

All statistics are based on contigs of size  $\geq$  5000 bp, unless otherwise noted  
(e.g., "# contigs ( $\geq$  0 bp)" and "Total length ( $\geq$  0 bp)" include all contigs).

## Misassemblies report

	MP_helen_r10_bins_1	MP_helen_r10_bins_2	MP_helen_r10_bins_3	MP_r10_bins_1	MP_r10_bins_2	MP_r10_bins_3	r1_medaka_r10_bins_1	r1_medaka_r10_bins_2	r1_medaka_r10_bins_3	r2_medaka_r10_bins_1	r2_medaka_r10_bins_2	r2_medaka_r10_bins_3	racon_r1_r10_bins_1	racon_r1_r10_bins_2	racon_r1_r10_bins_3	racon_r2_r10_bins_1	racon_r2_r10_bins_2	racon_r2_r10_bins_3	raw_r10_bins_1	raw_r10_bins_2	raw_r10_bins_3
# misassemblies	14	13	14	14	13	14	14	13	14	14	13	14	14	13	14	13	14	14	14	12	12
# contig misassemblies	14	13	14	14	13	14	14	13	14	14	13	14	14	13	14	13	14	14	14	12	12
# c. relocations	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# c. translocations	14	13	14	14	13	14	14	13	14	14	13	14	14	13	14	13	14	14	12	12	
# c. inversions	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# scaffold misassemblies	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# s. relocations	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# s. translocations	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# s. inversions	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# misassembled contigs	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
Misassembled contigs length	2717987	2718054	2718086	2719777	2719835	2719835	2719552	2719502	2719543	2719584	2719579	2719559	2720311	2720303	2720298	2720378	2720381	2720321	2718434	2718434	2718864
# possibly misassembled contigs	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
# possible misassemblies	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	18	16	19
# local misassemblies	10	10	9	9	10	9	9	9	11	9	9	10	9	9	11	8	10	9	9	20	20
# scaffold gap ext. mis.	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# scaffold gap loc. mis.	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
# misassemblies caused by fragmented reference	26	26	26	26	26	26	26	26	25	26	26	26	25	26	26	26	26	23	25	23	
# unaligned mis. contigs	6	6	6	6	6	6	6	6	6	6	6	6	6	6	6	6	6	5	5	5	
# mismatches	6852	6782	6822	7666	7678	7699	7737	7616	7744	7598	7565	7734	7888	7679	7646	8011	7763	7981	6761	6483	6800
# indels	1149	934	948	2984	2996	3012	2018	2047	2090	2032	2039	2099	2970	2948	2973	2932	2915	2999	4328	4207	4307
# indels (<= 5 bp)	1086	871	887	2914	2930	2942	1967	1995	2039	1981	1988	2048	2916	2893	2922	2880	2864	2947	4269	4147	4250
# indels (> 5 bp)	63	63	61	70	66	70	51	52	51	51	51	54	55	51	52	51	59	60	57		
Indels length	4903	4606	4601	6868	6836	6849	5363	5399	5444	5369	5381	5442	6706	6716	6642	6613	6559	6709	8430	8316	8350

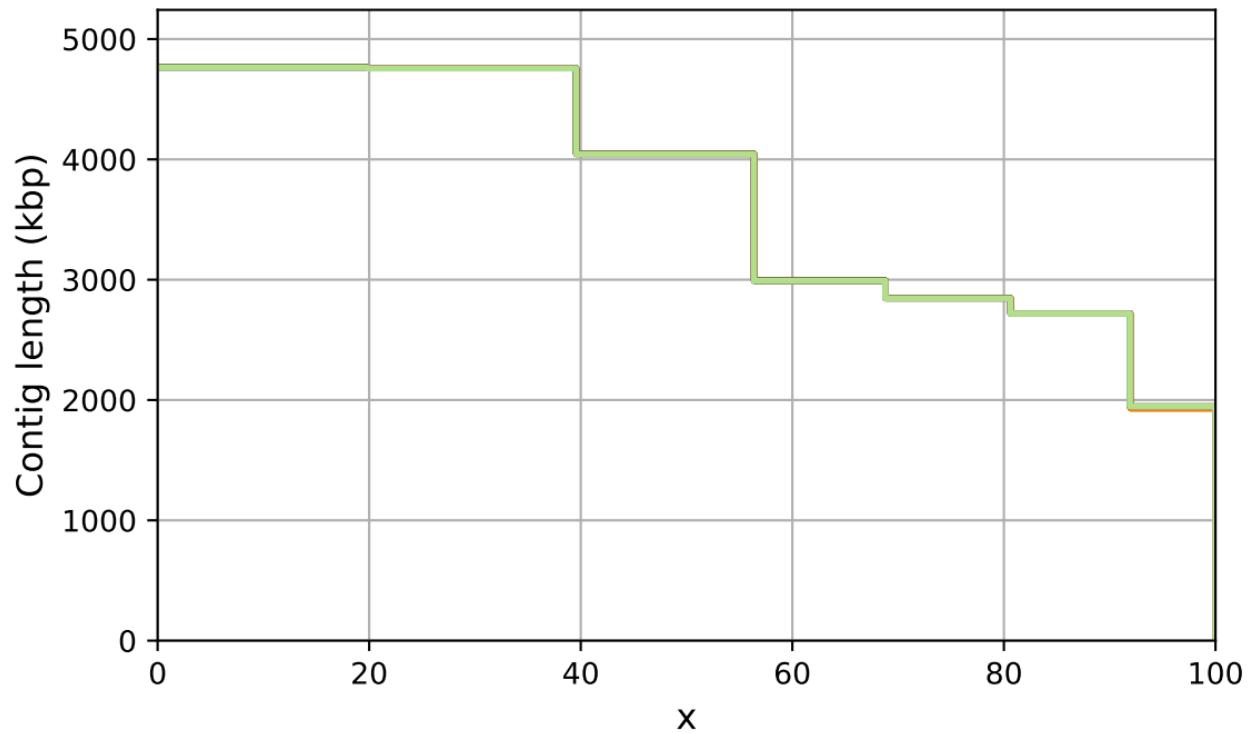
All statistics are based on contigs of size  $\geq 5000$  bp, unless otherwise noted  
(e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).

## Unaligned report

	MP_helen_r10_bins_1	MP_helen_r10_bins_2	MP_helen_r10_bins_3	MP_r10_bins_1	MP_r10_bins_2	MP_r10_bins_3	r1_medaka_r10_bins_1	r1_medaka_r10_bins_2	r1_medaka_r10_bins_3	racon_r1_r10_bins_1	racon_r1_r10_bins_2	racon_r1_r10_bins_3	racon_r2_r10_bins_1	racon_r2_r10_bins_2	racon_r2_r10_bins_3	raw_r10_bins_1	raw_r10_bins_2	raw_r10_bins_3	
# fully unaligned contigs	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Fully unaligned length	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
# partially unaligned contigs	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	
Partially unaligned length	21271543	21274547	21271970	21275539	21275911	21275190	21273039	21275906	21273107	21266527	21269791	21266383	21261081	21266891	21262972	21254935	21263159	21252871	21318428
# N's	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	

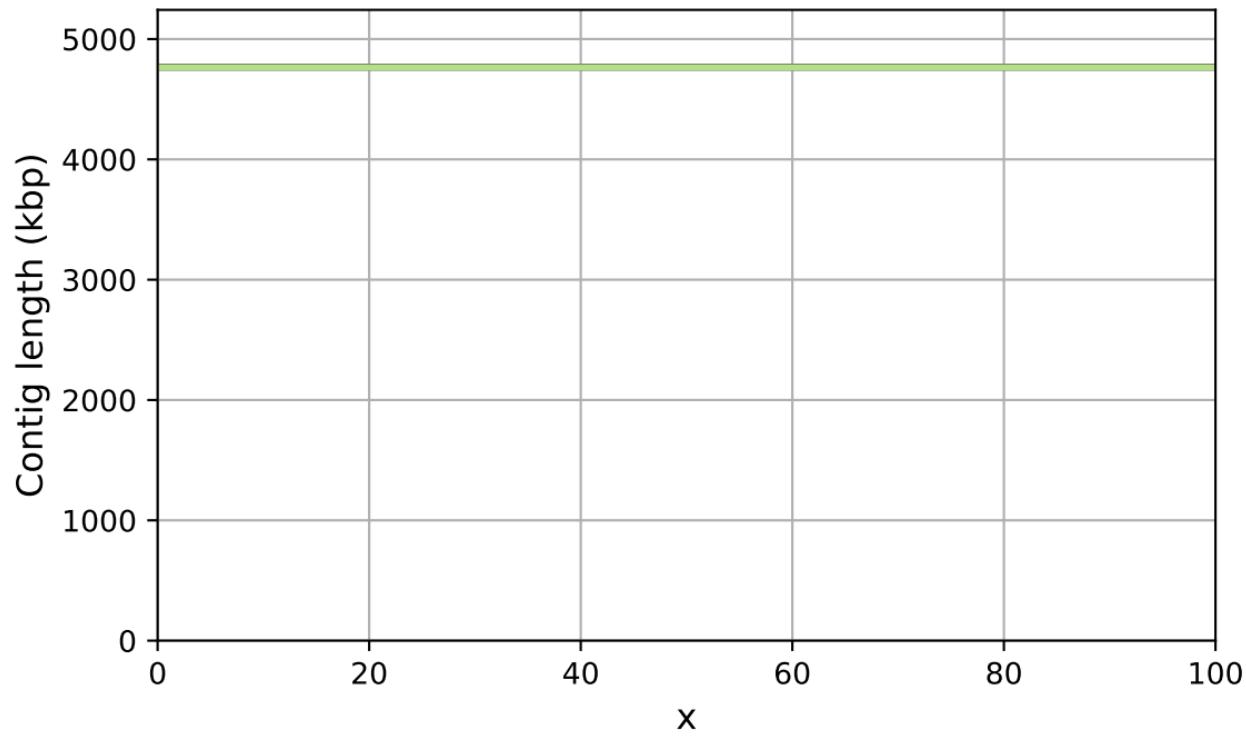
All statistics are based on contigs of size  $\geq 5000$  bp, unless otherwise noted  
(e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).

Nx



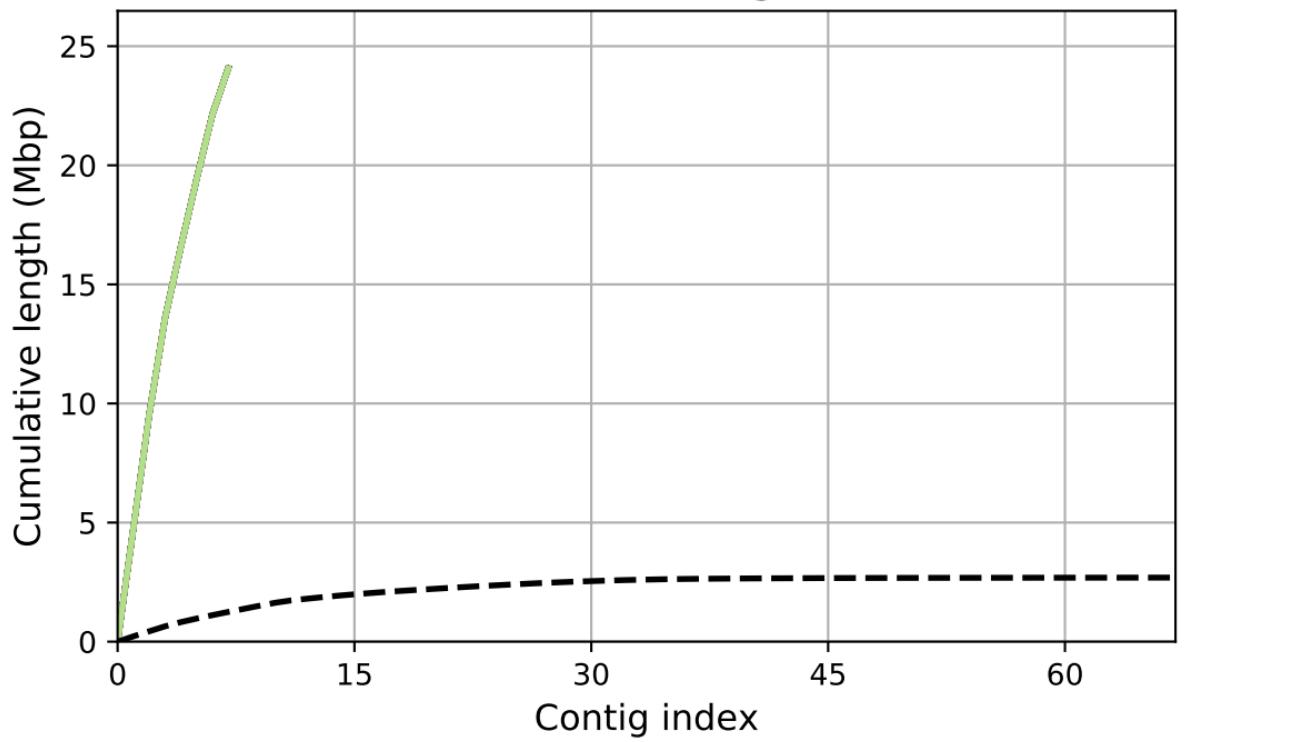
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP\_r10\_bins\_1
- r1\_medaka\_r10\_bins\_2
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- racon\_r1\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3

