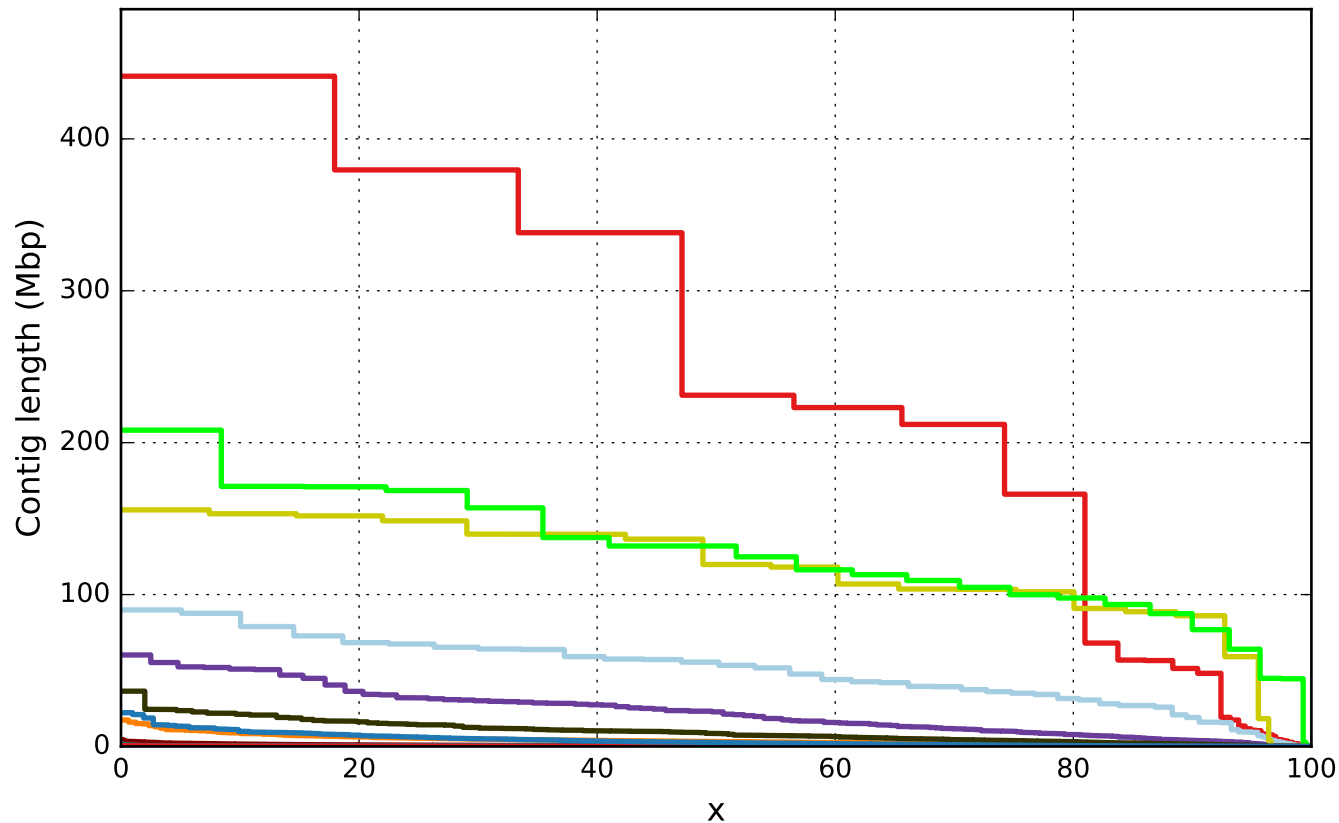


Report

	mSorAra1.pri.asm.20211103	GCA_000181395.1_ASM18139v1_genomic	GCA_004027635.1_Crolnd_v1_BIUU_genomic	GCF_000181275.1_SorAra2.0_genomic	GCA_000296755.1_EriEur2.0_genomic	GCA_004363575.1_SolPar_v1_BIUU_genomic	GCF_000260355.1_ConCri1.0_genomic	GCA_002901085.1_ASM290108v1_genomic	GCA_019455555.1_Gpyr_1.0_genomic	GCF_014898055.1_MPIMG_talOcc4_genomic	GCA_004024925.1_ScaAqu_v1_BIUU_genomic	GCA_021461705.1_Cryptotis_parva_assembly_1.0_genomic	GCF_024139225.1_mSunEtr1.pri.cur_genomic	GCA_004024945.1_UroGra_v1_BIUU_genomic	GCA_026122425.1_SC2_V1_genomic
# contigs (>= 0 bp)	233	847680	2415412	12845	5802	176618	2040	40482	12306	4425	640296	39388	147	723835	3379
# contigs (>= 1000 bp)	233	771929	533455	12845	5802	22577	2040	40482	3275	4425	55714	38273	147	103814	3379
# contigs (>= 5000 bp)	233	75731	120290	2172	2559	14072	467	31912	1469	3685	30305	34459	142	47328	3379
# contigs (>= 10000 bp)	232	18699	54248	1078	2045	12022	266	27716	1117	2681	25869	30863	130	35104	3379
# contigs (>= 25000 bp)	213	507	11136	585	1770	9115	189	20179	848	944	18927	22020	108	20757	3379
# contigs (>= 50000 bp)	164	63	1505	473	1663	6726	144	13180	680	92	12364	13069	70	10908	3379
Total length (>= 0 bp)	2459336321	2133117389	3082217326	2423158183	2715703478	2109877870	1769662895	2049415269	1828347170	2098020470	2068871057	2196365350	2471666657	2112710047	2283226199
Total length (>= 1000 bp)	2459336321	2066879886	2402584627	2423158183	2715703478	2062807578	1769662895	2049415269	1822166123	2098020470	1901288684	2195647555	2471666657	1914657881	2283226199
Total length (>= 5000 bp)	2459336321	664873857	1532259342	2404465319	2709454854	2044701388	1766318581	2028343025	1817972561	2095917748	1849728396	2184794356	2471650960	1792905057	2283226199
Total length (>= 10000 bp)	2459326403	259905613	1070390752	2396748103	2705583874	2029881143	1764918546	1997595767	1815584124	2088709202	1817877759	2158359884	2471550903	1705763405	2283226199
Total length (>= 25000 bp)	2458942042	19472995	417167120	2389897034	2701480748	1981673732	1763844726	1871204344	1811265490	2059841332	1701908320	2005110559	2471160777	1471659153	2283226199
Total length (>= 50000 bp)	2457086062	5414065	98299390	2385054282	2697629261	1895673329	1762011911	1616982459	1805352079	2032050082	1463336420	1685520253	2469745763	1118879868	2283226199
# contigs	233	160314	199640	3663	3186	34588	827	34782	1874	4024	36165	144	59453	3379	
Largest contig	441239200	392637	334926	60238417	17429815	4518367	89887394	919807	36404611	155735125	758314	1631536	208244741	675744	22233027
Total length	2459336321	971098796	1837639356	2410424072	2711841657	2051806763	1767713640	2038902213	1819516938	2097278780	1867122067	2191539156	2471658457	1839970050	2283226199
GC (%)	43.23	41.13	38.76	42.91	41.53	39.87	41.54	39.87	41.65	41.91	41.64	42.79	39.81	39.99	41.93
N50	231235302	7153	12307	22794405	3267260	425252	55520359	111636	8555737	119794413	106858	112495	131952995	67231	2625934
N75	166085329	4315	6391	9970888	1649127	177767	35080050	58362	3836593	103423330	56035	53654	99890541	31321	874277
L50	4	45685	40587	36	248	1231	13	5281	65	8	5184	5192	8	7477	209
L75	7	89382	92965	78	541	3099	24	11556	146	12	11177	12261	14	17485	586
# N's per 100 kbp	2392.21	0.00	28.66	9584.55	14104.85	25.28	4923.26	13.07	1495.92	17.36	38.77	0.00	430.64	44.83	5404.91

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).

