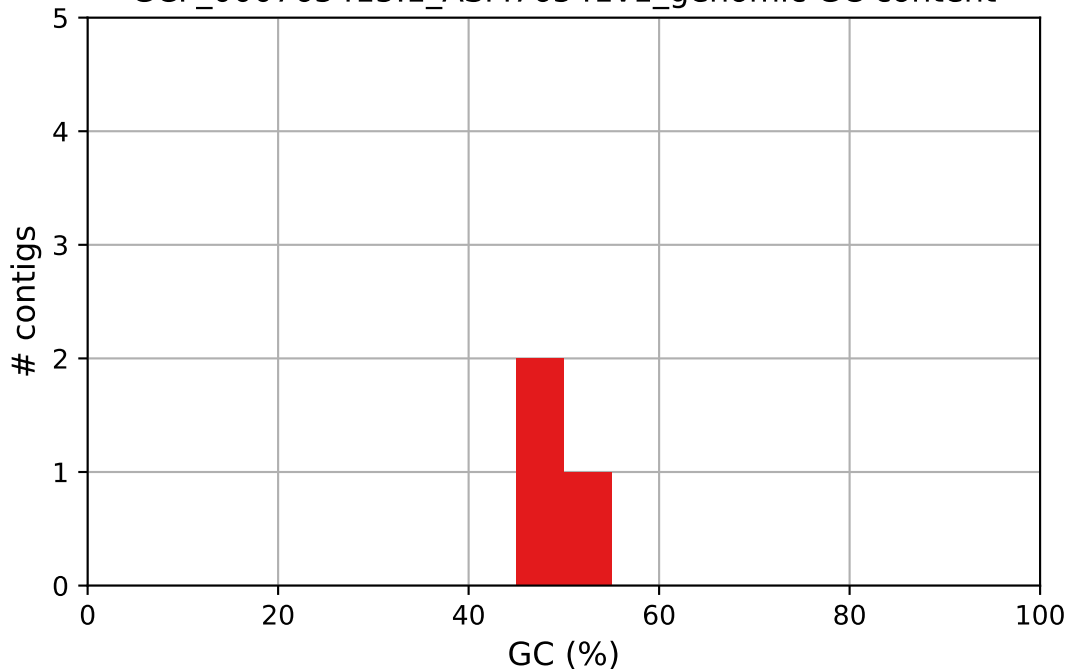
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GCF_00231

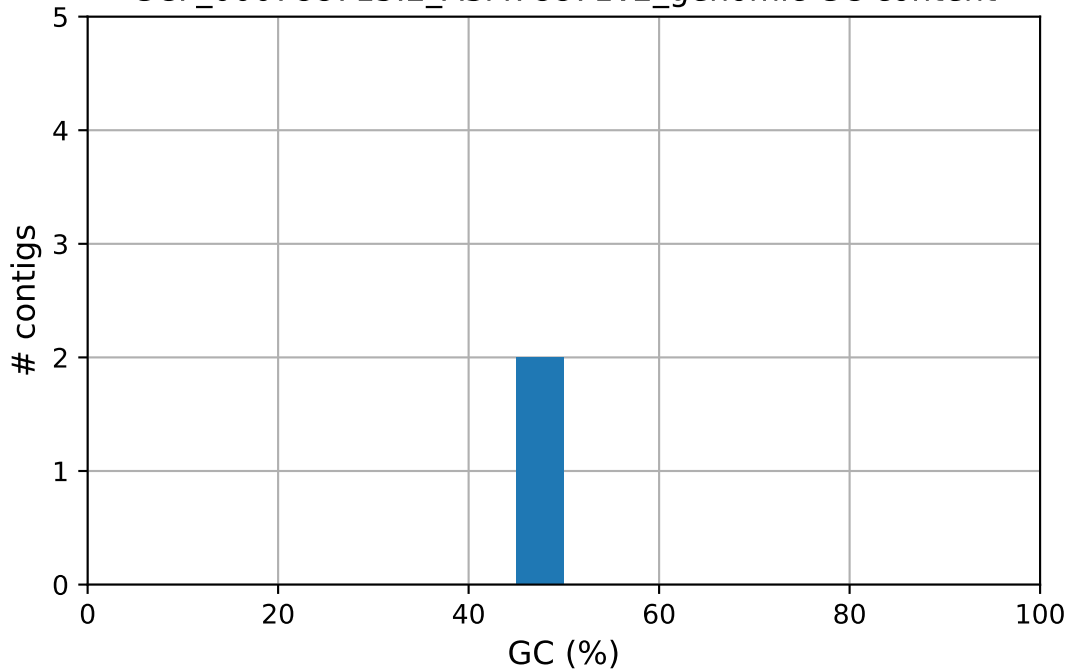
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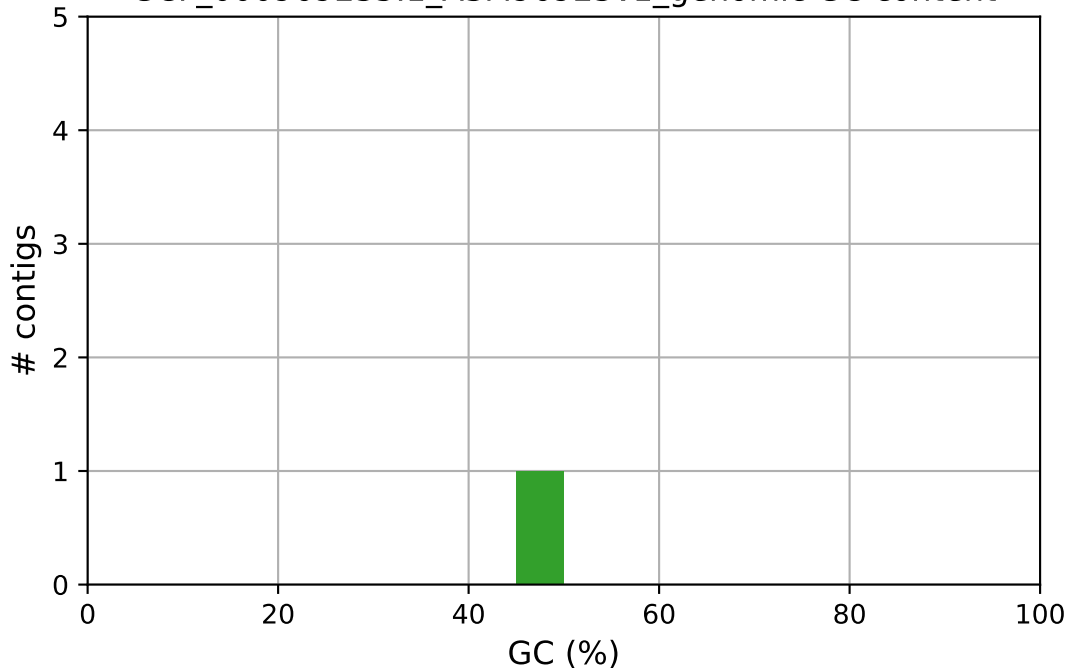
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GCF_000788715.2_ASM78871v2_genomic GC content