NGx



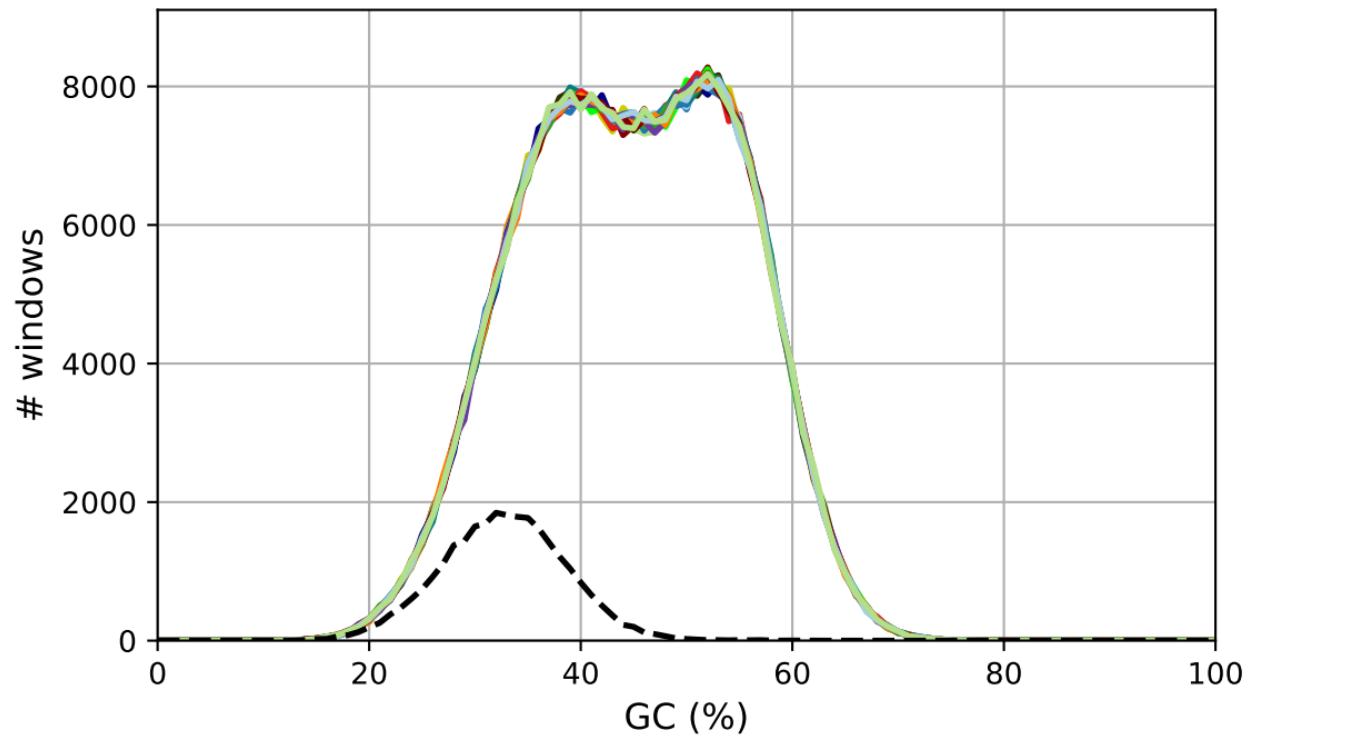
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins 1
- r1\_medaka\_r10\_bins\_2
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- racon\_r1\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3

Cumulative length



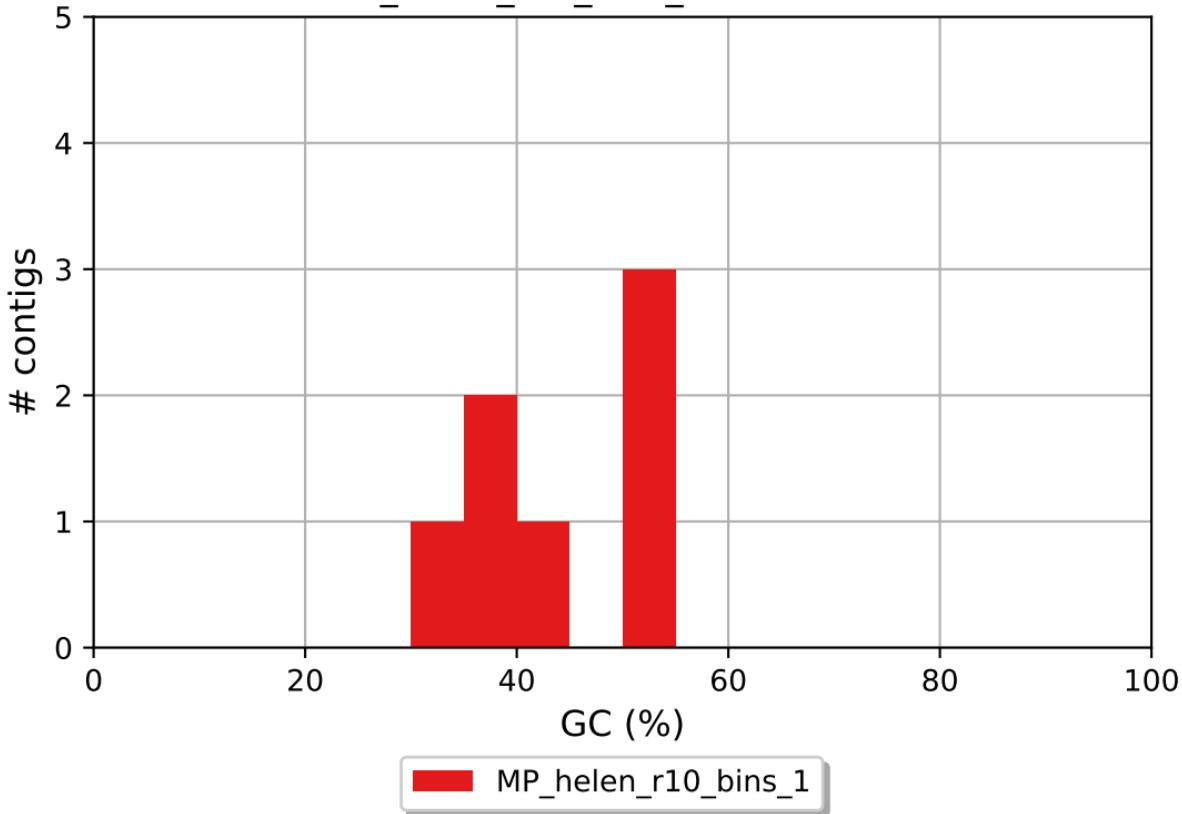
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP\_r10\_bins\_1
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- r2\_medaka\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3
- raw\_r10\_bins\_1

## GC content

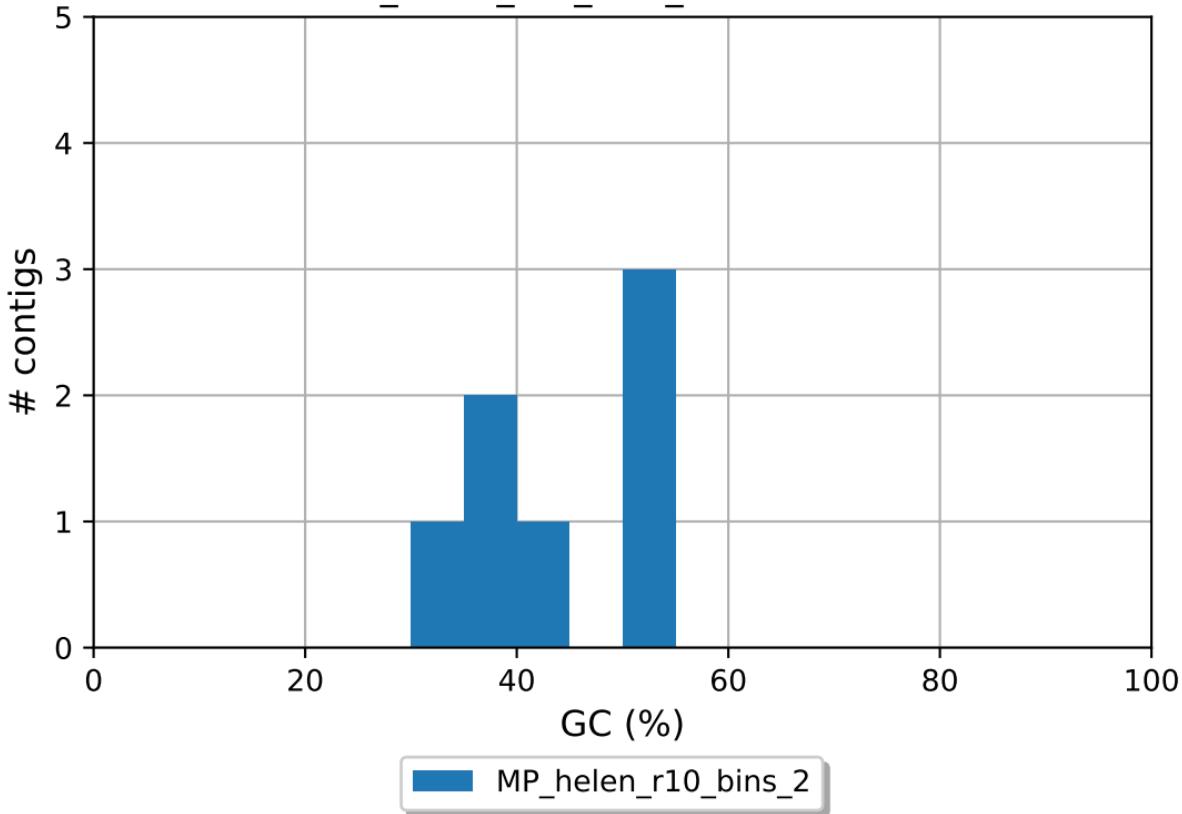


- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins 1
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- r2 medaka r10 bins 3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3
- raw r10 bins 1