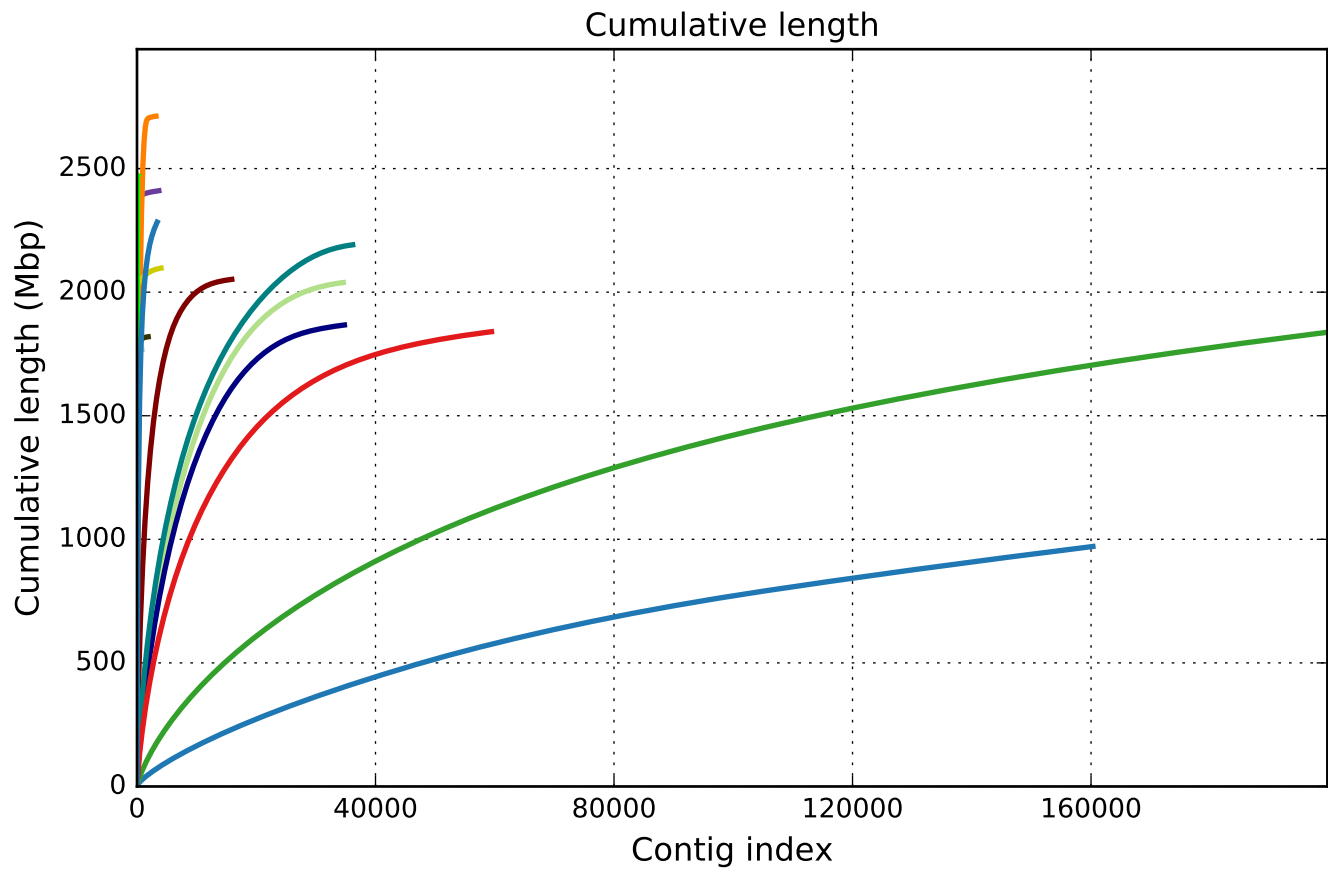
Nx



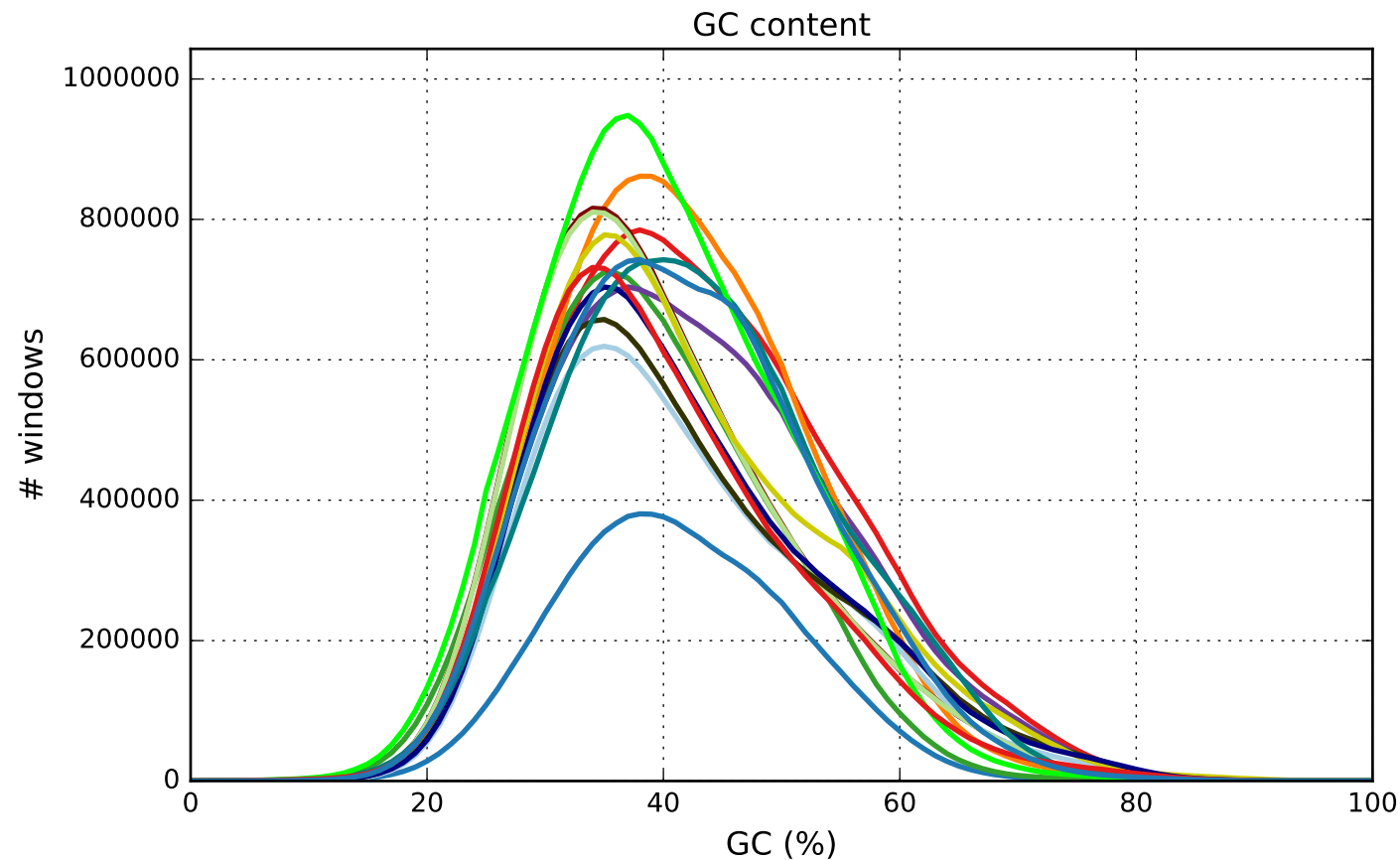
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— GCA_002901085.1_ASM290108v1_genomic
— GCA_019455555.1_Gpyr_1.0_genomic

— GCA_004024925.1_s
— GCA_021461705.1_c