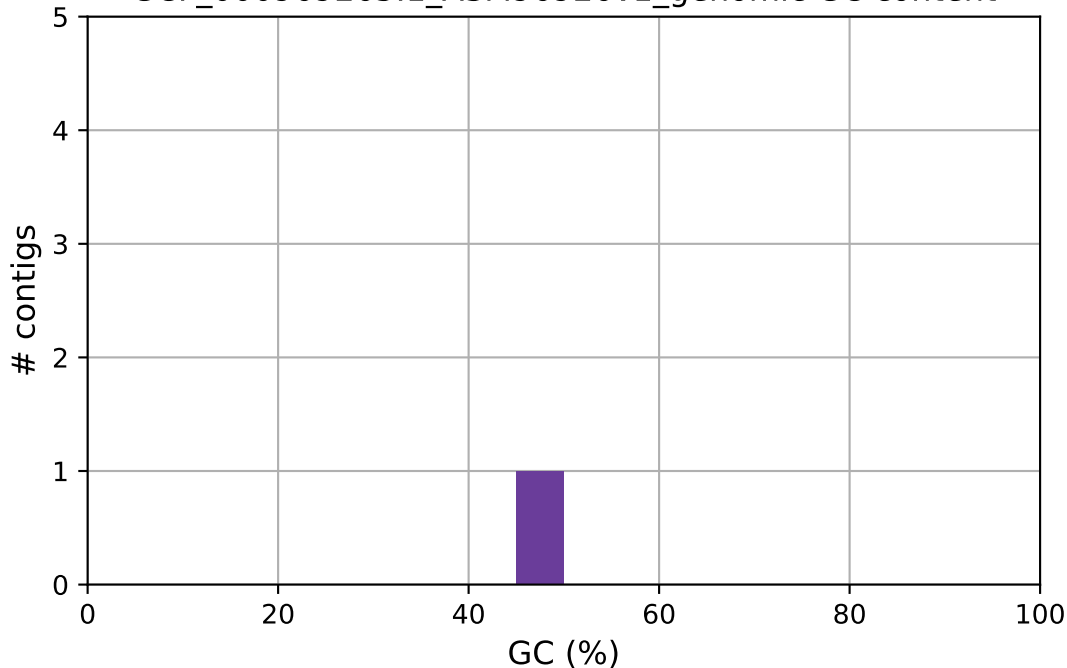
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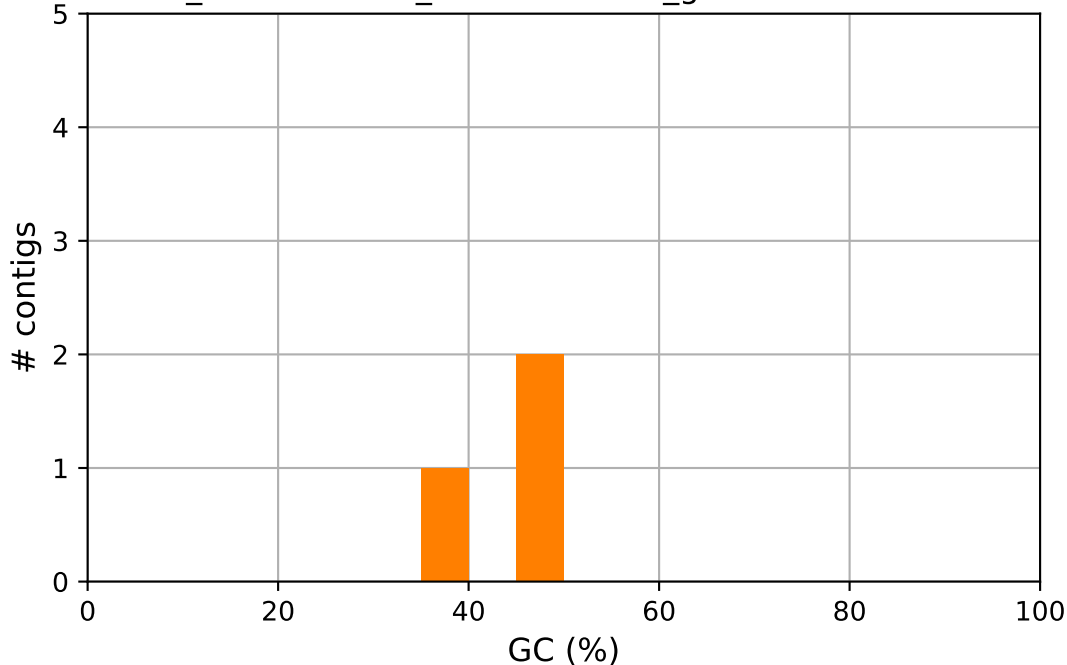
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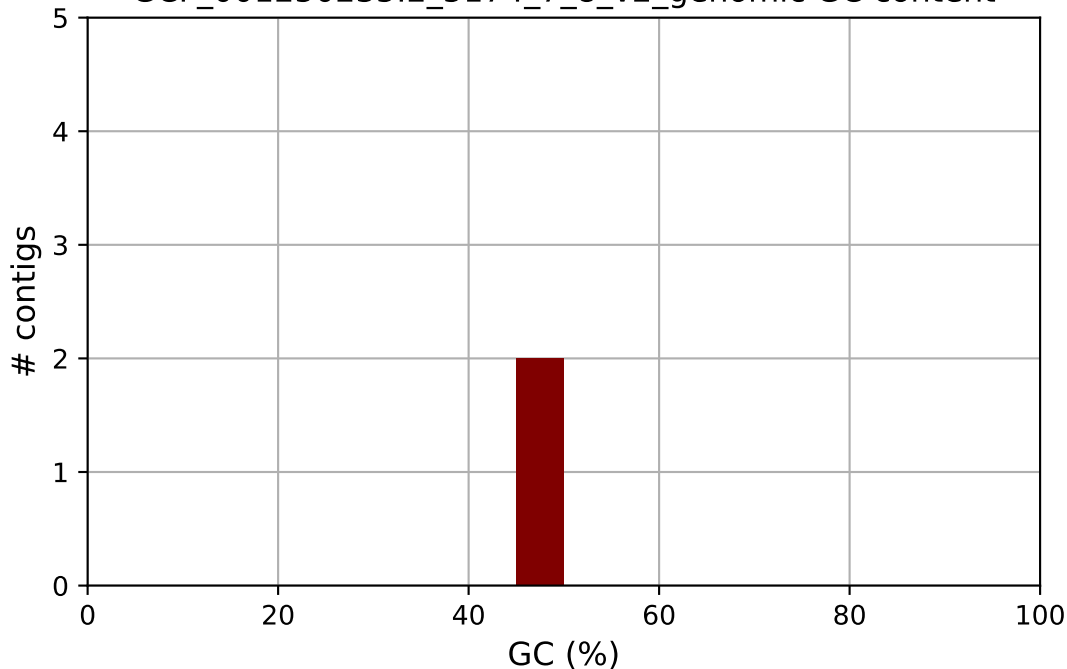
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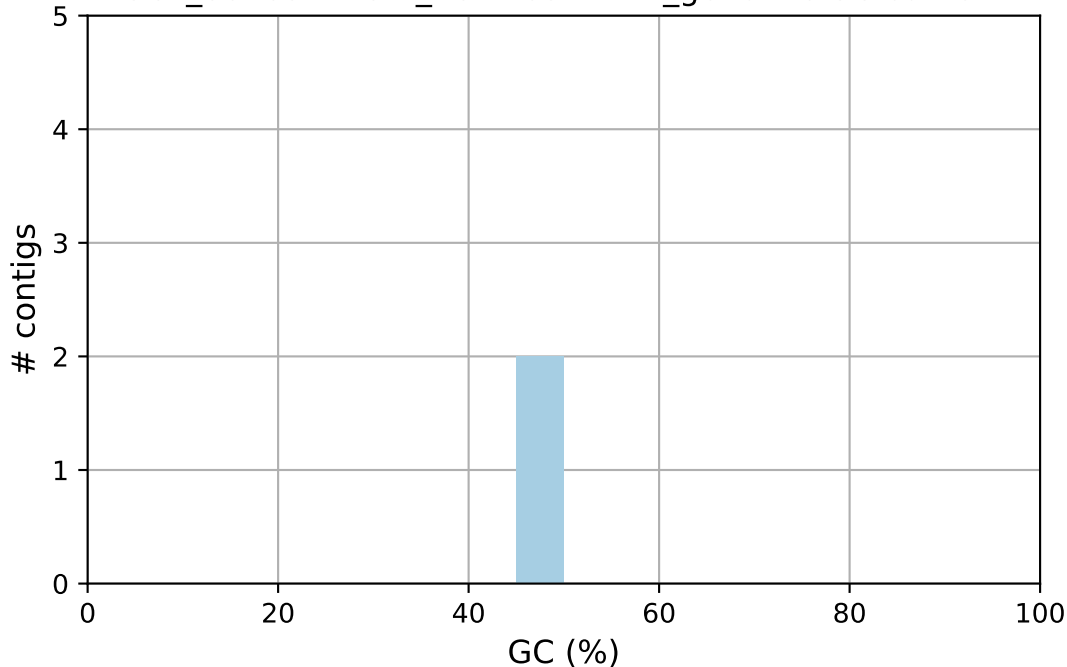