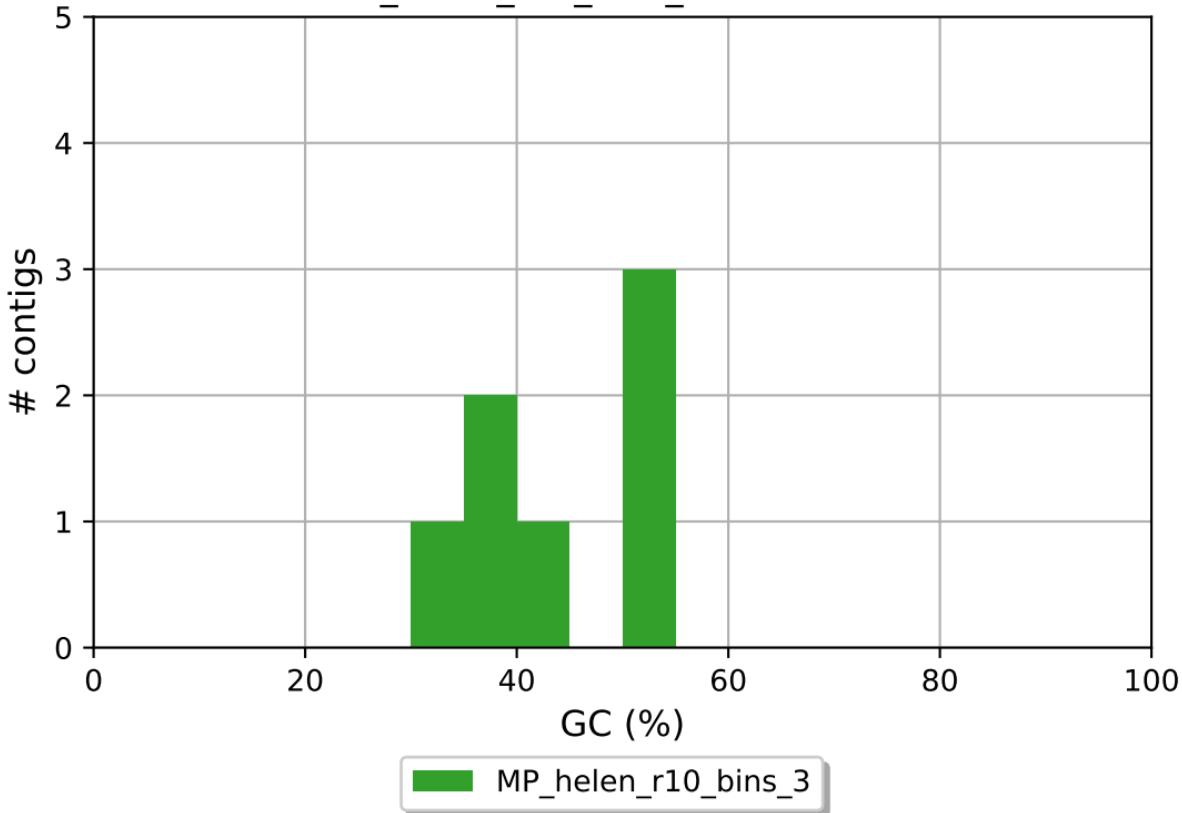
MP\_helen\_r10\_bins\_1 GC content



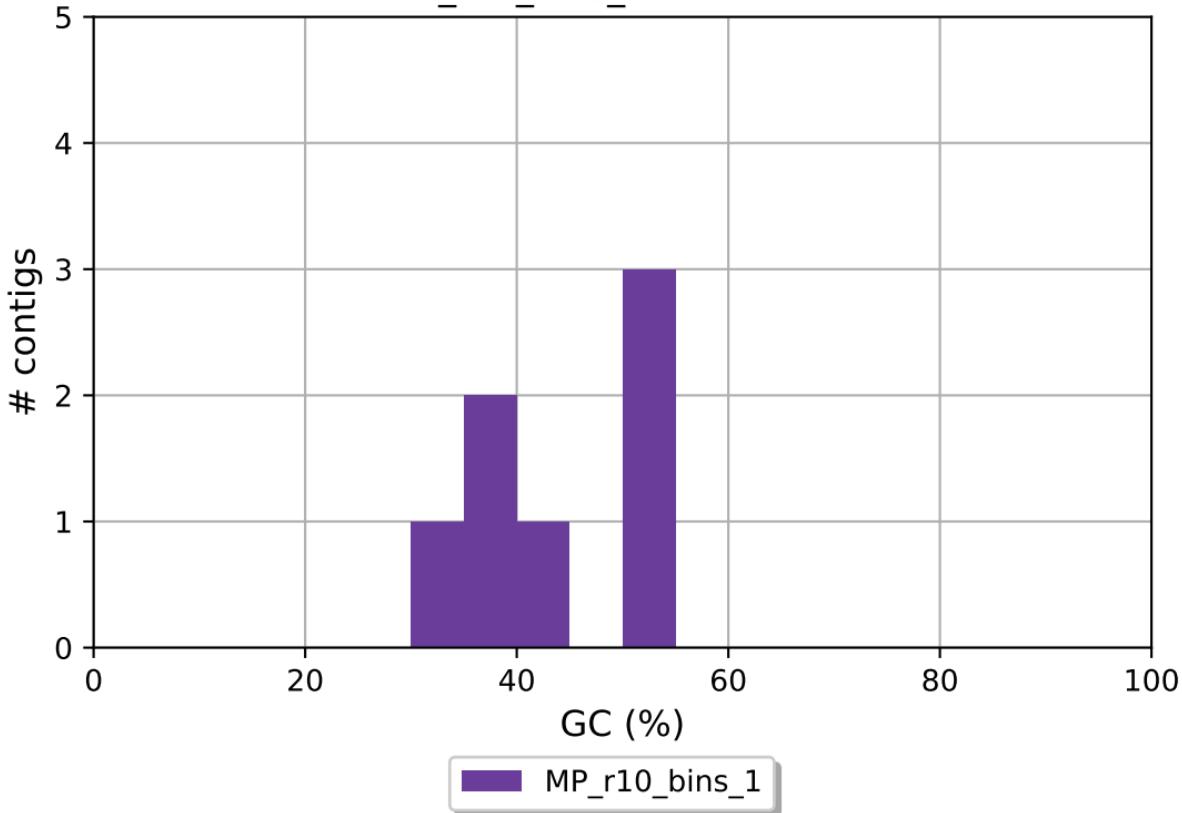
MP\_helen\_r10\_bins\_2 GC content



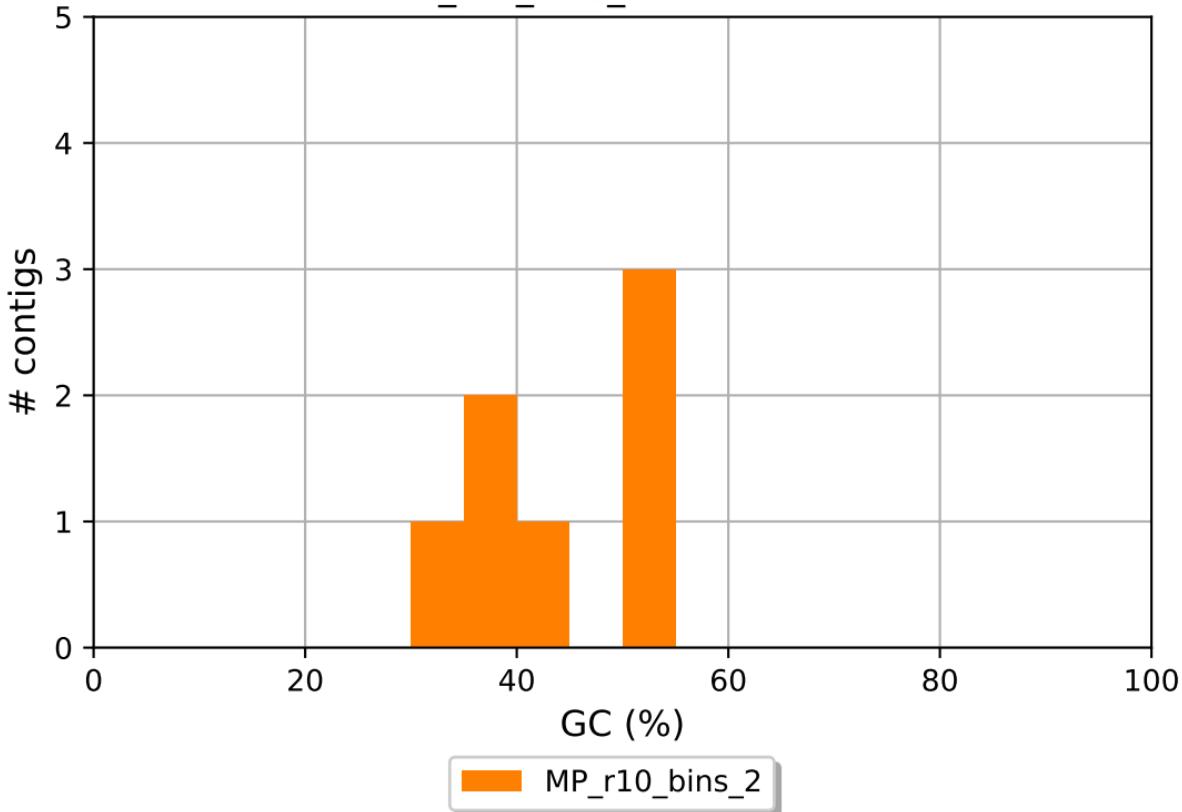
MP\_helen\_r10\_bins\_3 GC content



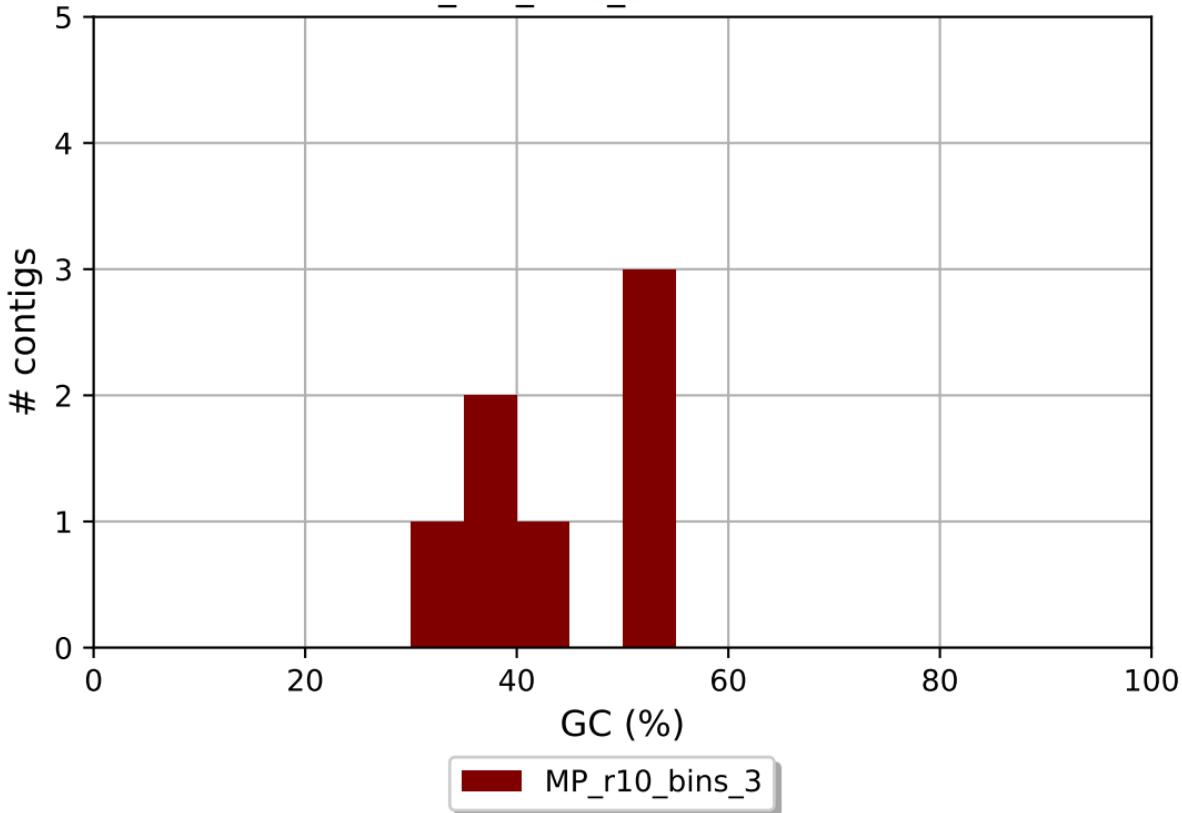
MP\_r10\_bins\_1 GC content