— GCF_024139225.1_r
— GCA_004024945.1_s



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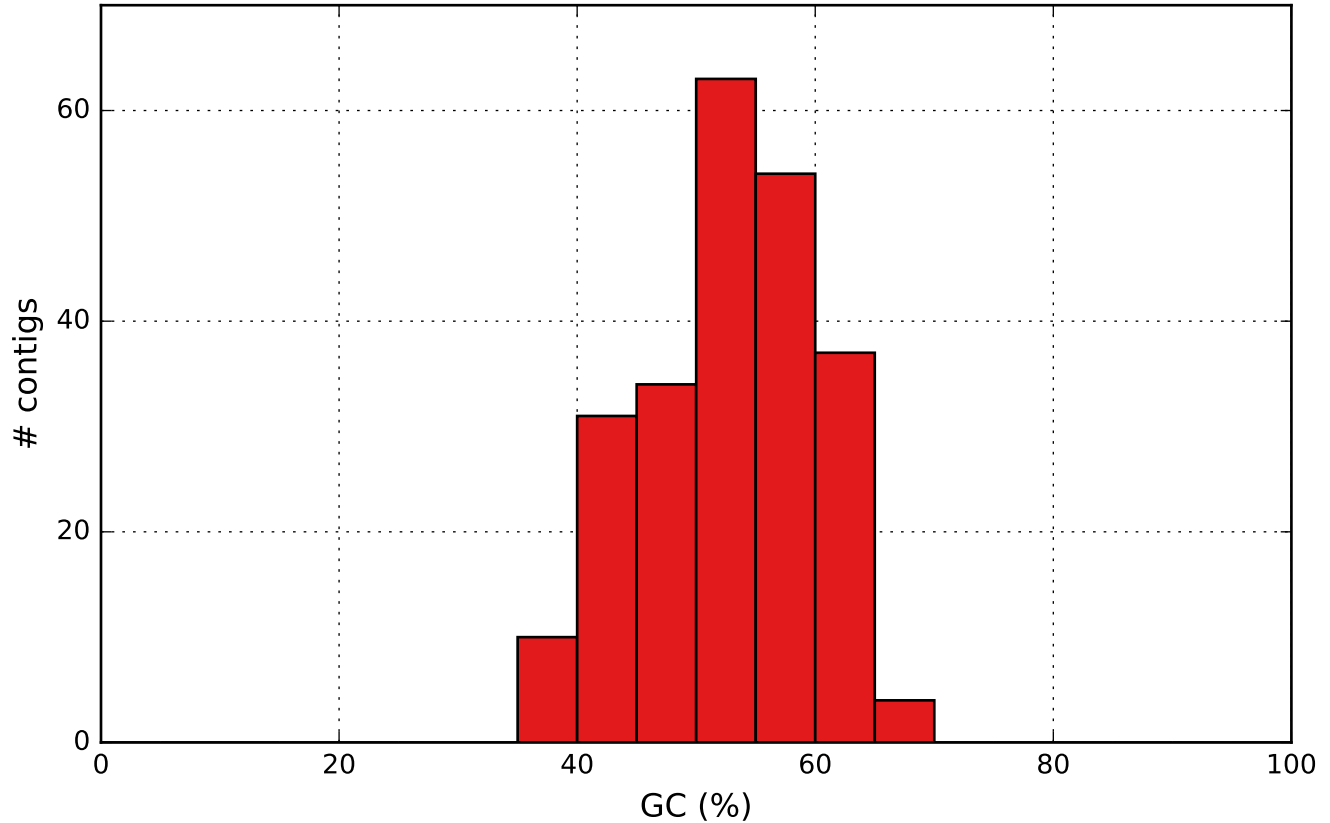


_genomic
UU_genomic
omic

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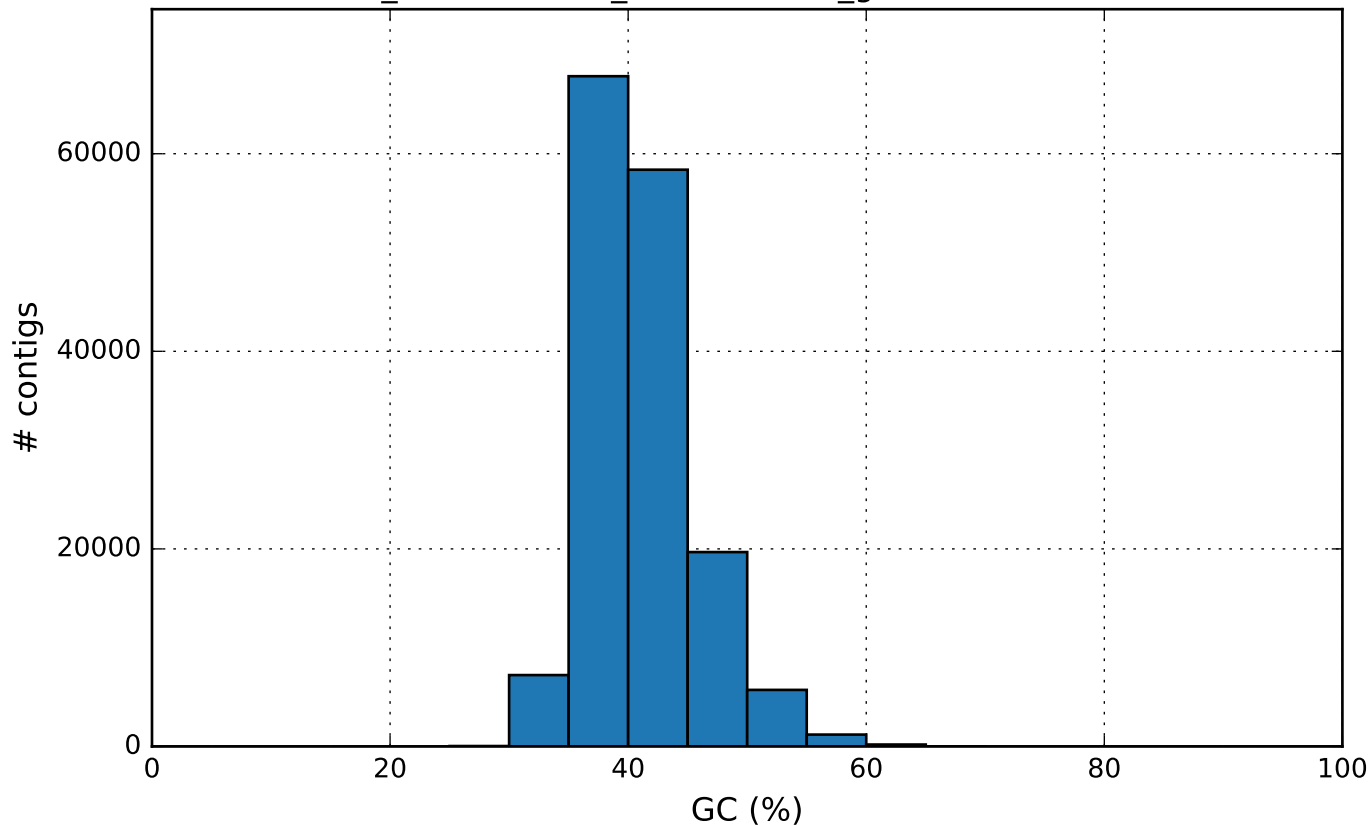