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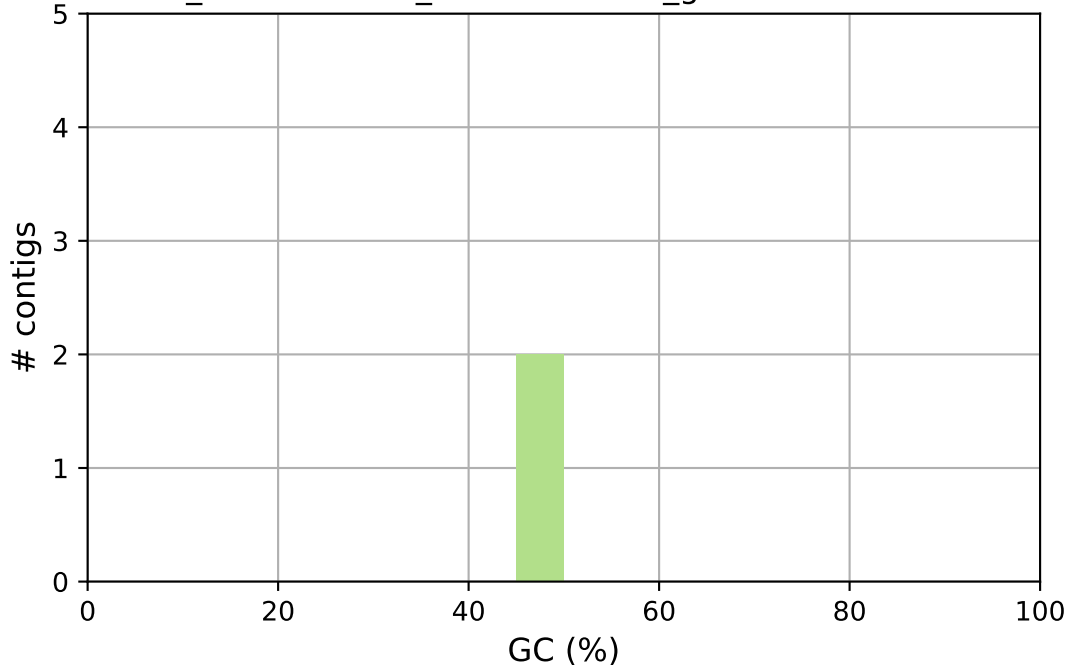
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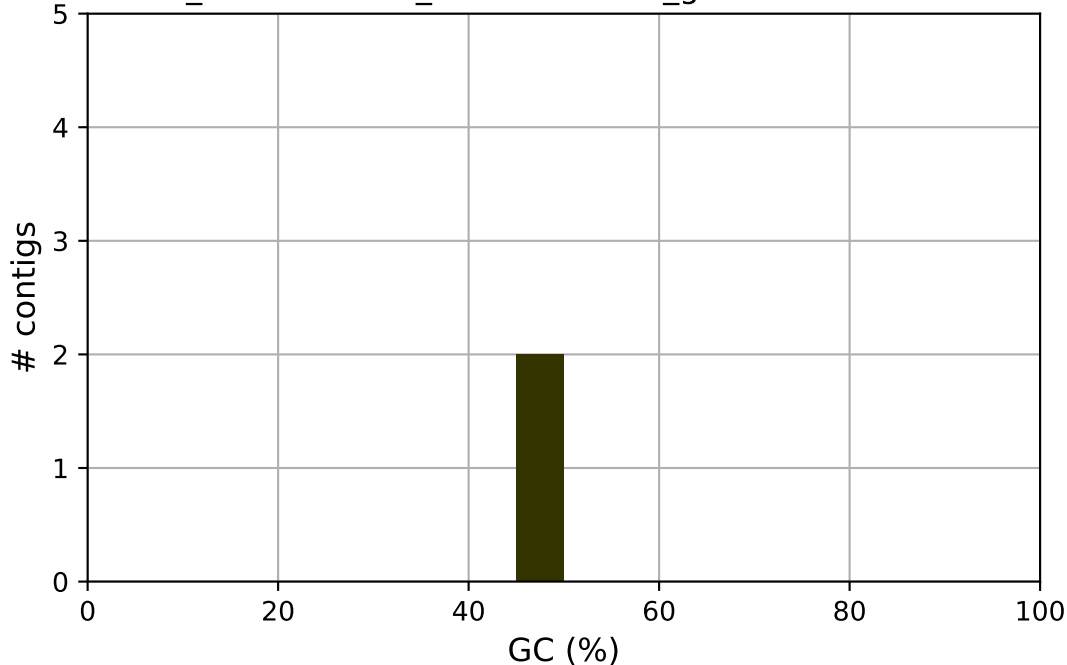
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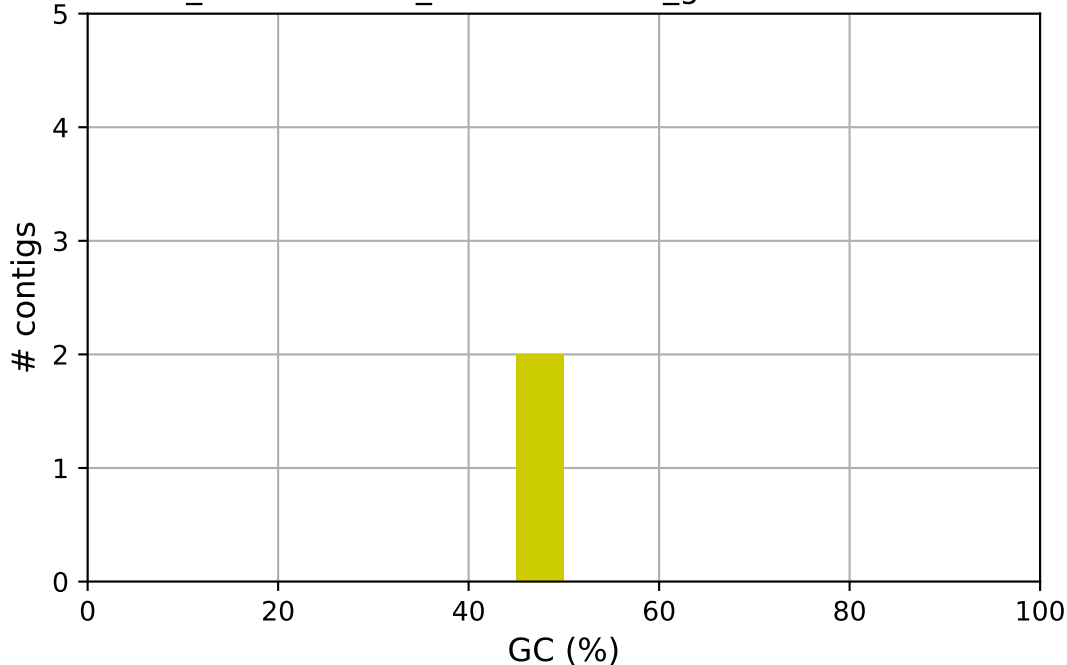