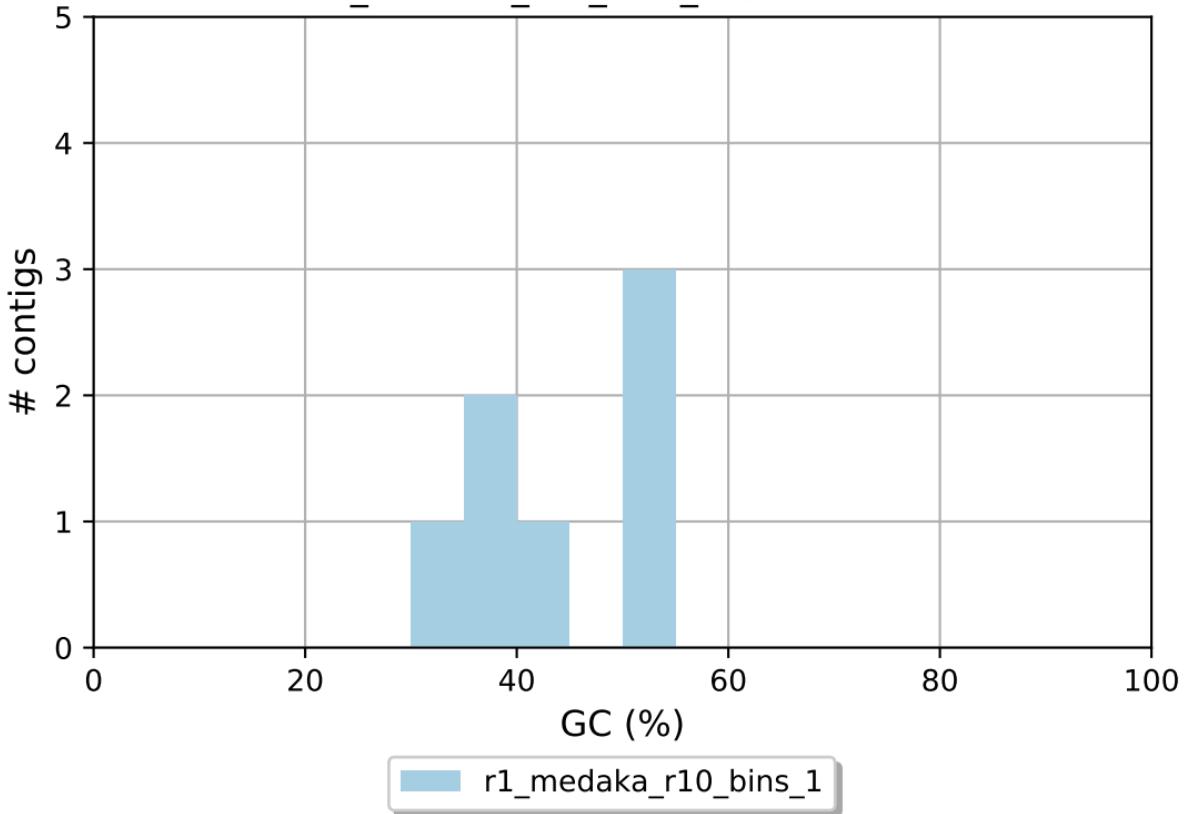
MP\_r10\_bins\_2 GC content



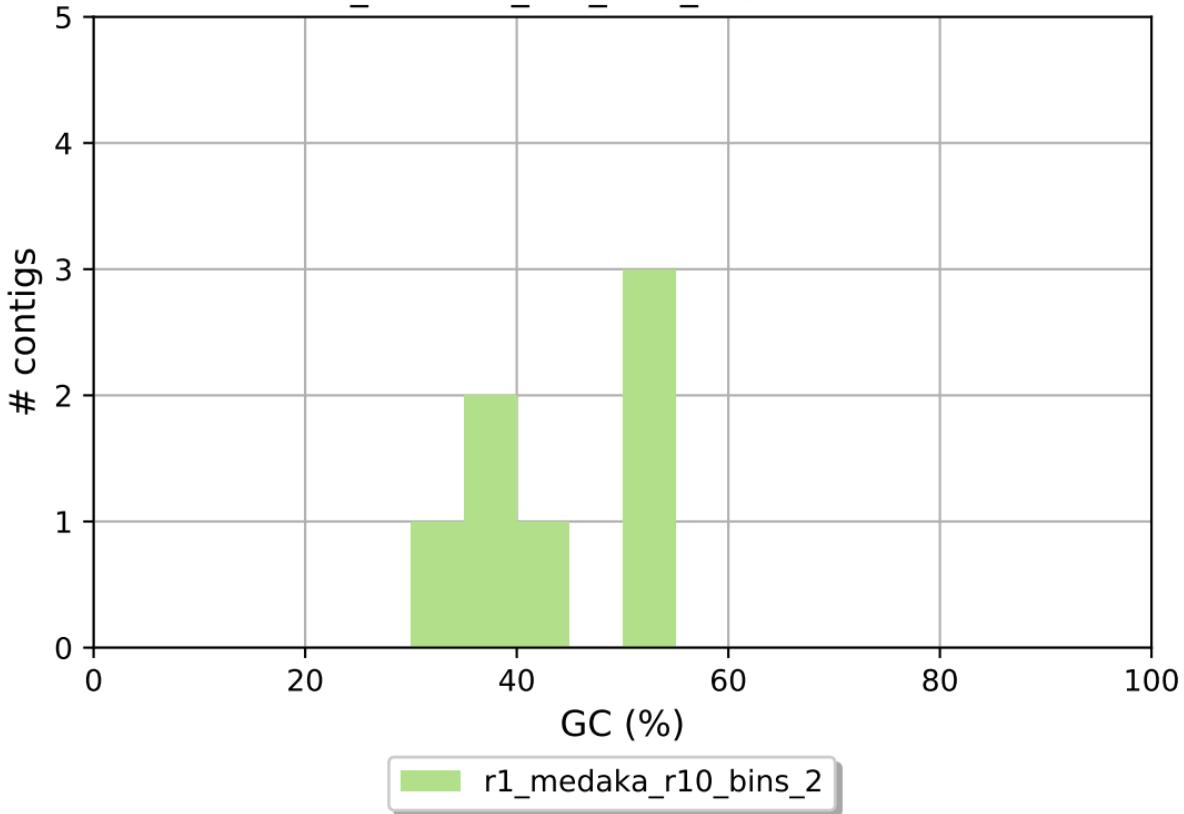
MP\_r10\_bins\_3 GC content



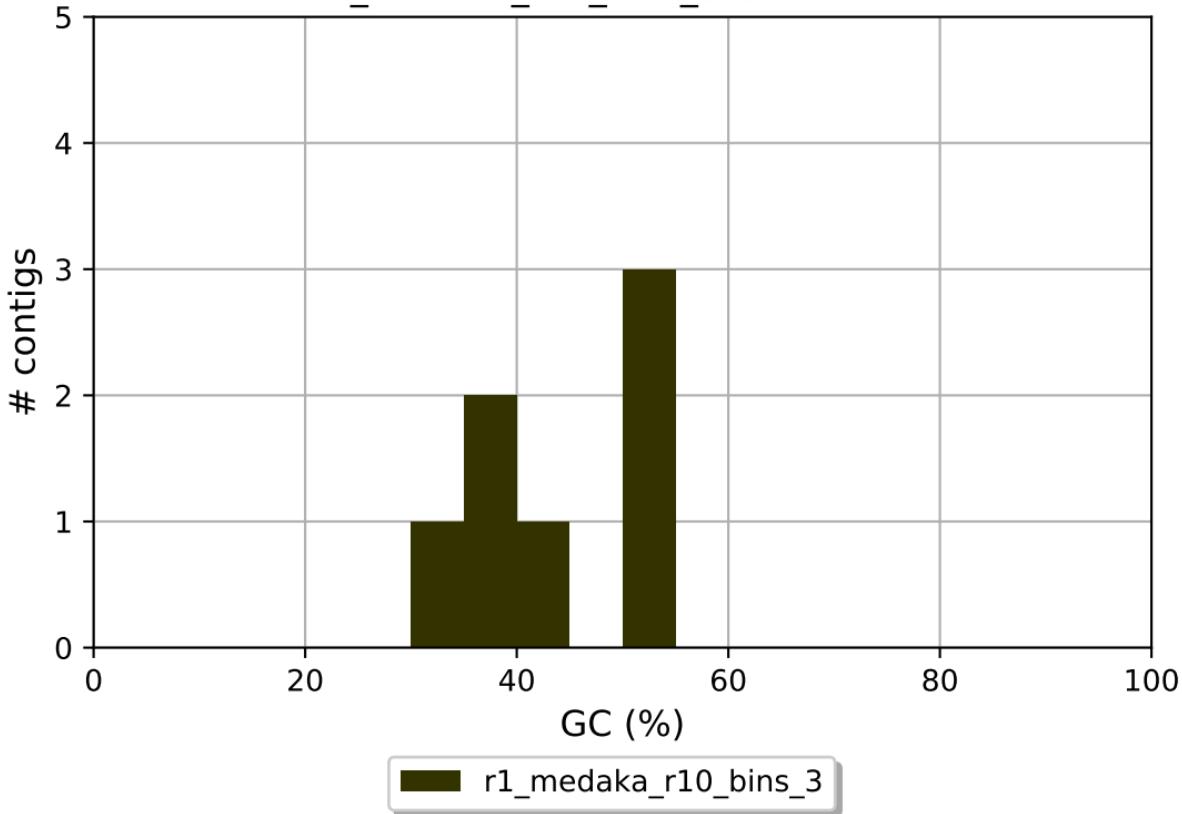
r1\_medaka\_r10\_bins\_1 GC content



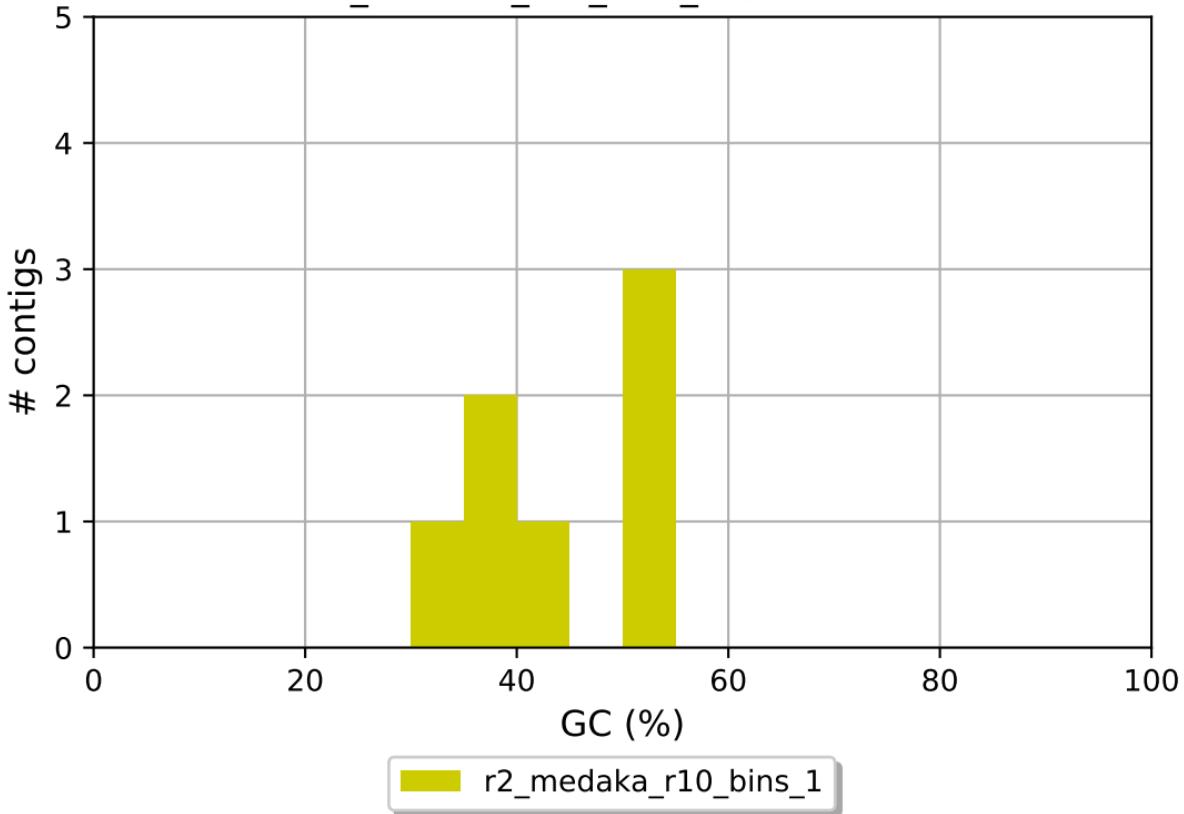
r1\_medaka\_r10\_bins\_2 GC content