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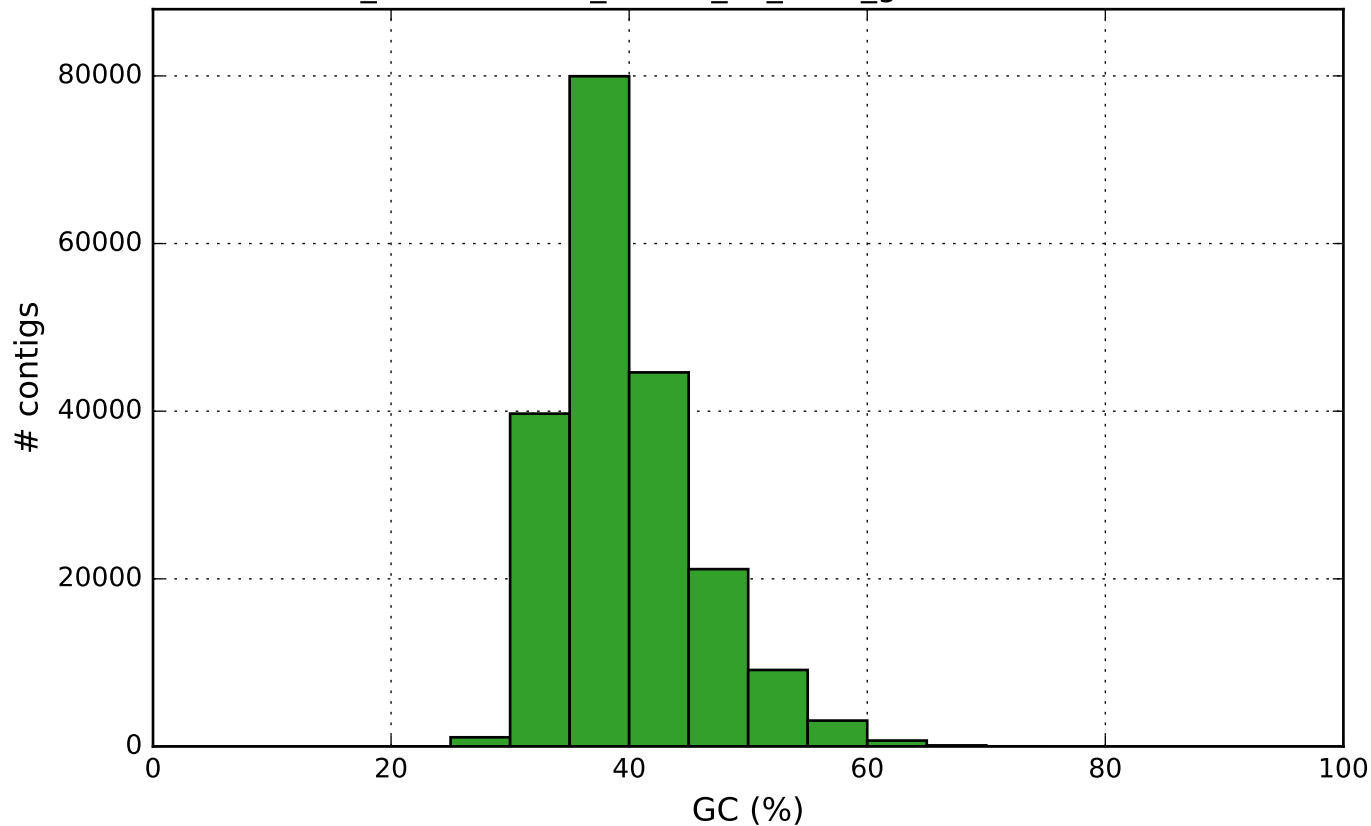
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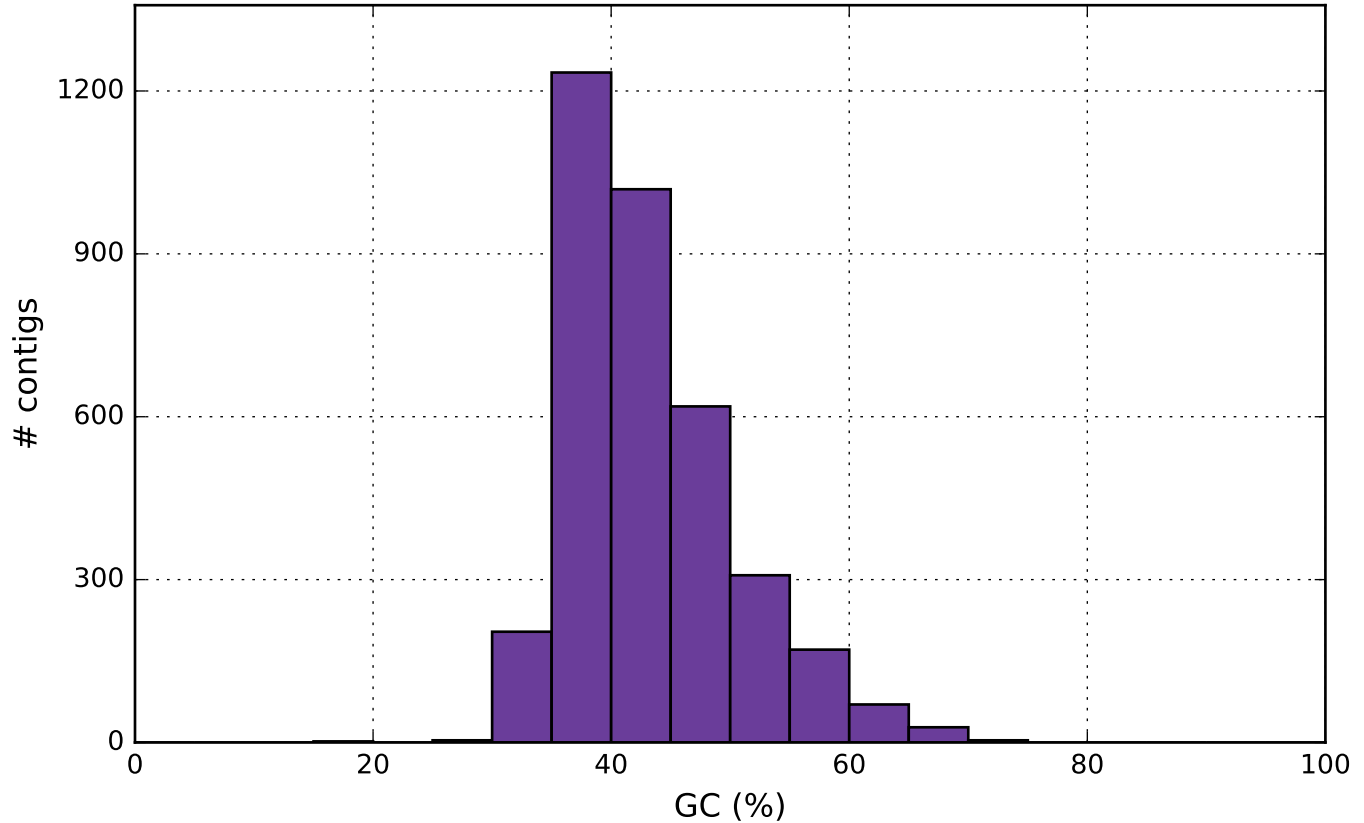
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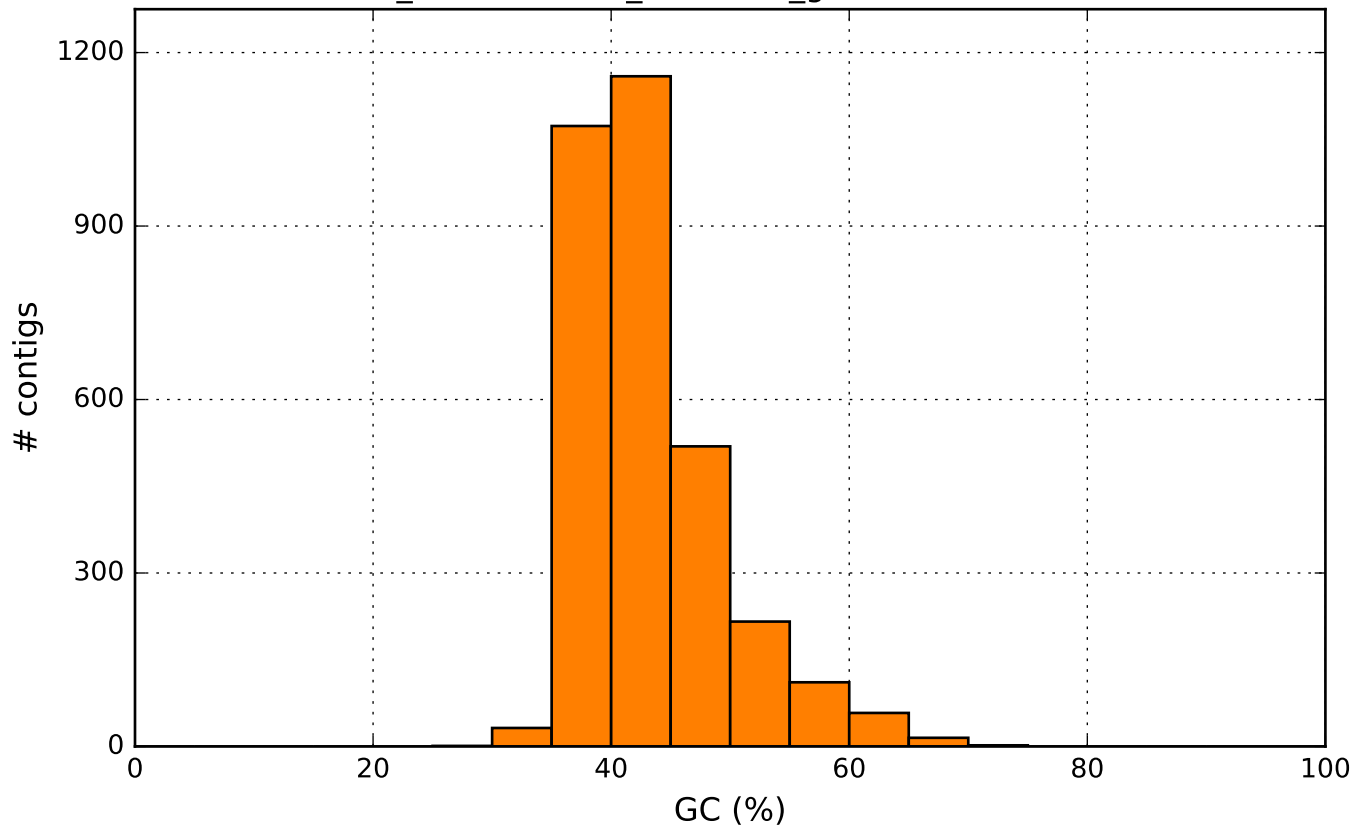