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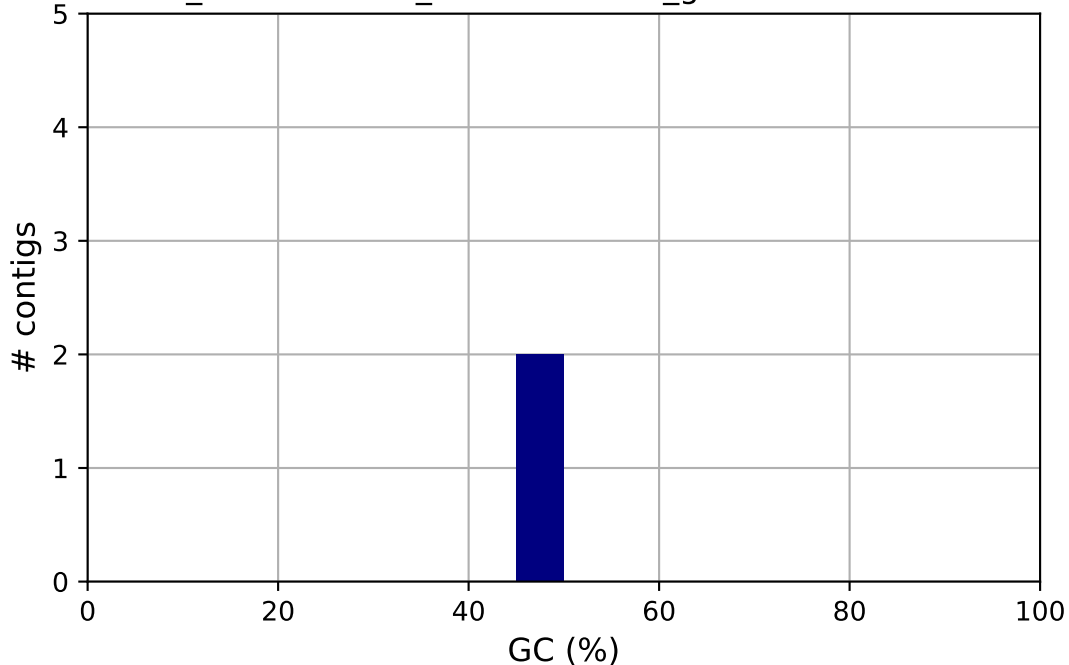
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GCF_001887435.1_ASM188743v1_genomic GC content



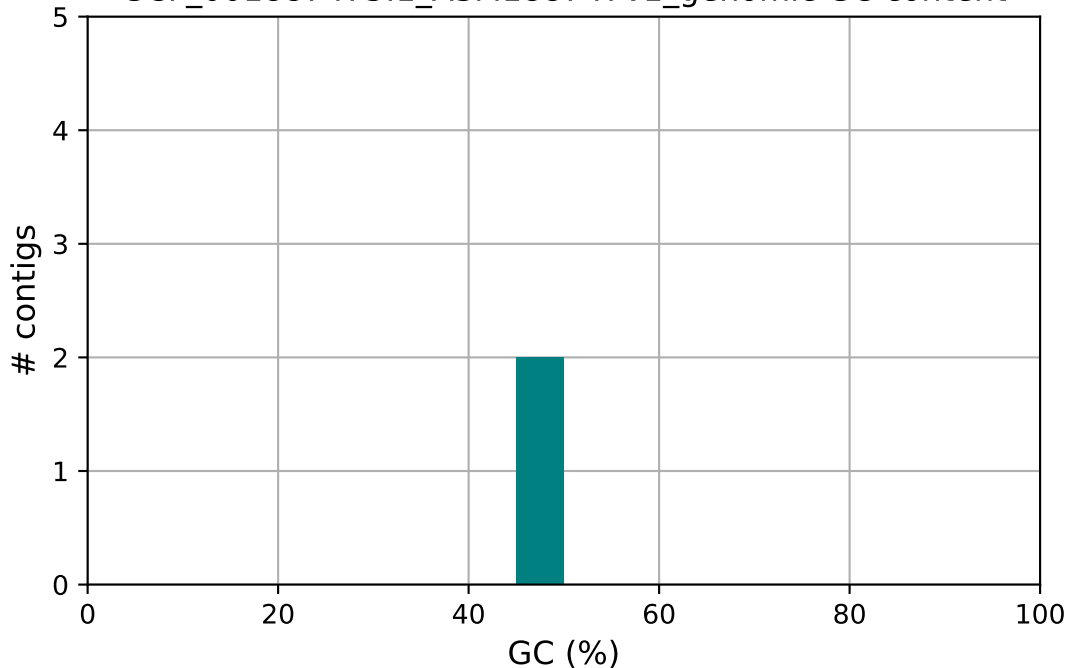
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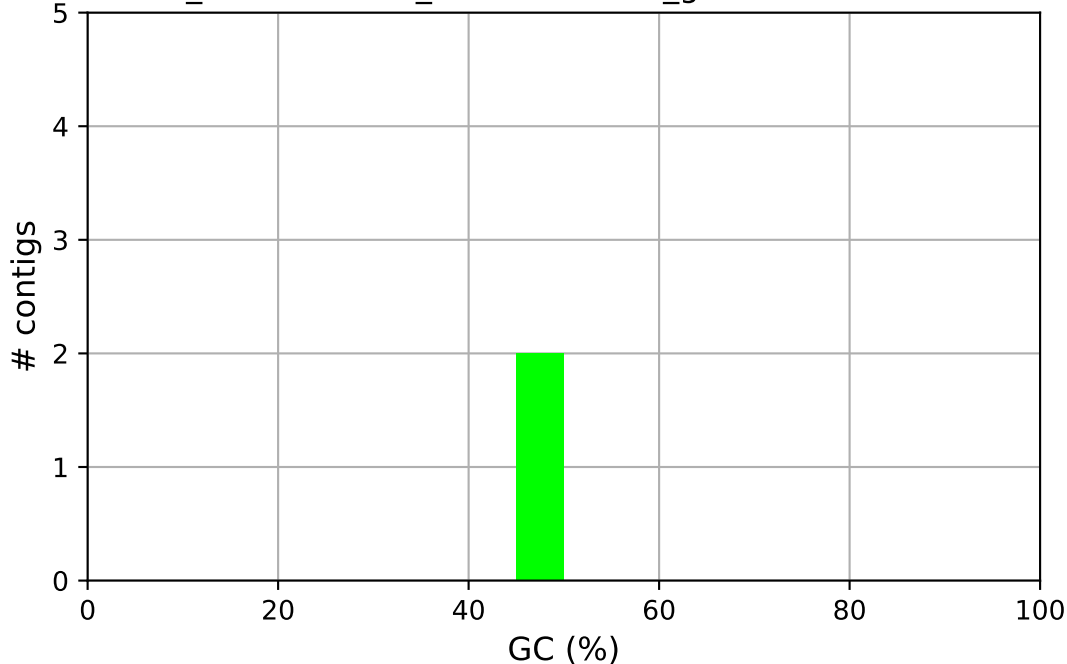