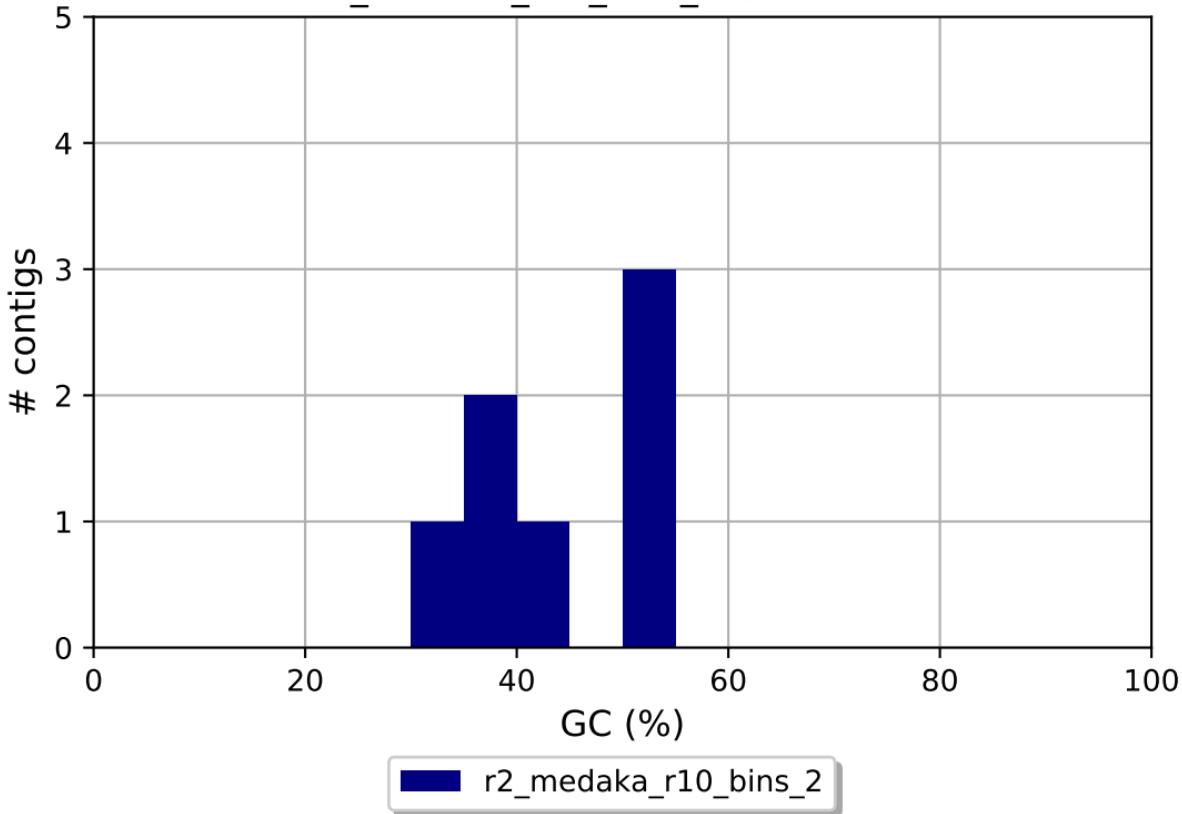
r1\_medaka\_r10\_bins\_3 GC content



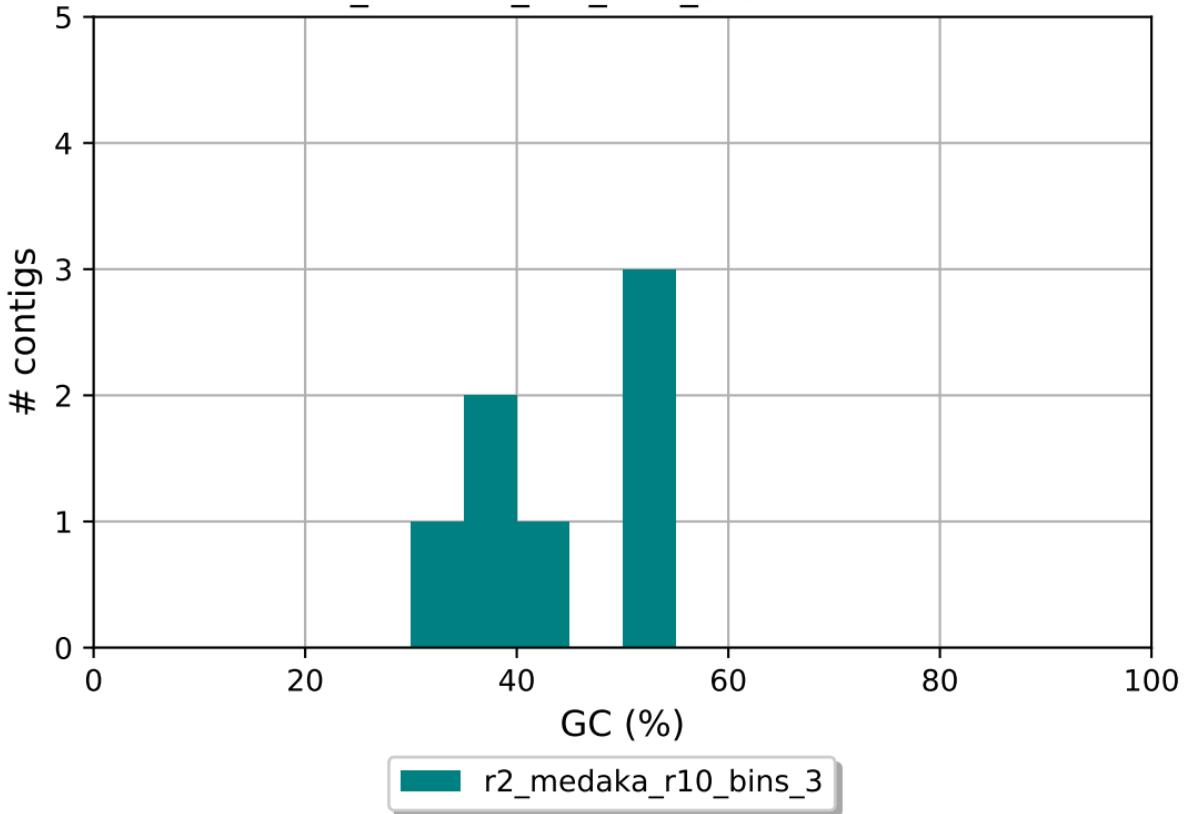
r2\_medaka\_r10\_bins\_1 GC content



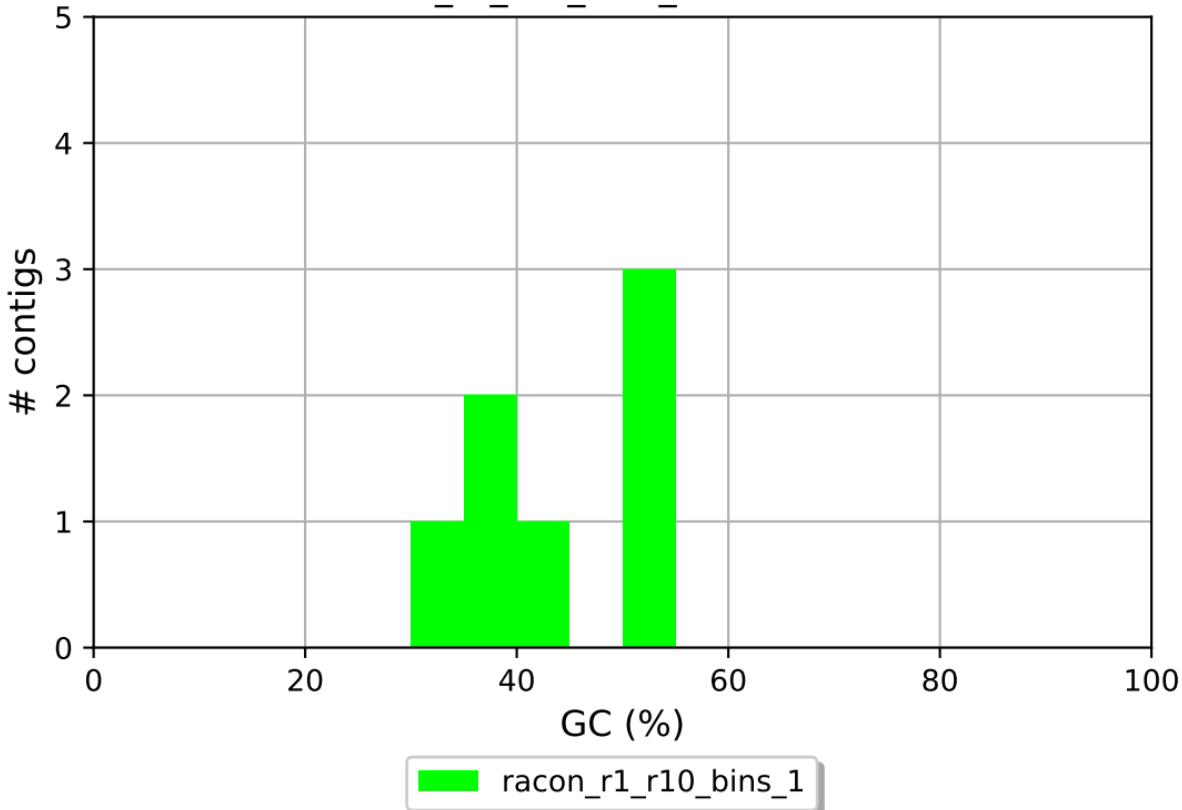
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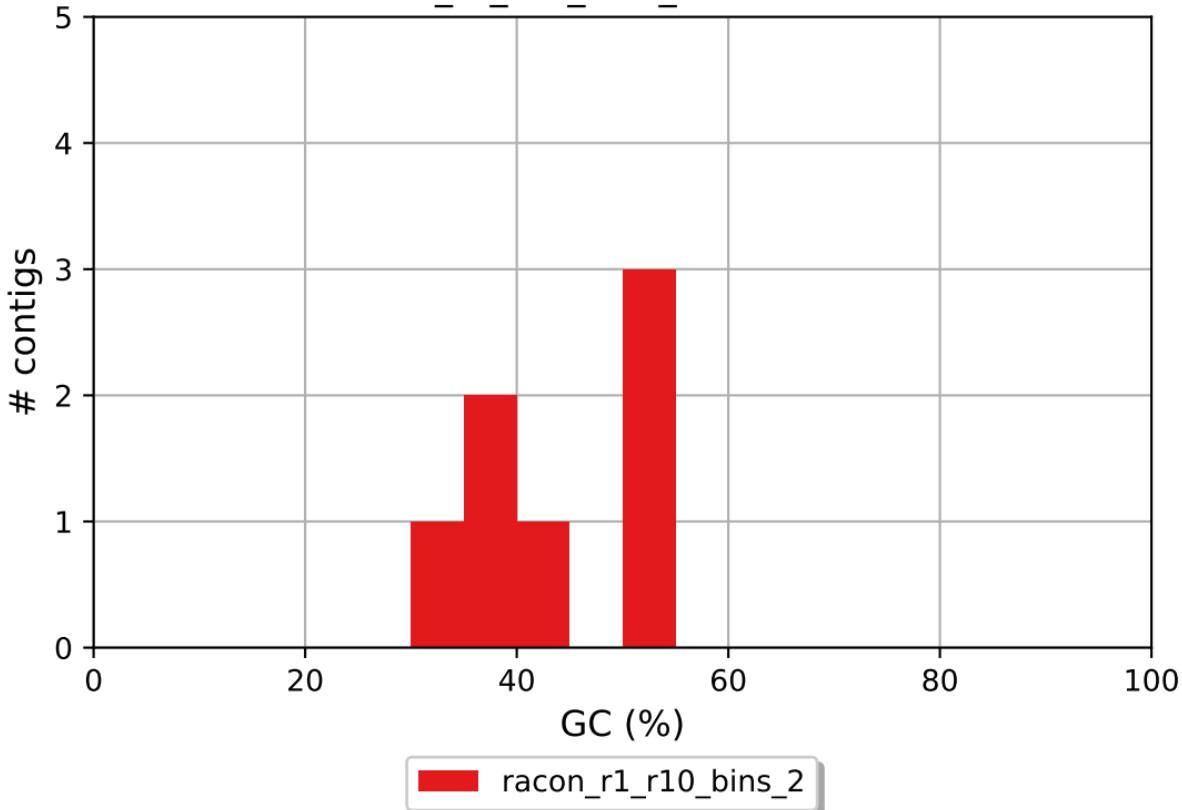
r2\_medaka\_r10\_bins\_3 GC content