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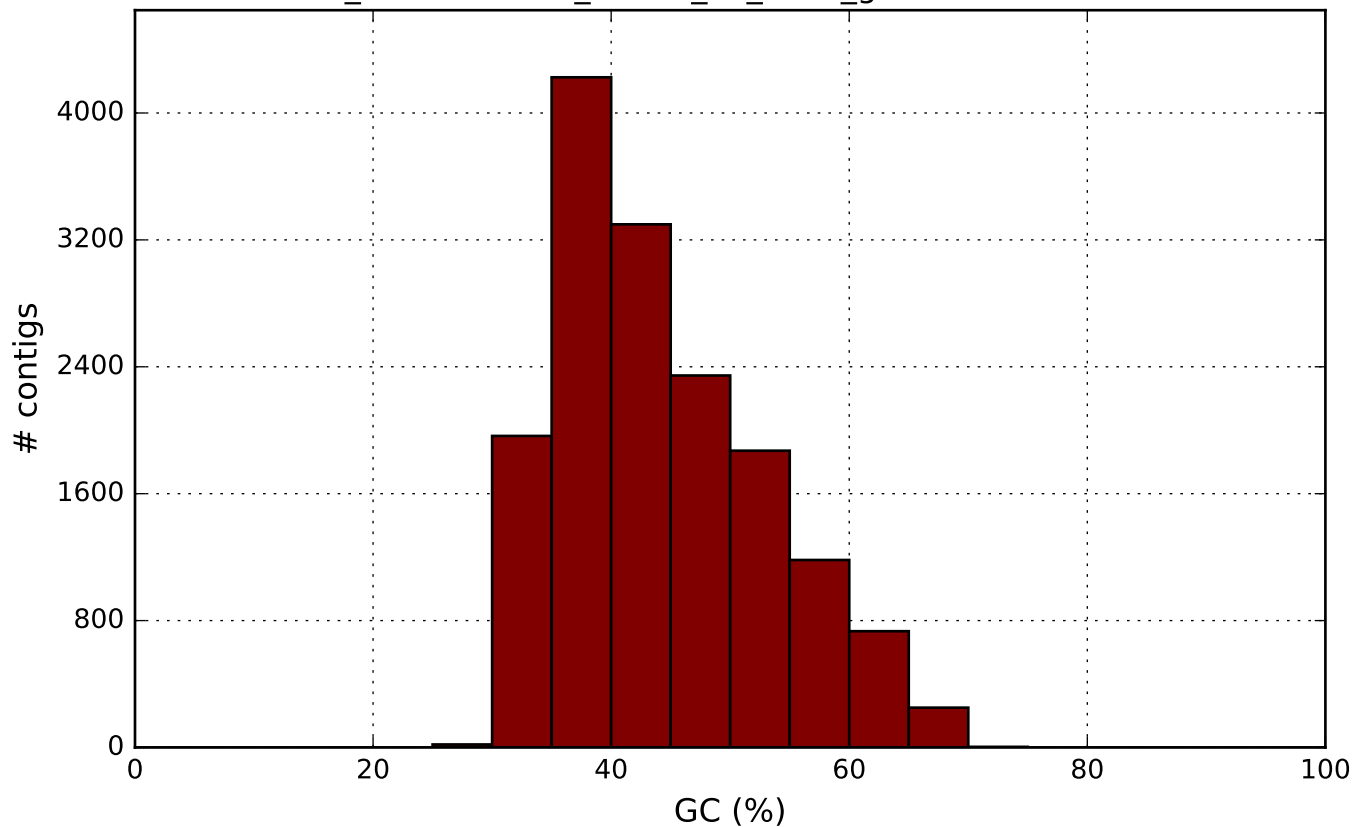
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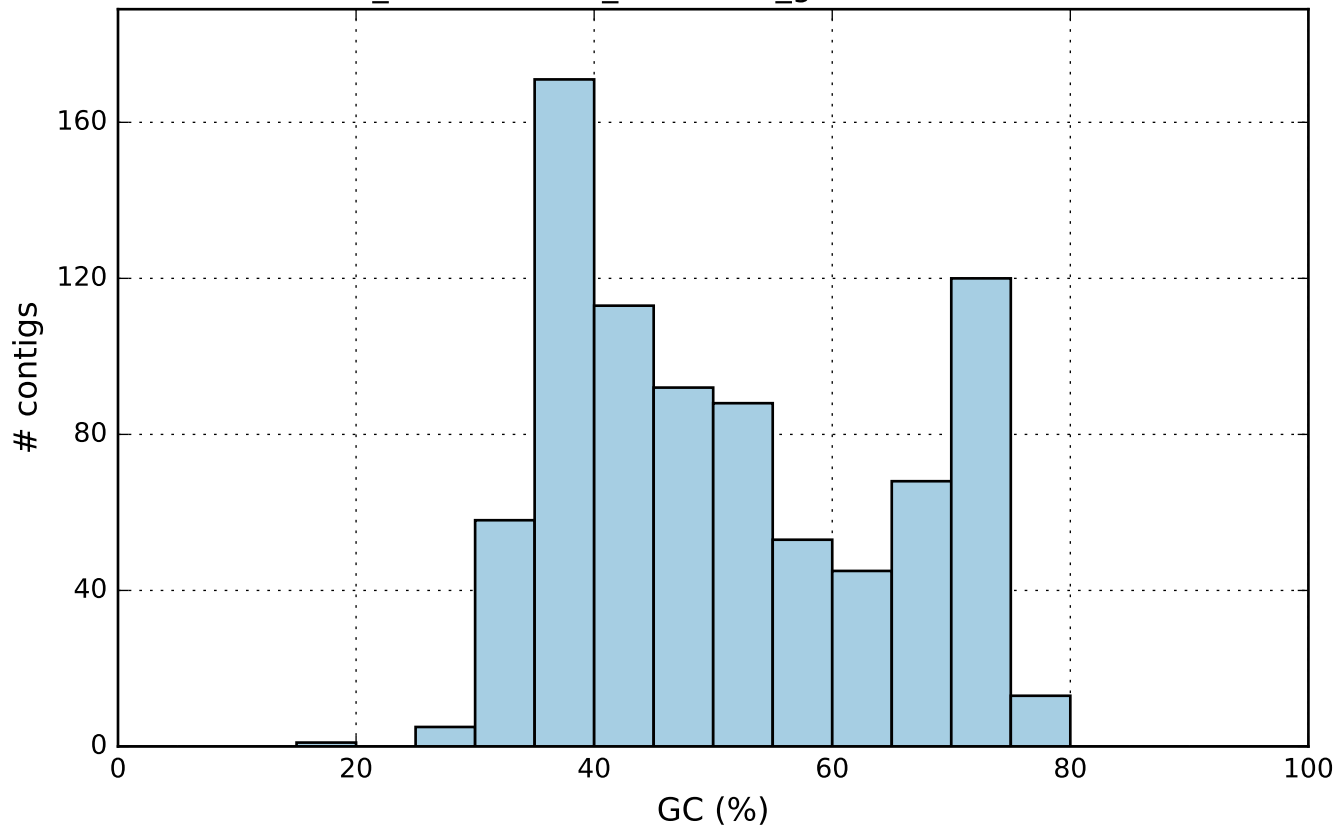