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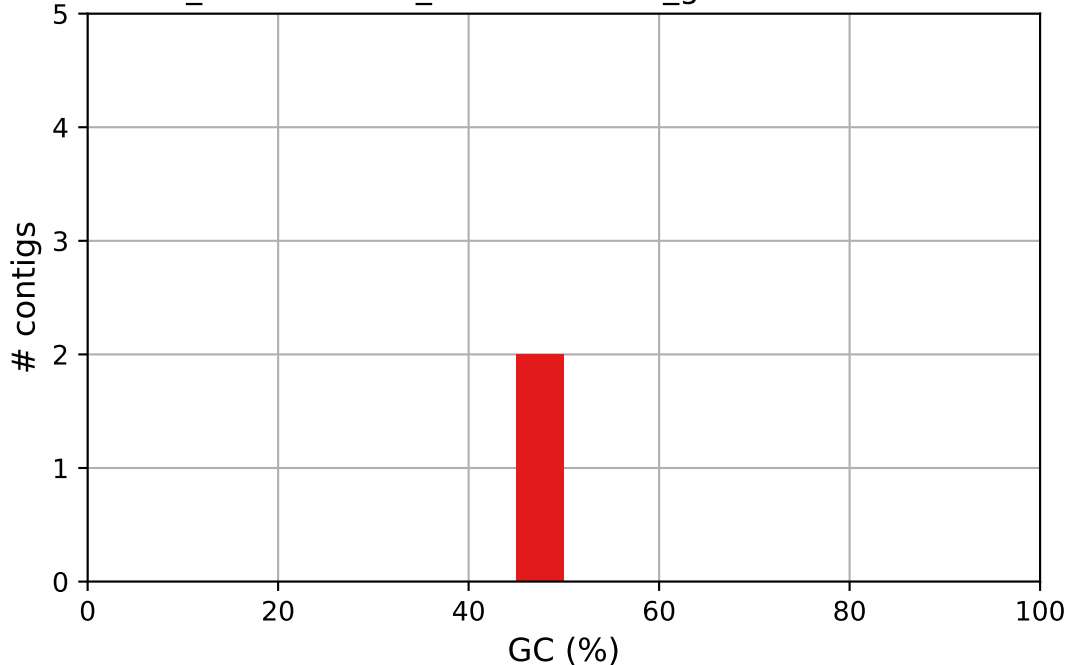
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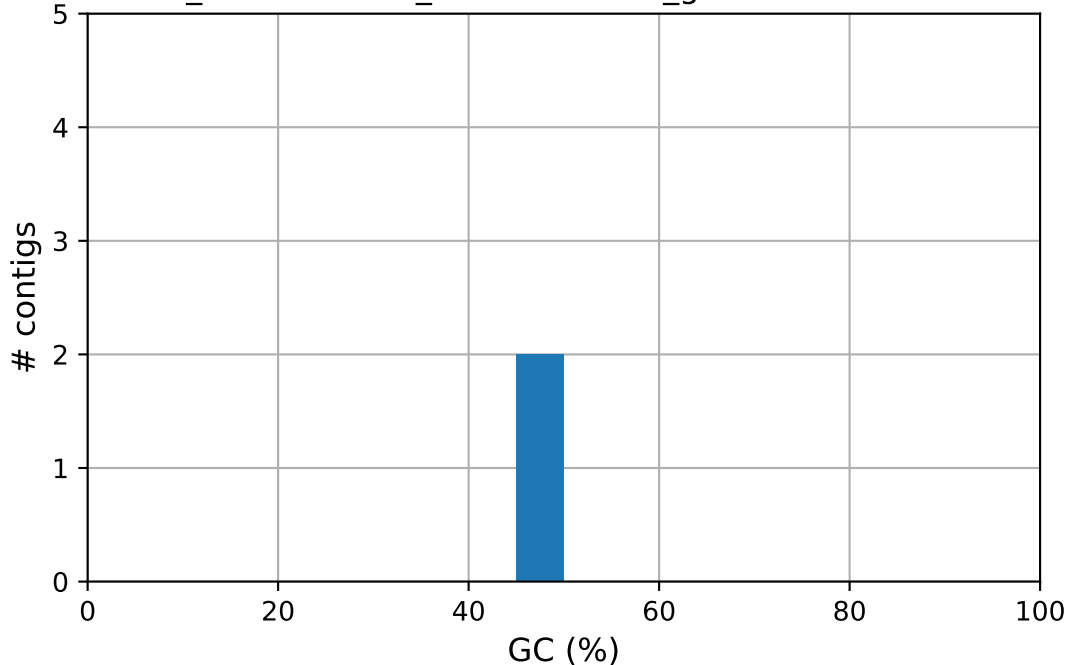
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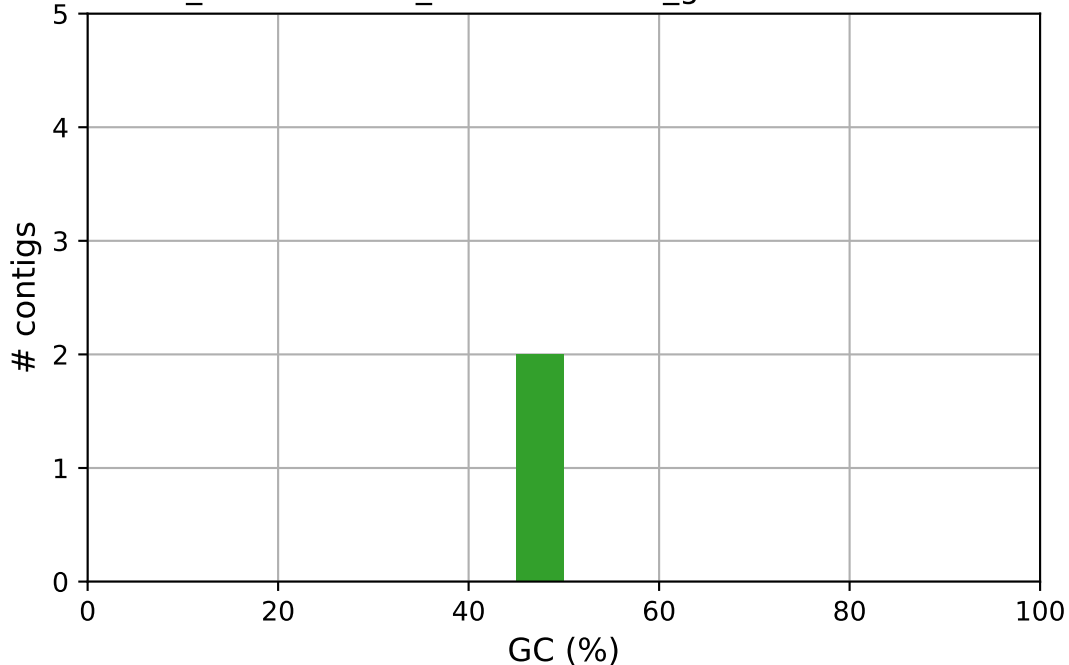
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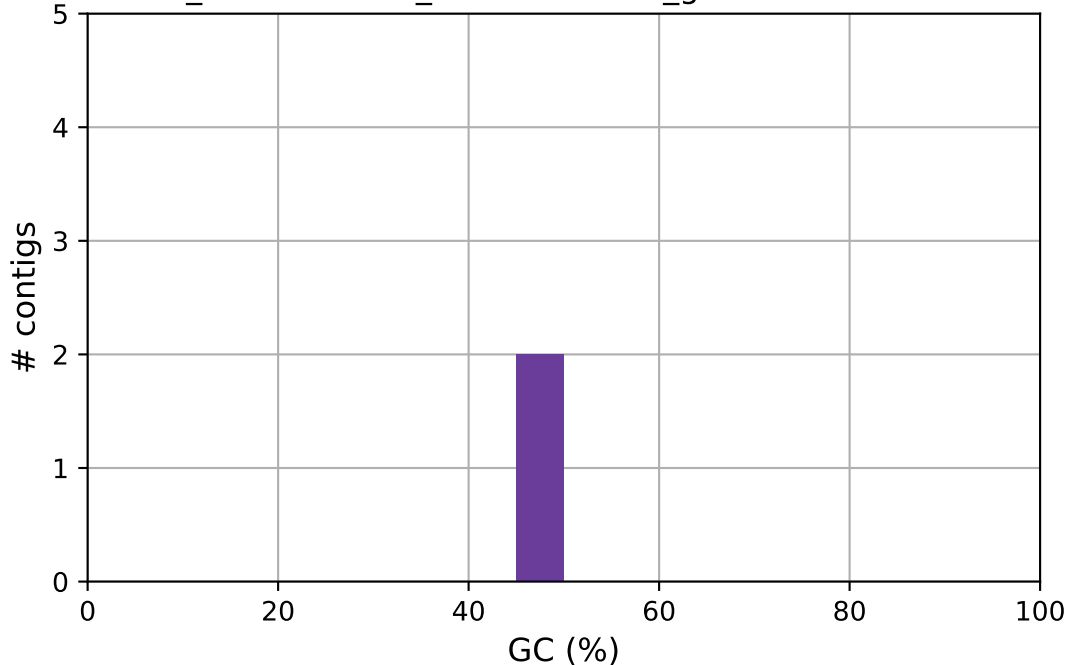