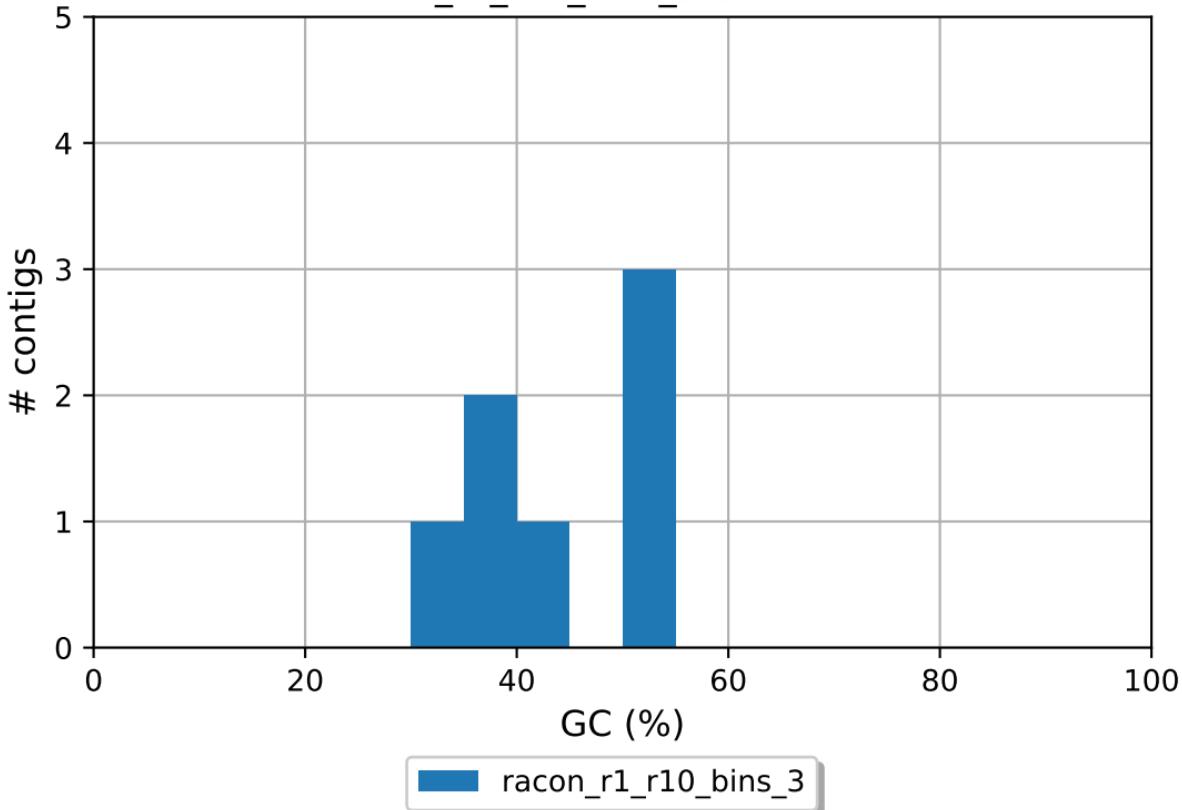
### racon\_r1\_r10\_bins\_1 GC content



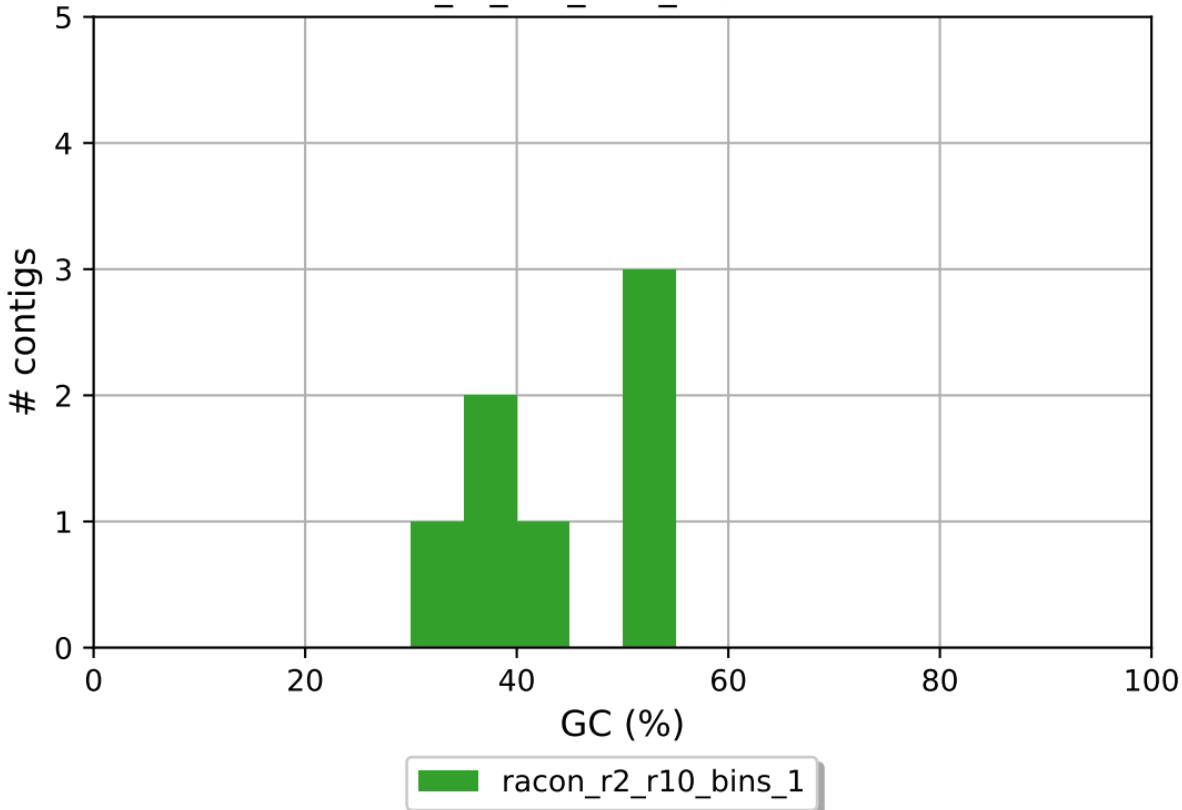
### racon\_r1\_r10\_bins\_2 GC content



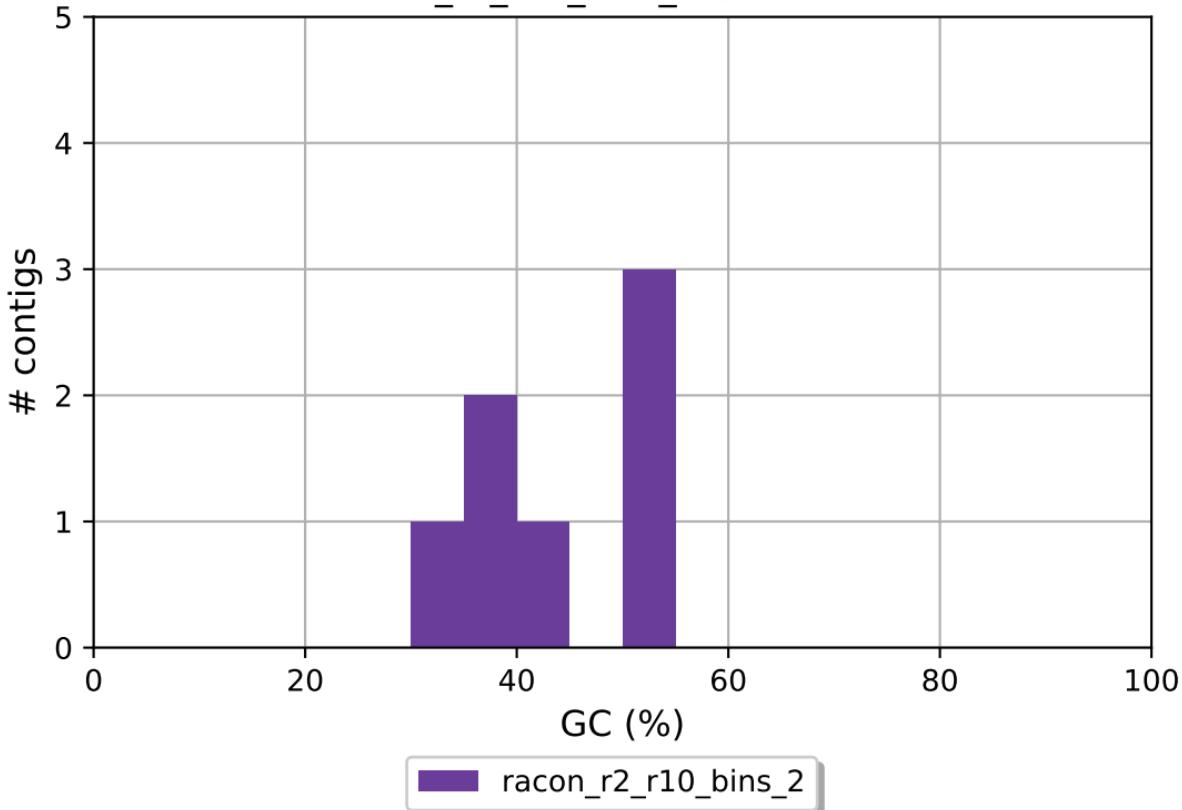
### racon\_r1\_r10\_bins\_3 GC content



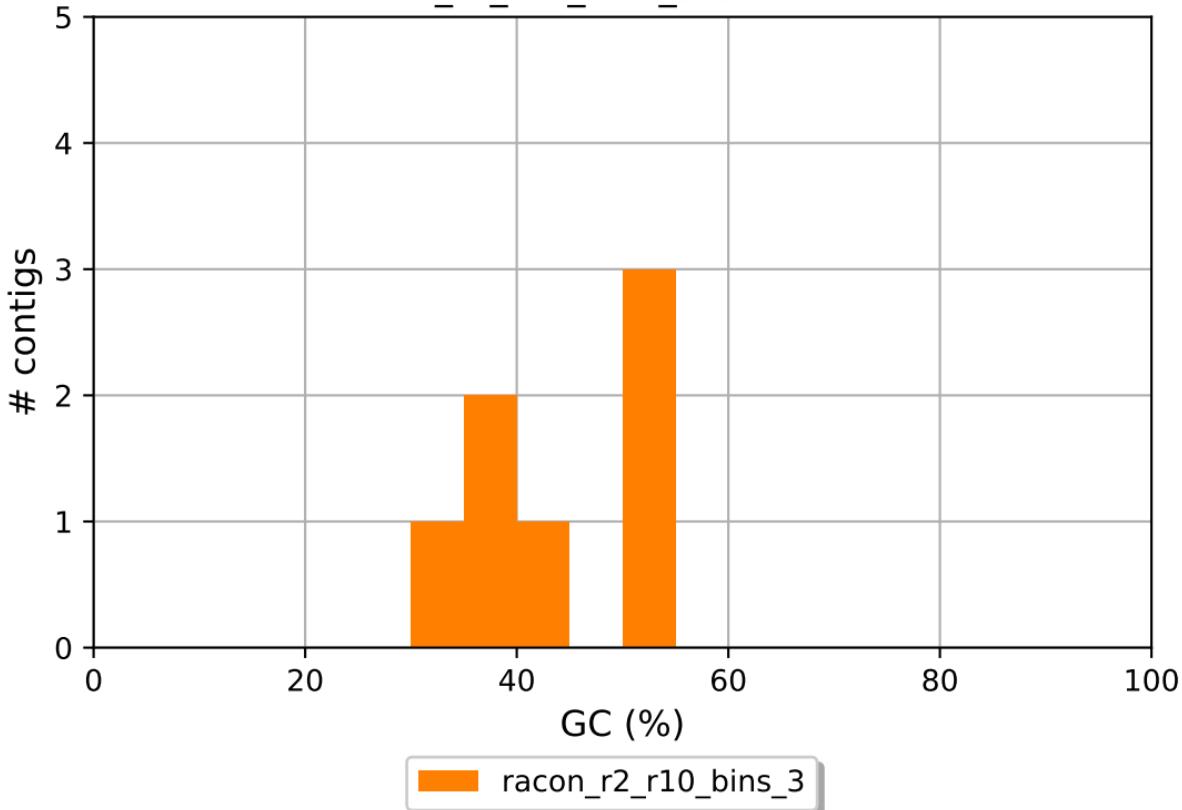
### racon\_r2\_r10\_bins\_1 GC content



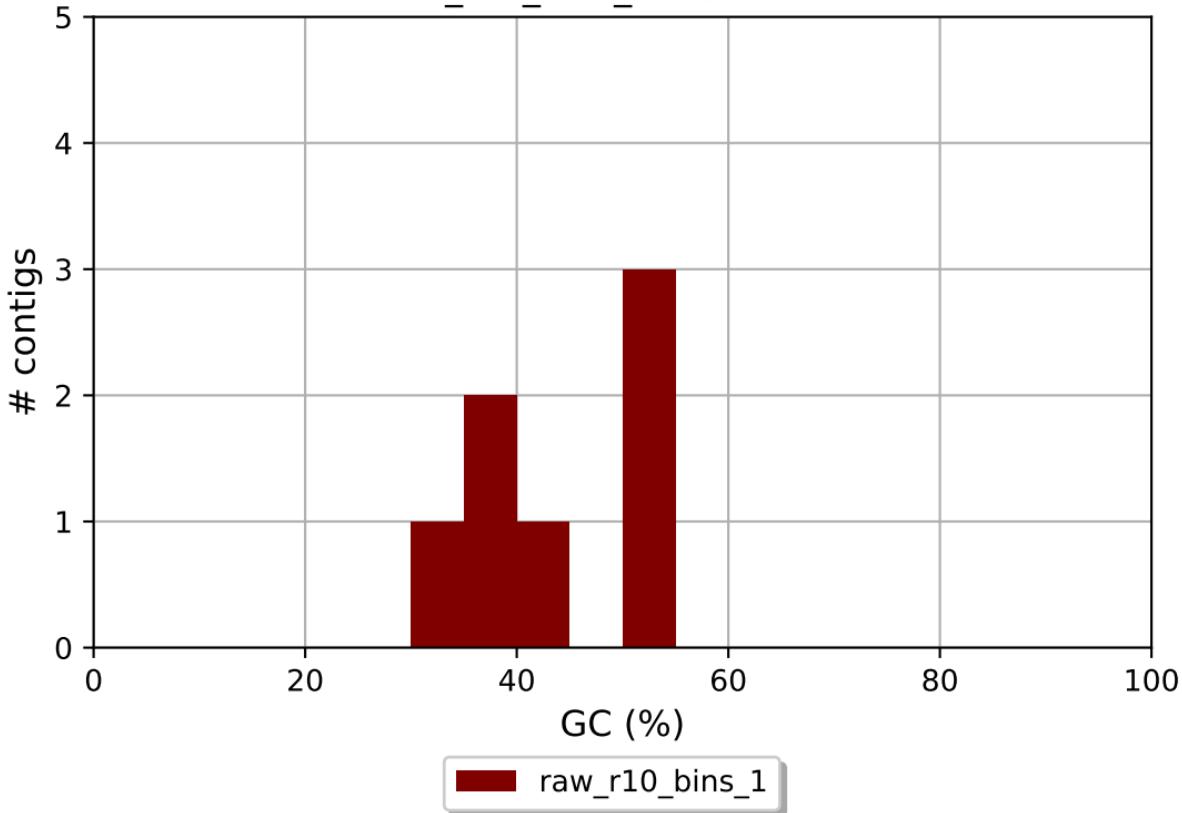
### racon\_r2\_r10\_bins\_2 GC content



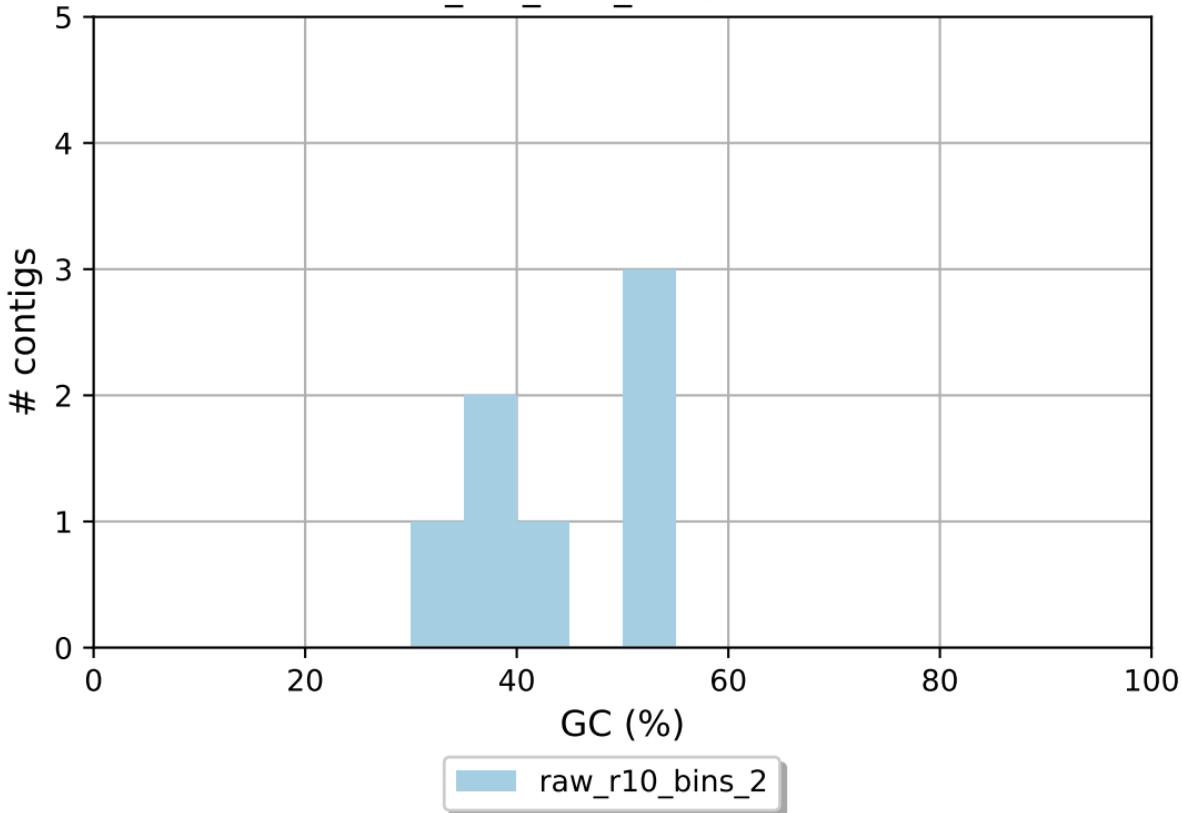
racon\_r2\_r10\_bins\_3 GC content



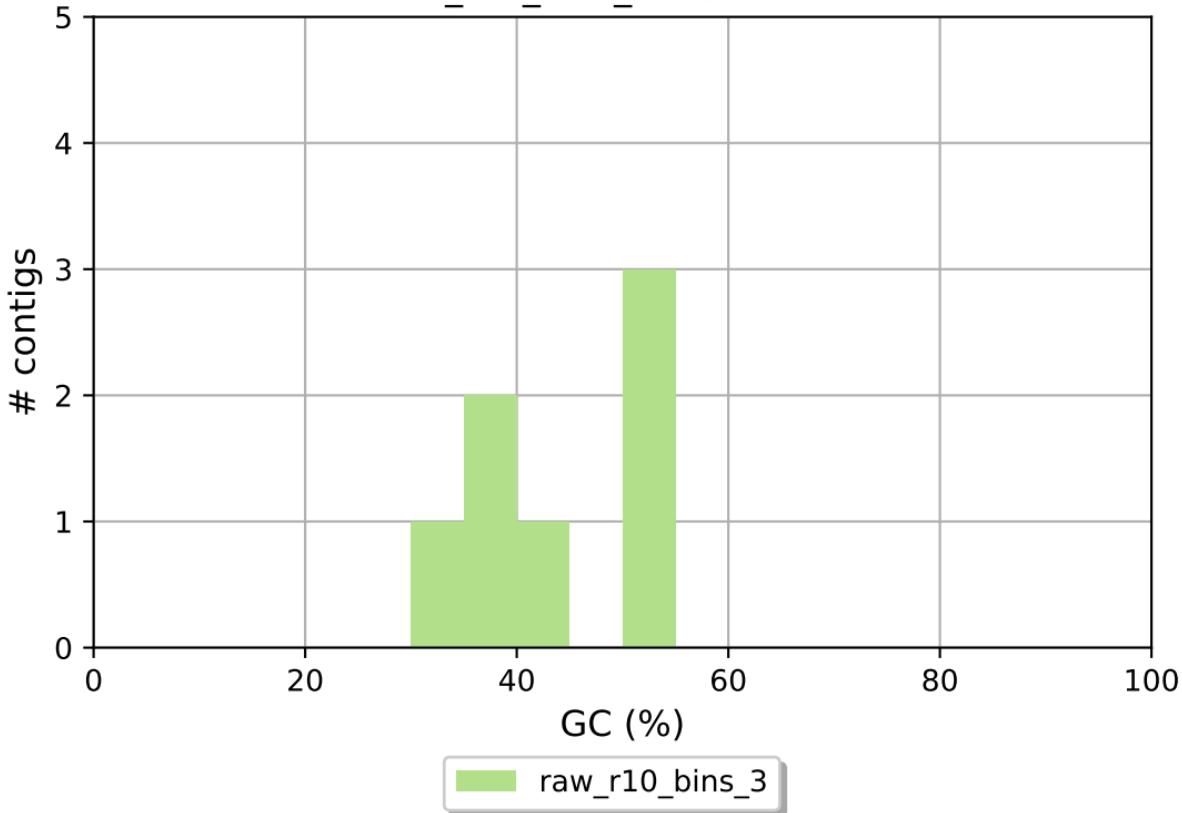
raw\_r10\_bins\_1 GC content



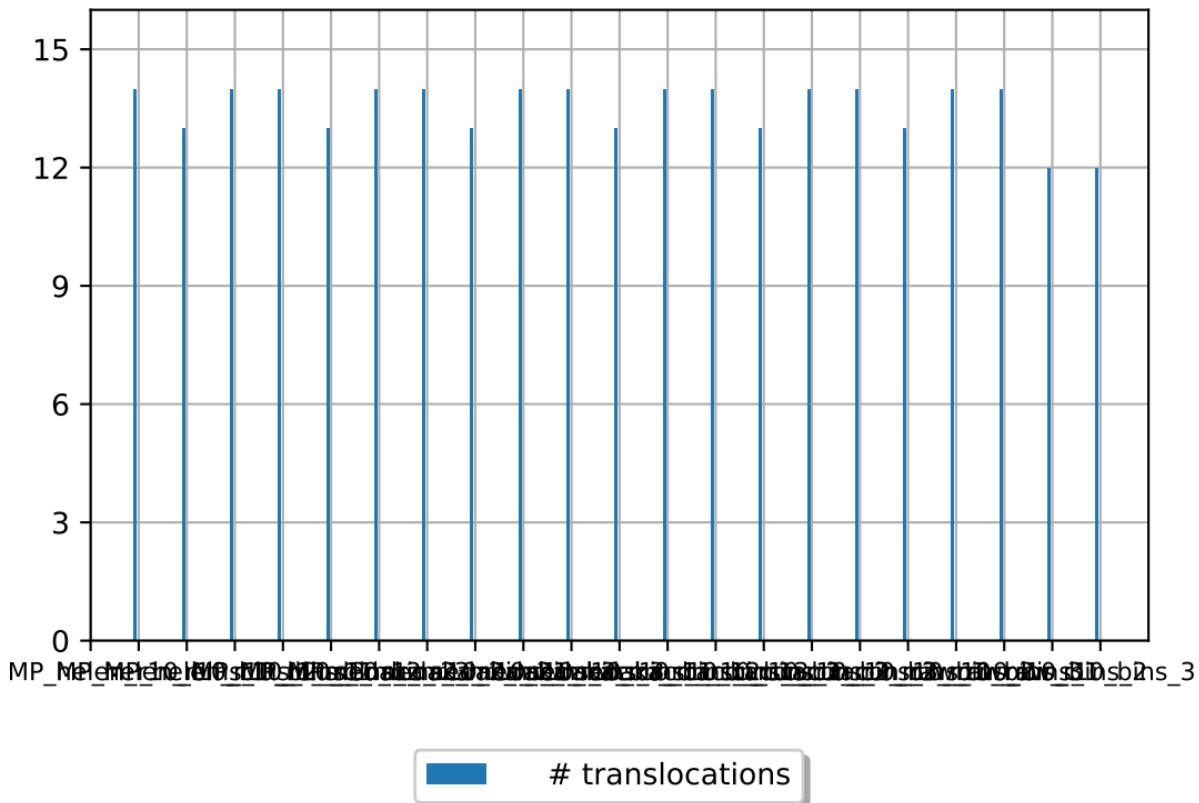
raw\_r10\_bins\_2 GC content



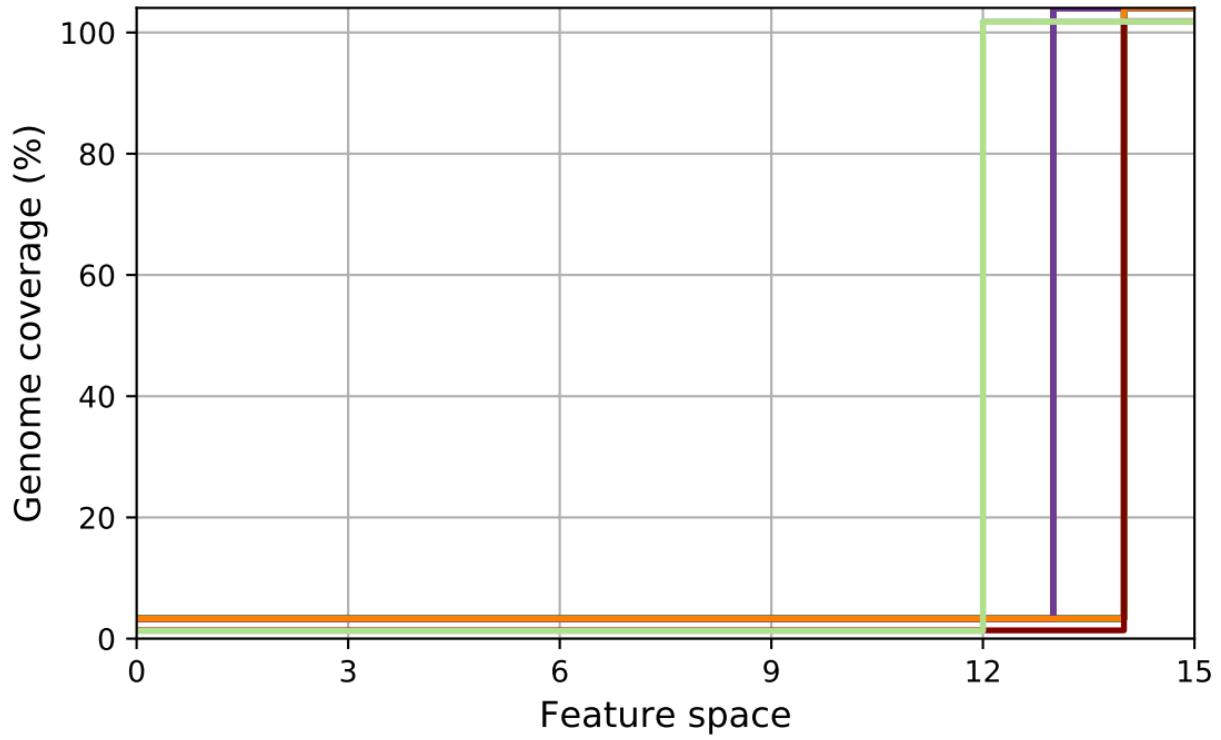
raw\_r10\_bins\_3 GC content



## Misassemblies



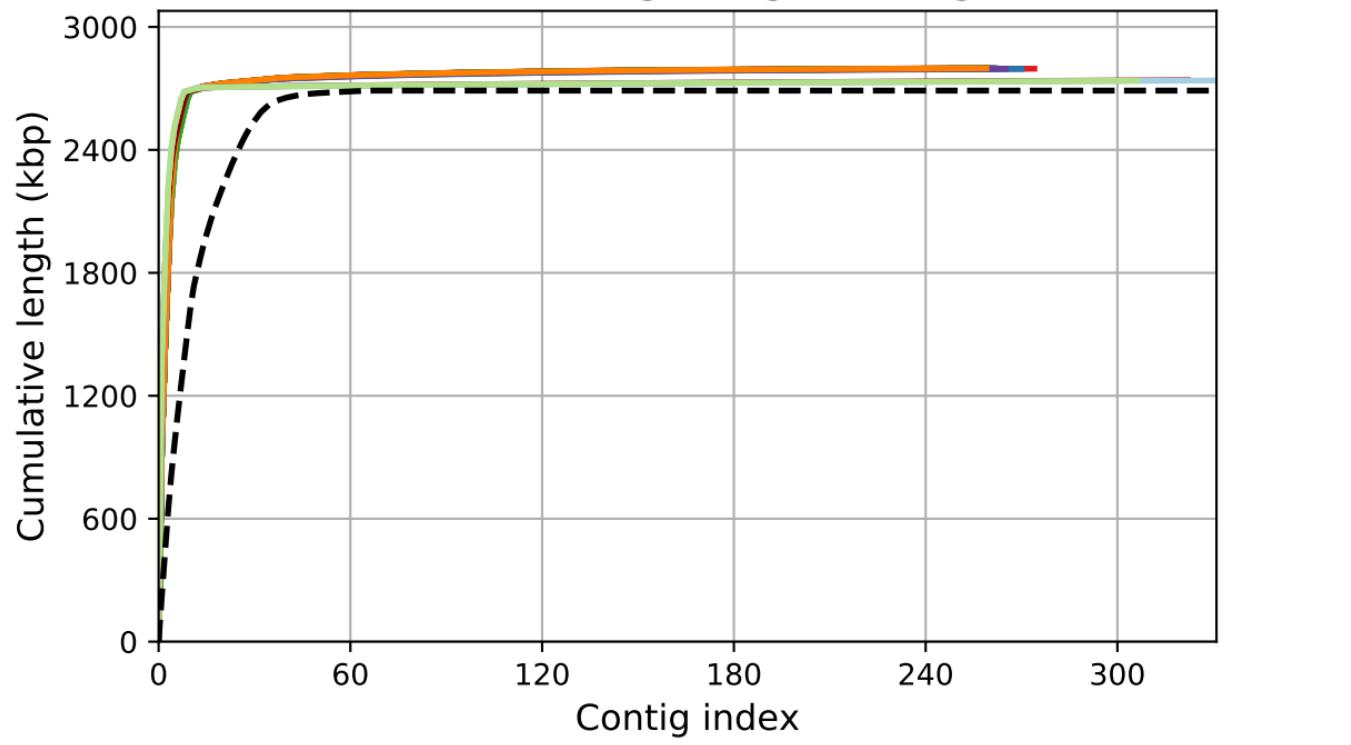