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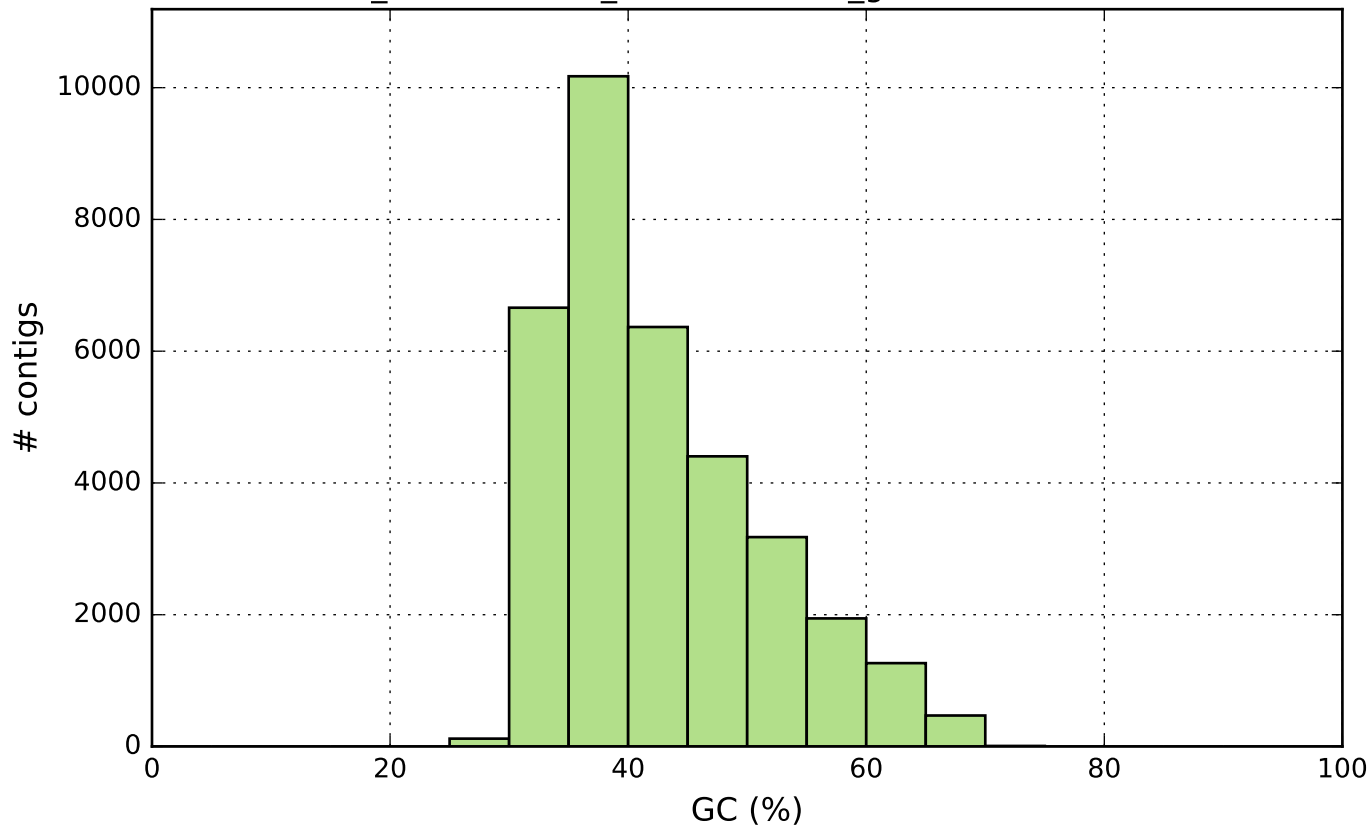
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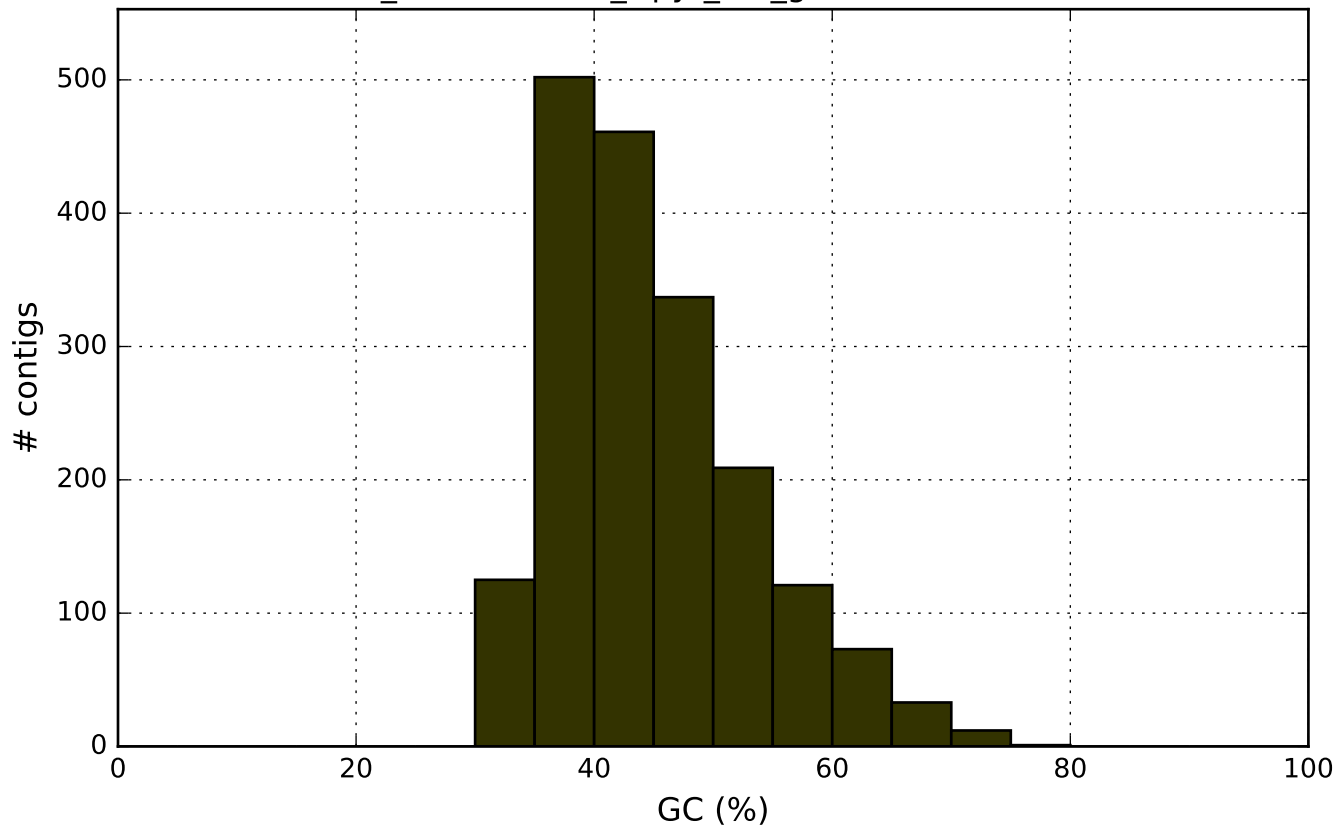
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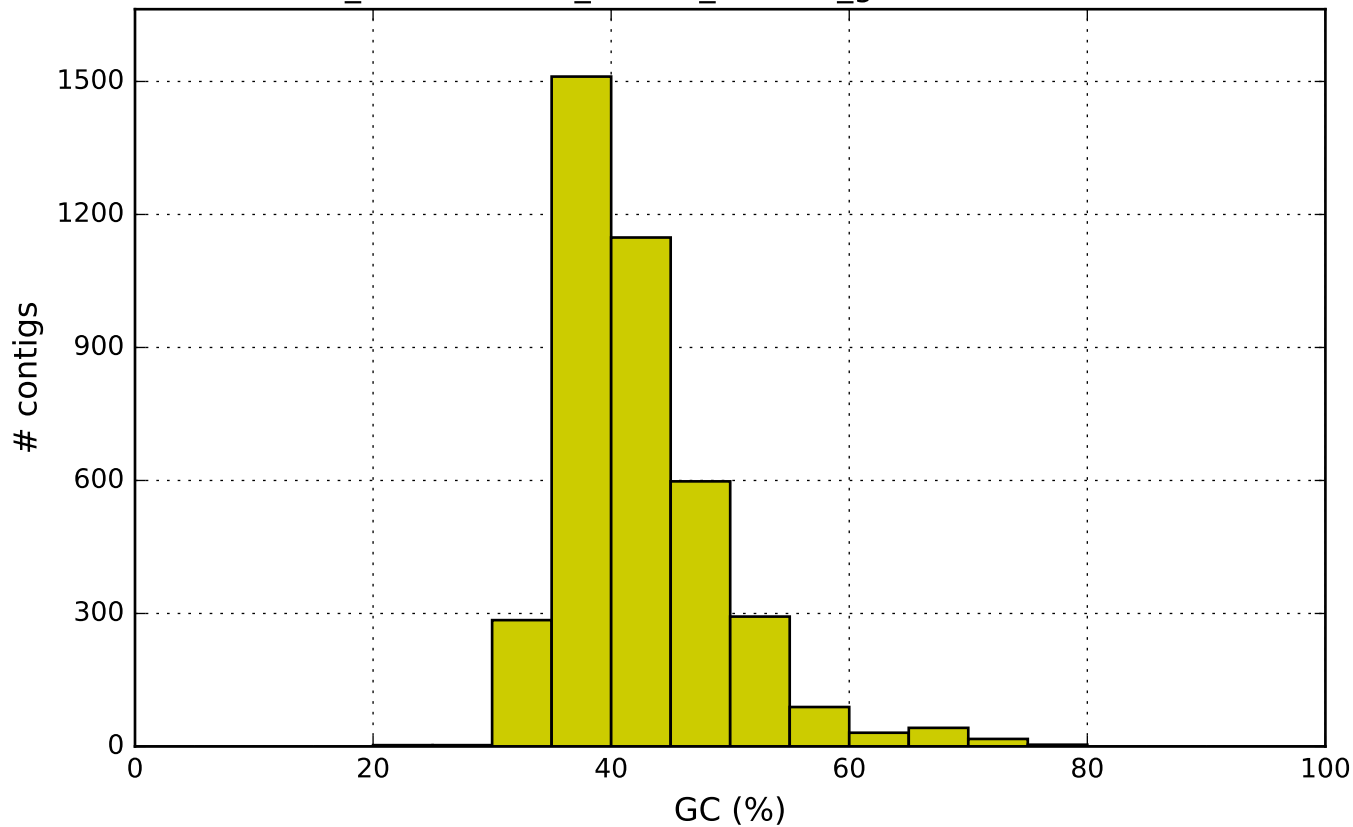