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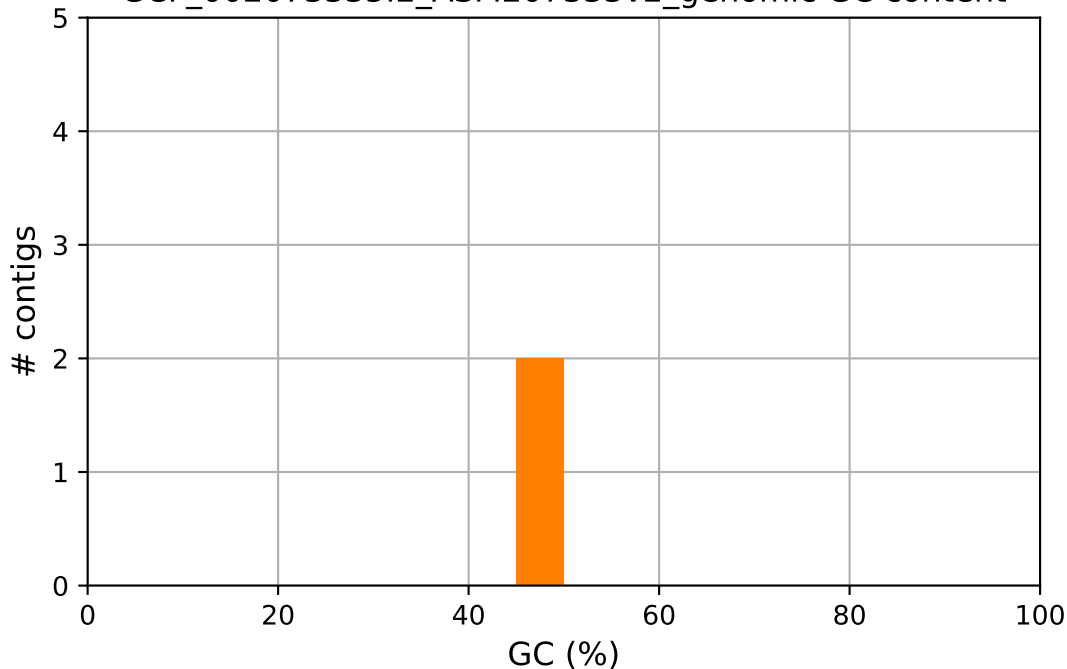
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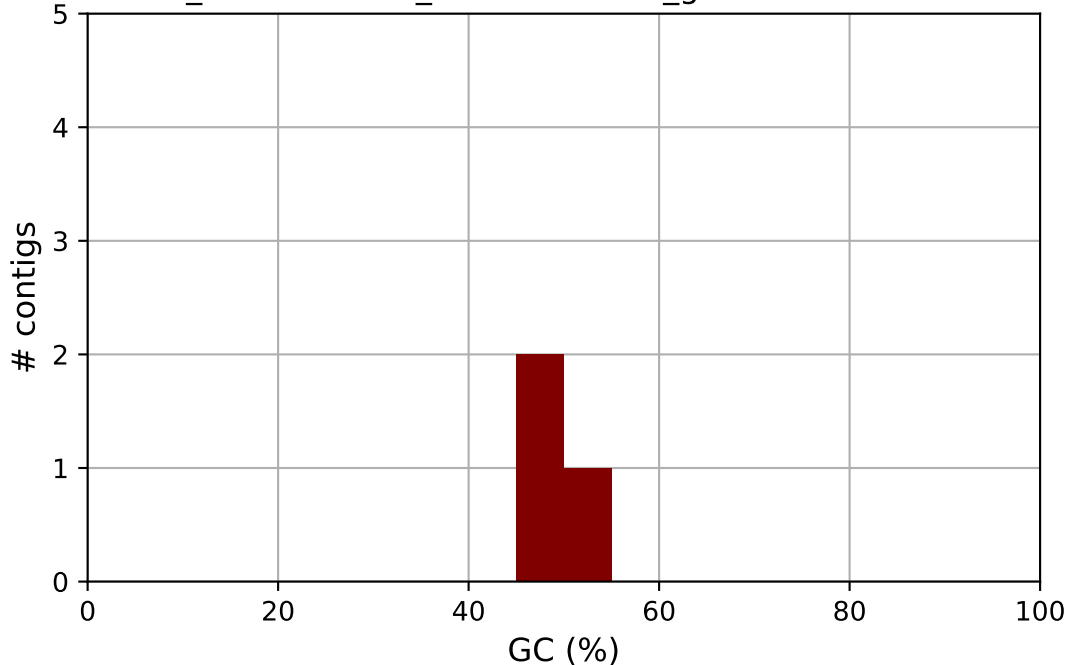
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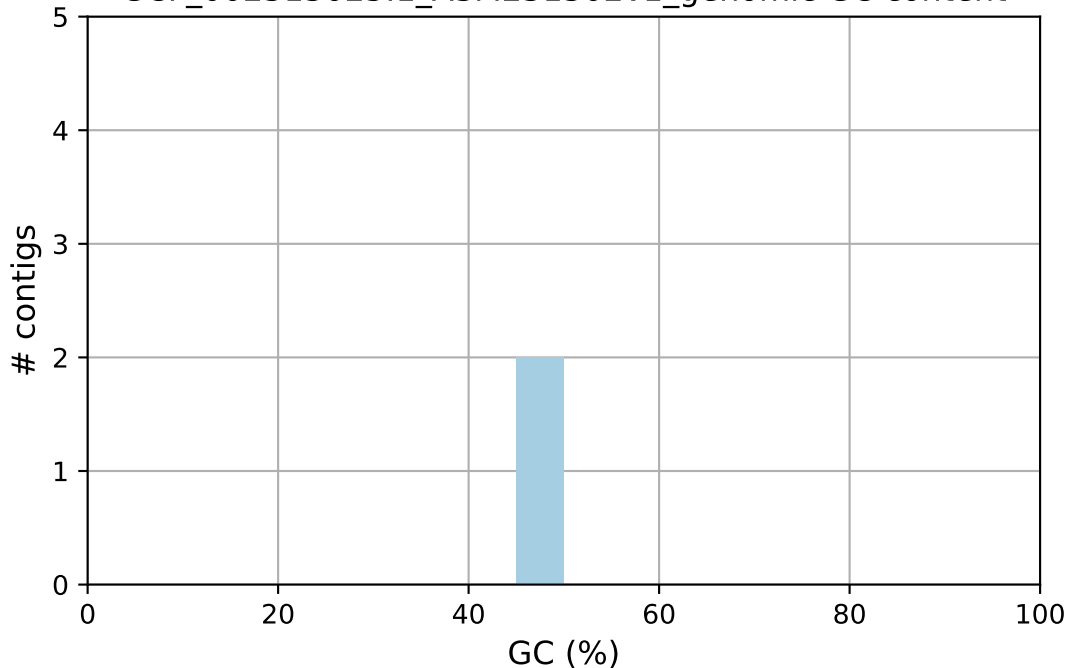