### FRCurve (misassemblies)



Legend:

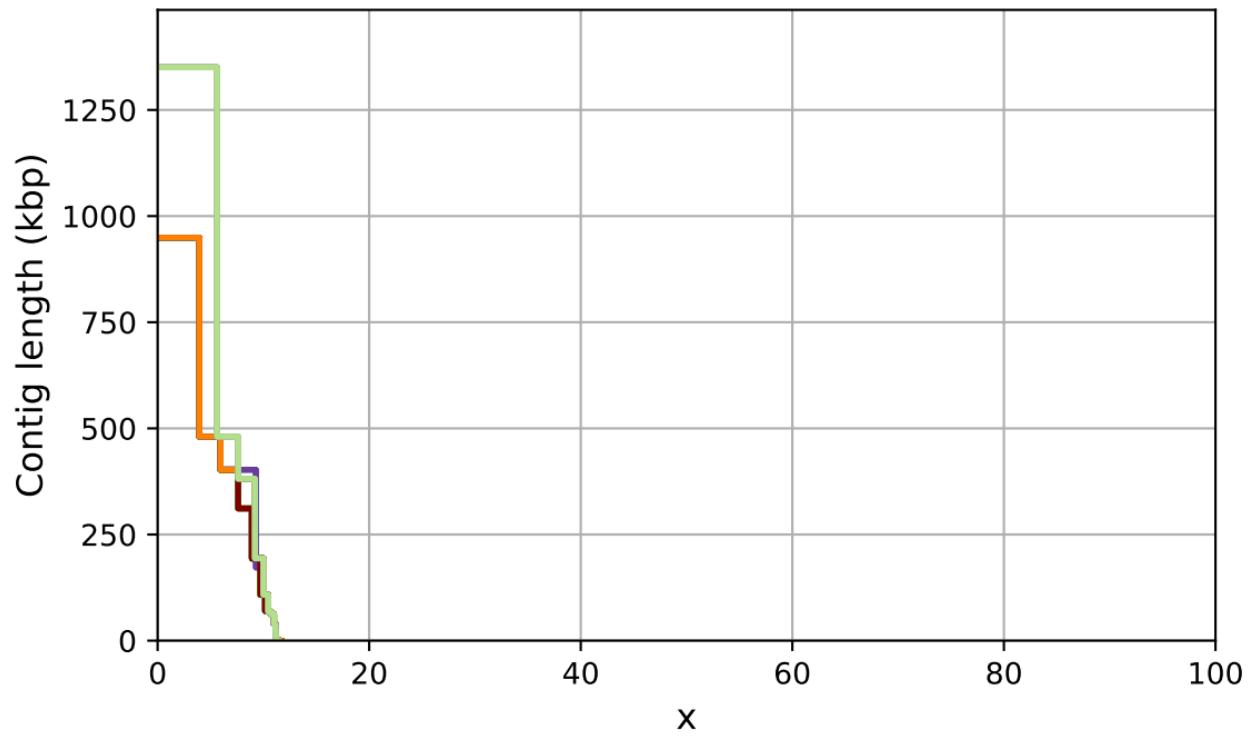
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins\_1
- r1\_medaka\_r10\_bins\_2
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- racon\_r1\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3

Cumulative length (aligned contigs)



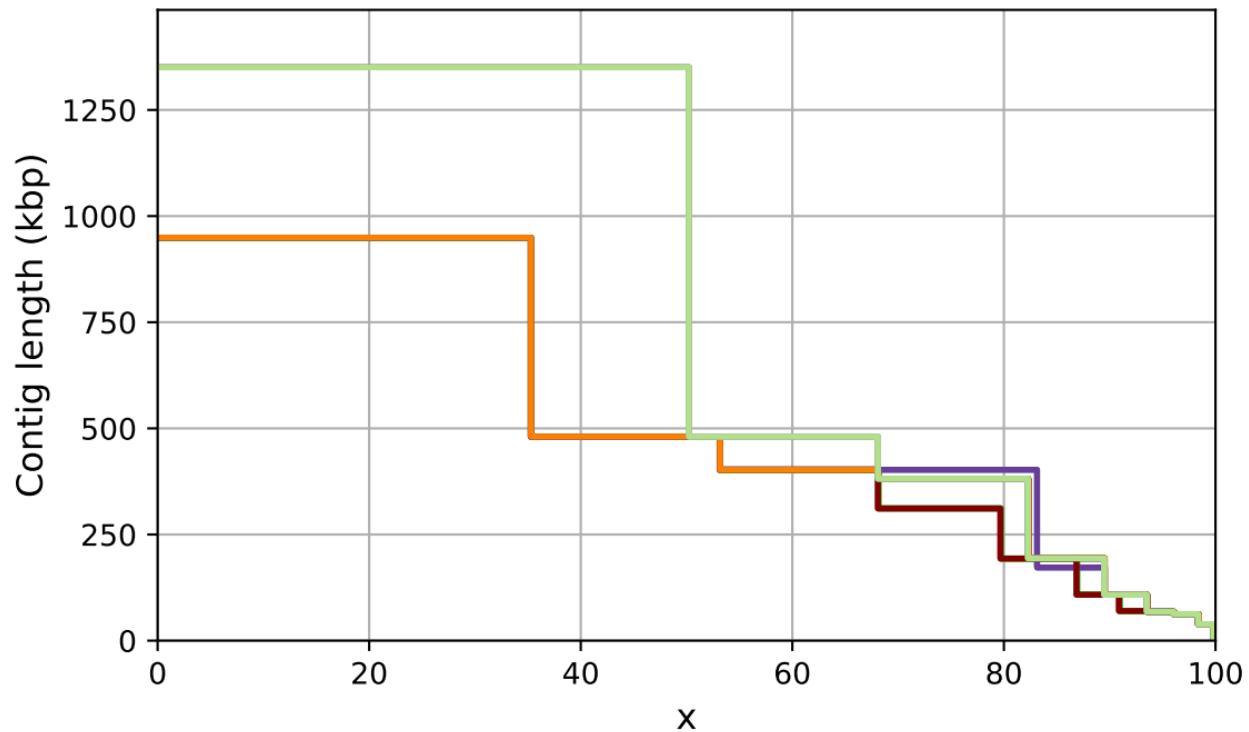
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins 1
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- r2\_medaka\_r10\_bins\_3
- raw r10 bins 1

# NAX



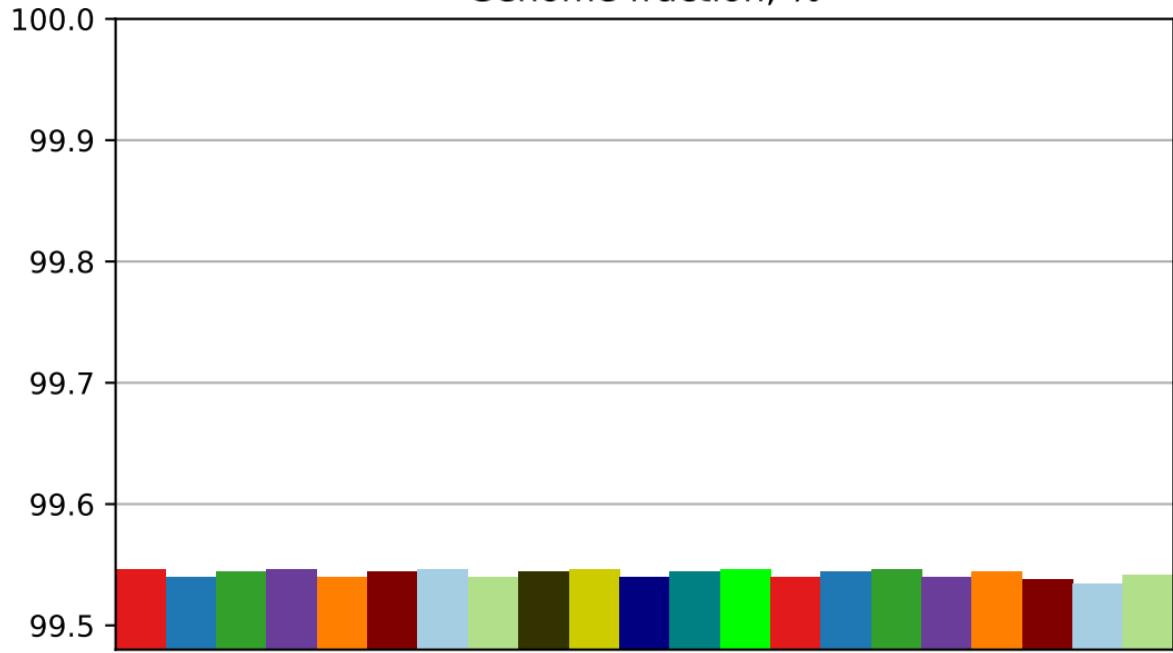
- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins 1
- r1\_medaka\_r10\_bins\_2
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- racon\_r1\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3

# NGAx



- MP\_helen\_r10\_bins\_1
- MP\_helen\_r10\_bins\_2
- MP\_helen\_r10\_bins\_3
- MP r10 bins 1
- r1\_medaka\_r10\_bins\_2
- r1\_medaka\_r10\_bins\_3
- r2\_medaka\_r10\_bins\_1
- r2\_medaka\_r10\_bins\_2
- racon\_r1\_r10\_bins\_3
- racon\_r2\_r10\_bins\_1
- racon\_r2\_r10\_bins\_2
- racon\_r2\_r10\_bins\_3

Genome fraction, %



- █ MP\_helen\_r10\_bins\_1
- █ MP\_helen\_r10\_bins\_2
- █ MP\_helen\_r10\_bins\_3
- █ MP r10 bins 1
- █ r1\_medaka\_r10\_bins\_2
- █ r1\_medaka\_r10\_bins\_3
- █ r2\_medaka\_r10\_bins\_1
- █ r2\_medaka\_r10\_bins\_2
- █ racon\_r1\_r10\_bins\_3
- █ racon\_r2\_r10\_bins\_1
- █ racon\_r2\_r10\_bins\_2
- █ racon\_r2\_r10\_bins\_3
- █ racon\_r2\_r10\_bins\_4