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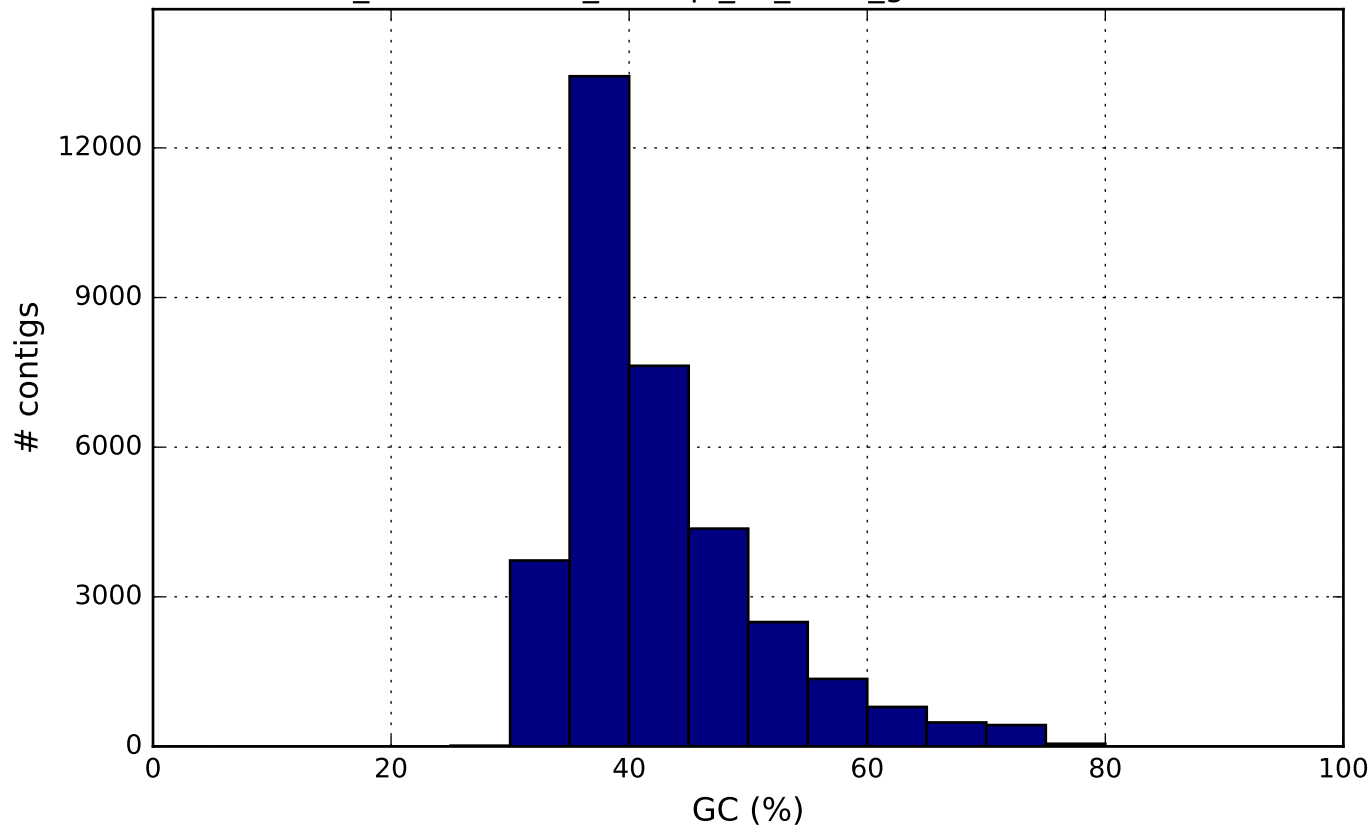
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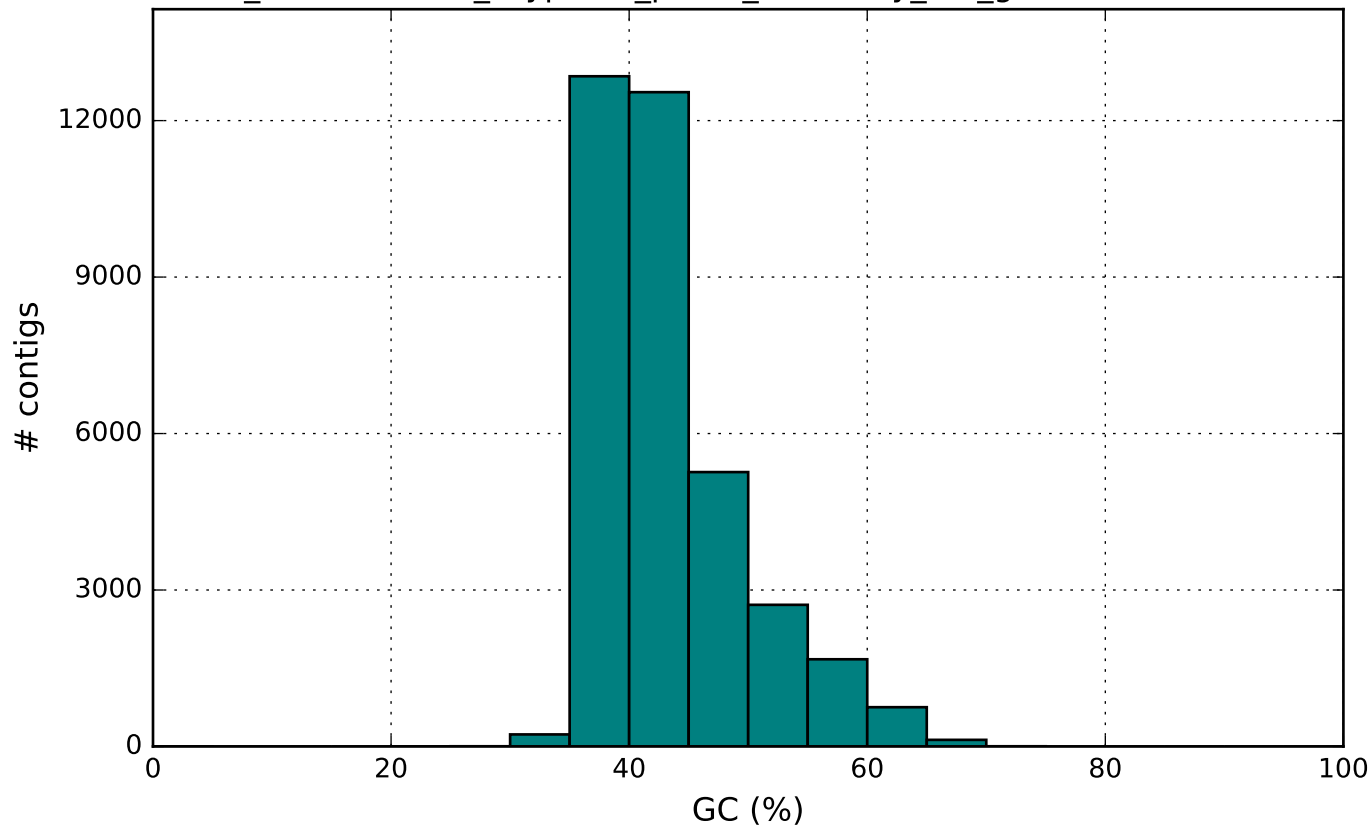
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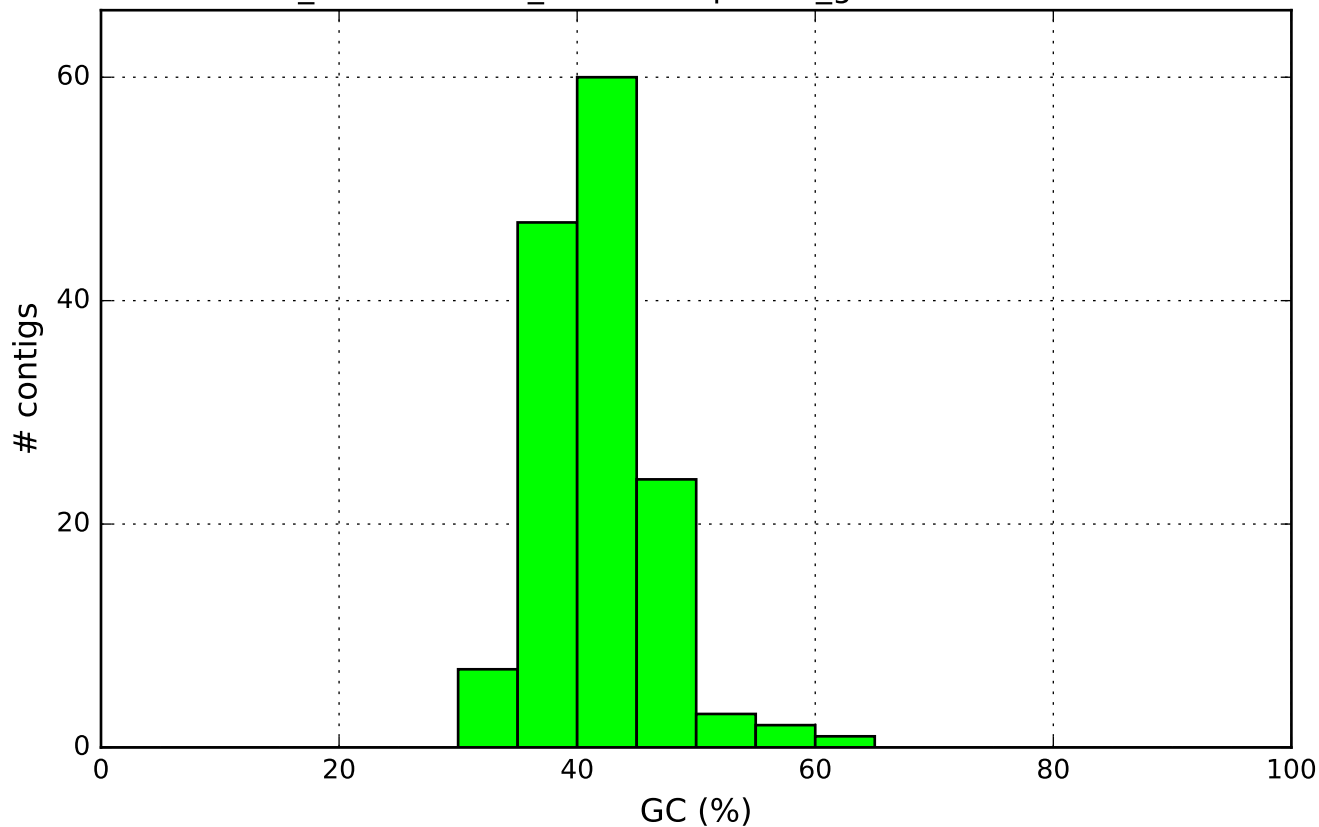