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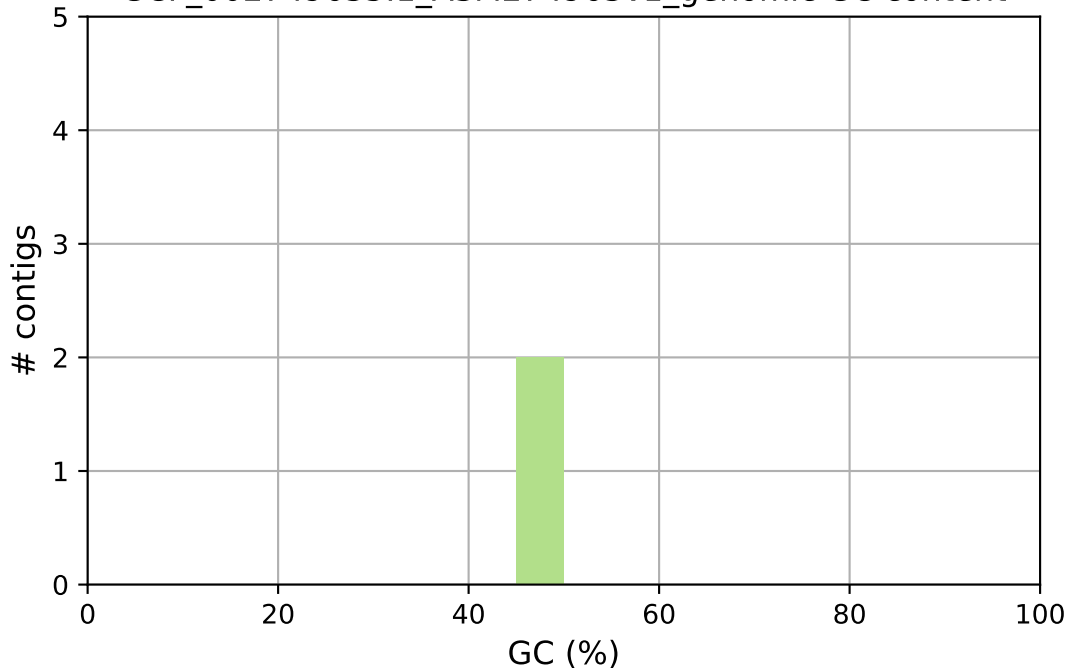
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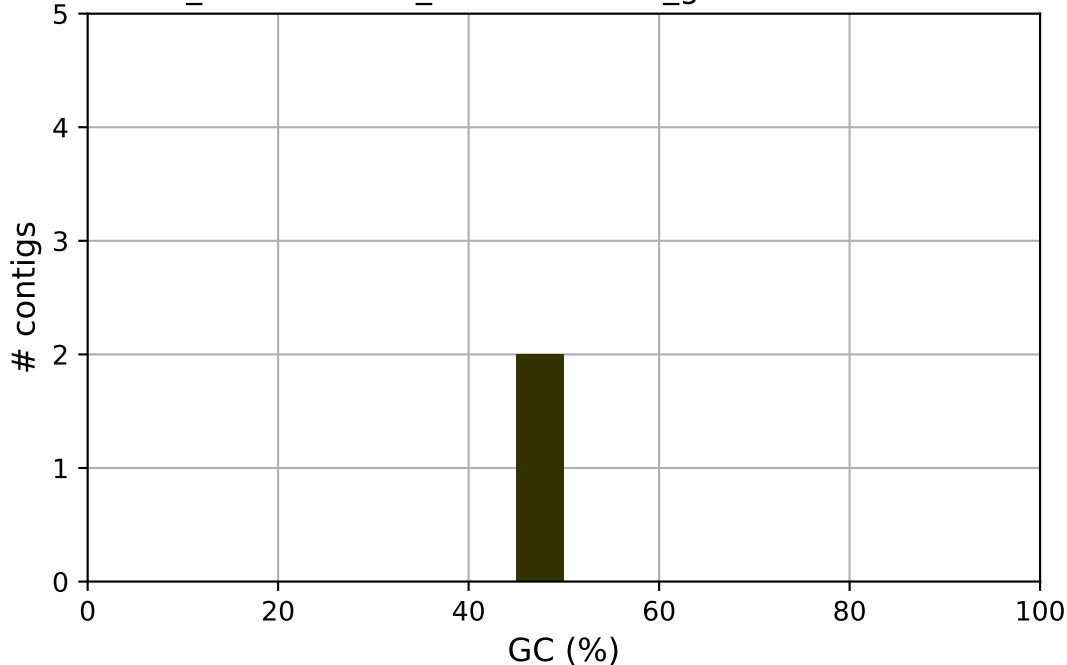
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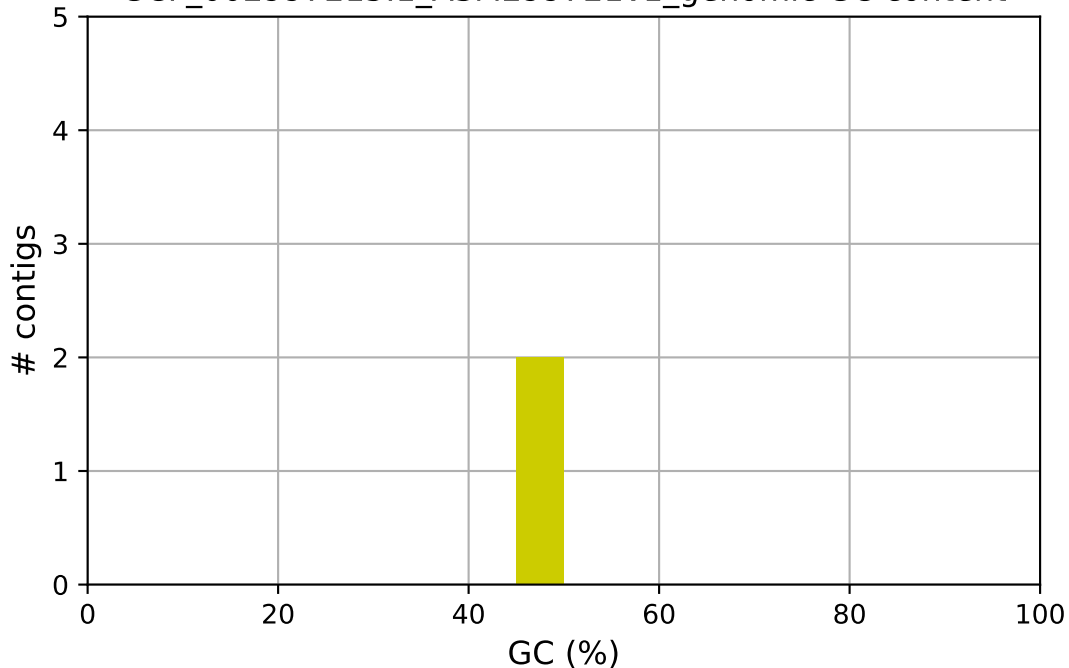