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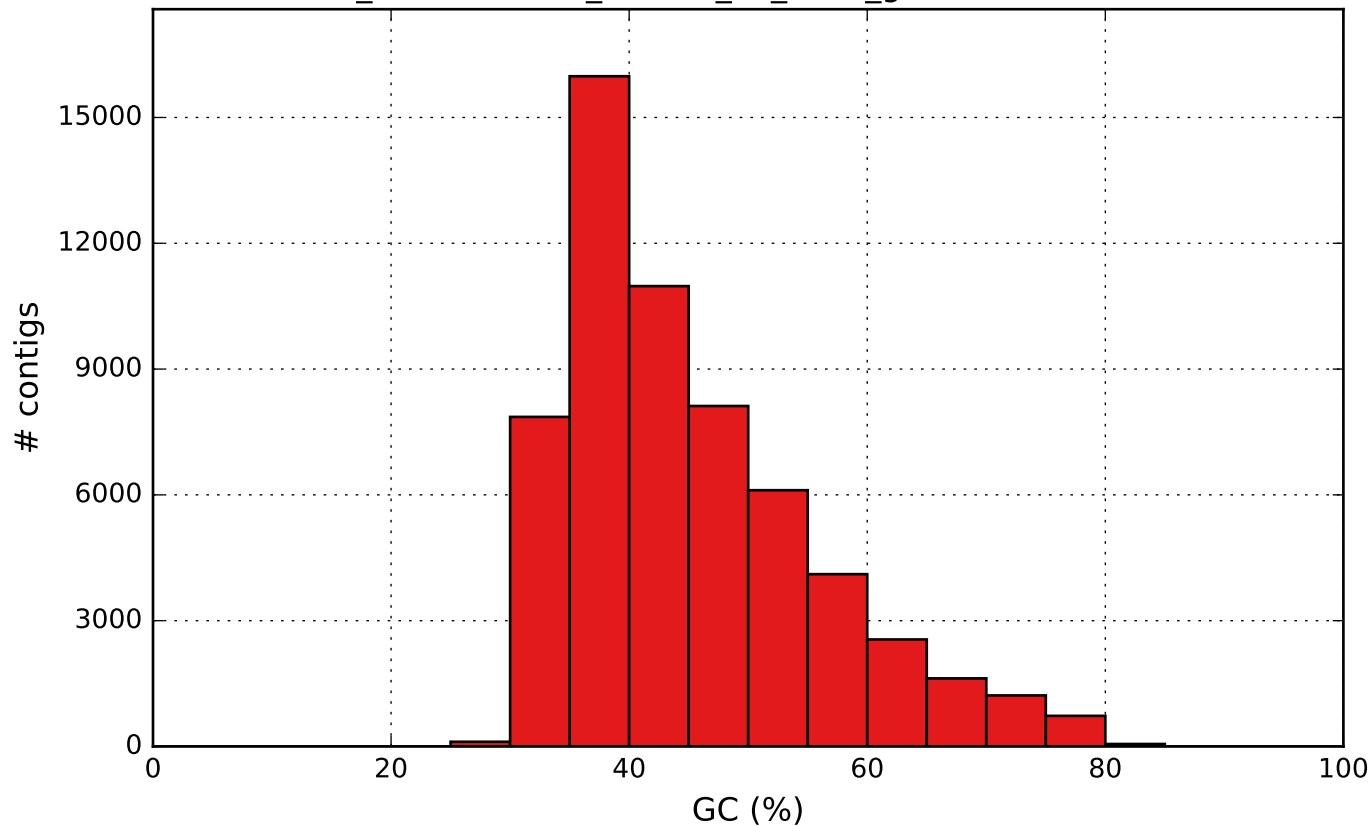
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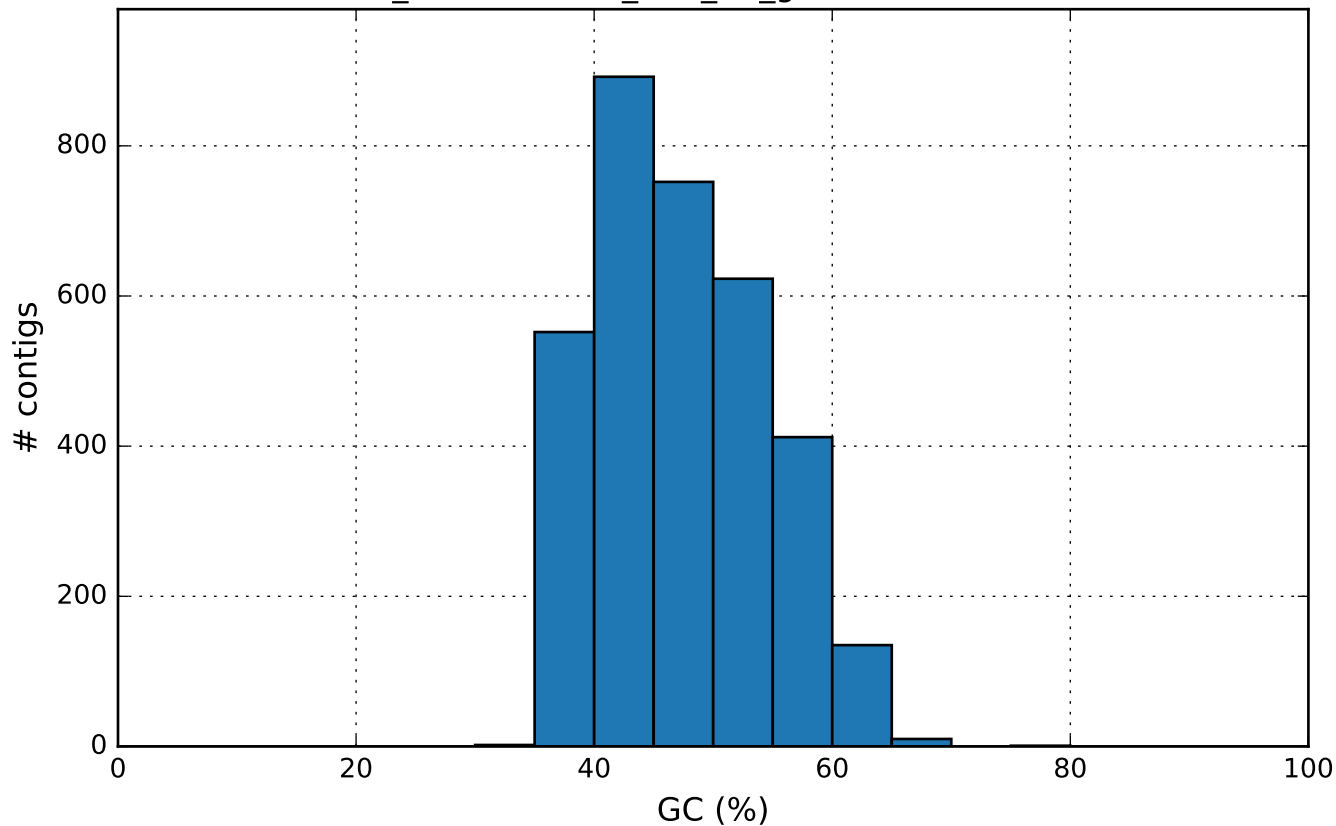
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