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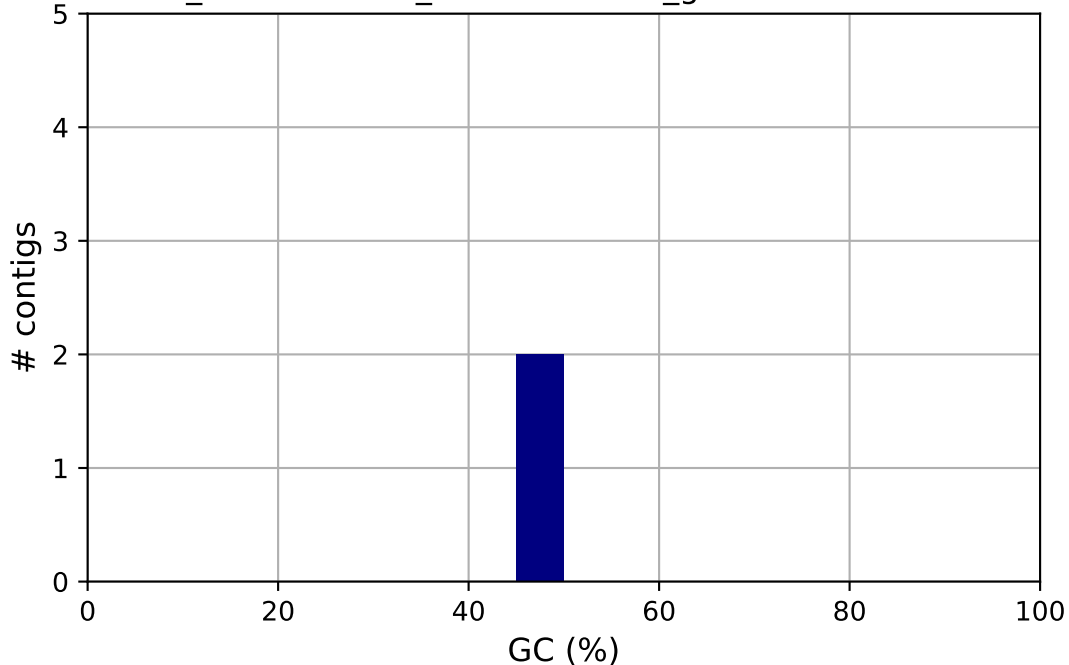
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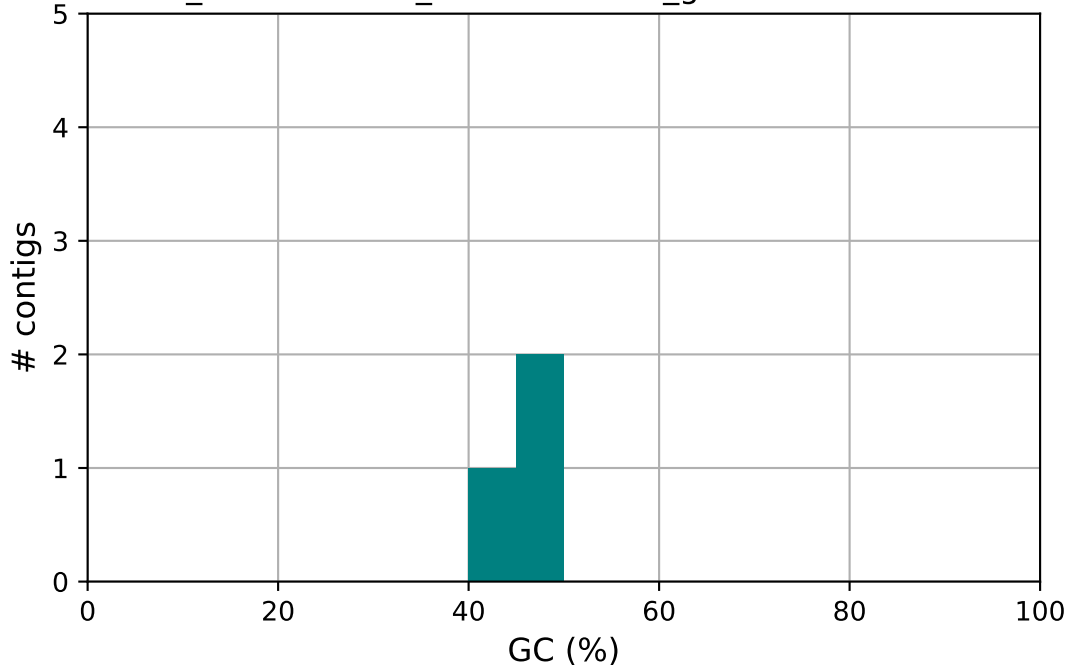
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GCF_003063785.1_ASM306378v1_genomic

GCF_008369605.1_ASM836960v1_genomic GC content



GCF_008369605.1_ASM836960v1_genomic