Field Run Report

Run Information

Run Name	R_2012_07_17_14_31_13_ FLX04070166_Administrator_797A_3_265_2_3_1A_1B_797A_1_265_2_1
Analysis Name	D_2012_07_18_09_54_01_brk07_fullProcessingAmplicons
Instr. SW Rev	2.6 (20110516_1007)
gsRunProcessor Rev.	2.6
Regions	8
Cycle Count	200
PTP ID	772491
PTP Type	70 x 75 , 35 uM
ICL Script Filename	200x_TACG_70x75_XLR70KIT.icl
Instr. Serial Number	FLX04070166
Instrument Model	GSFLX
Instrument Configuration	FLX+
User Bar Codes	064824490019387192002201405351588001938544201220120523654100193852320032013 064234180019385202007201305641870001938660200420130523368200193856120092013 053515960019387442004201305233534001938594200120137724910

Quality Filter Metrics

Control (ATGC)

Region	Raw Wells	Total Keypass (% raw)	ATGC Keypass	Passed Filter (% Keypass)	Avg Length	Total Bases	Dot Failed (% Keypass)	Mixed Failed (% Keypass)	ShortQuality (% Keypass)	Primer Failed (% Keypass)
1	221,814	217,802 (98.2 %)	1,494	78 (5.2 %)	532	41,517 bp	0 (.0 %)	4 (.3 %)	1,412 (94.5 %)	0 (.0 %)
2	133,455	130,681 (97.9 %)	2,754	834 (30.3 %)	517	431,297 bp	1 (.0 %)	1 (.0 %)	1,918 (69.6 %)	0 (.0 %)
3	129,602	126,902 (97.9 %)	3,559	1,170 (32.9 %)	516	603,251 bp	0 (.0 %)	1 (.0 %)	2,387 (67.1 %)	0 (.0 %)
4	142,151	138,211 (97.2 %)	4,135	1,066 (25.8 %)	511	544,897 bp	1 (.0 %)	7 (.2 %)	3,061 (74.0 %)	0 (.0 %)
5	202,049	197,913 (98.0 %)	1,714	513 (29.9 %)	525	269,527 bp	5 (.3 %)	2 (.1 %)	1,193 (69.6 %)	0 (.0 %)
6	94,253	91,164 (96.7 %)	4,393	2,944 (67.0 %)	519	1,528,786 bp	2 (.0 %)	2 (.0 %)	1,445 (32.9 %)	0 (.0 %)
7	144,069	141,249 (98.0 %)	4,478	1,573 (35.1 %)	515	809,904 bp	0 (.0 %)	2 (.0 %)	2,903 (64.8 %)	0 (.0 %)
8	152,544	150,064 (98.4 %)	4,360	782 (17.9 %)	522	408,426 bp	1 (.0 %)	3 (.1 %)	3,574 (82.0 %)	0 (.0 %)
TOTAL	**1,219,937**	1,193,986 (97.9 %)	**26,887**	**8,960**		**4,637,605 bp**	**10**	**22**	**17,893**	**0**

Control (CATG)

Region	Raw Wells	Total Keypass (% raw)	CATG Keypass	Passed Filter (% Keypass)	Avg Length	Total Bases	Dot Failed (% Keypass)	Mixed Failed (% Keypass)	ShortQuality (% Keypass)	Primer Failed (% Keypass)
1	221,814	217,802 (98.2 %)	819	12 (1.5 %)	506	6,067 bp	0 (.0 %)	101 (12.3 %)	702 (85.7 %)	0 (.0 %)
2	133,455	130,681 (97.9 %)	1,992	156 (7.8 %)	507	79,075 bp	0 (.0 %)	17 (.9 %)	1,818 (91.3 %)	0 (.0 %)
3	129,602	126,902 (97.9 %)	2,610	174 (6.7 %)	505	87,805 bp	0 (.0 %)	40 (1.5 %)	2,392 (91.6 %)	0 (.0 %)
4	142,151	138,211 (97.2 %)	3,005	173 (5.8 %)	500	86,434 bp	2 (.1 %)	147 (4.9 %)	2,657 (88.4 %)	0 (.0 %)
5	202,049	197,913 (98.0 %)	1,110	44 (4.0 %)	503	22,125 bp	2 (.2 %)	30 (2.7 %)	1,029 (92.7 %)	0 (.0 %)
6	94,253	91,164 (96.7 %)	3,033	536 (17.7 %)	512	274,466 bp	1 (.0 %)	5 (.2 %)	2,487 (82.0 %)	0 (.0 %)
7	144,069	141,249 (98.0 %)	3,051	136 (4.5 %)	503	68,381 bp	2 (.1 %)	17 (.6 %)	2,890 (94.7 %)	0 (.0 %)
8	152,544	150,064 (98.4 %)	2,752	64 (2.3 %)	480	30,730 bp	10 (.4 %)	76 (2.8 %)	2,582 (93.8 %)	0 (.0 %)
TOTAL	**1,219,937**	1,193,986 (97.9 %)	**18,372**	**1,295**		**655,083 bp**	**17**	**433**	**16,557**	**0**

Library (GACT)

Region	Raw Wells	Total Keypass (% raw)	GACT Keypass	Passed Filter (% Keypass)	Avg Length	Total Bases	Dot Failed (% Keypass)	Mixed Failed (% Keypass)	ShortQuality (% Keypass)	Primer Failed (% Keypass)
1	221,814	217,802 (98.2 %)	215,489	2,365 (1.1 %)	178	421,723 bp	4,526 (2.1 %)	41,275 (19.2 %)	166,002 (77.0 %)	325 (.2 %)
5	202,049	197,913 (98.0 %)	195,089	3,600 (1.8 %)	255	916,271 bp	4,817 (2.5 %)	44,242 (22.7 %)	141,157 (72.4 %)	206 (.1 %)
TOTAL	**1,219,937**	1,193,986 (97.9 %)	**410,578**	**5,965**		**1,337,994 bp**	**9,343**	**85,517**	**307,159**	**531**

Library (TCAG)

Region	Raw Wells	Total Keypass (% raw)	TCAG Keypass	Passed Filter (% Keypass)	Avg Length	Total Bases	Dot Failed (% Keypass)	Mixed Failed (% Keypass)	ShortQuality (% Keypass)	Primer Failed (% Keypass)
2	133,455	130,681 (97.9 %)	125,935	20,658 (16.4 %)	416	8,588,708 bp	1,249 (1.0 %)	30,605 (24.3 %)	73,208 (58.1 %)	0 (.0 %)
3	129,602	126,902 (97.9 %)	120,733	3,458 (2.9 %)	426	1,472,732 bp	129 (.1 %)	8,881 (7.4 %)	108,042 (89.5 %)	0 (.0 %)
4	142,151	138,211 (97.2 %)	131,071	5,351 (4.1 %)	404	2,162,183 bp	825 (.6 %)	11,686 (8.9 %)	112,950 (86.2 %)	0 (.0 %)
6	94,253	91,164 (96.7 %)	83,738	27,553 (32.9 %)	417	11,501,720 bp	762 (.9 %)	13,665 (16.3 %)	41,613 (49.7 %)	1 (.0 %)
7	144,069	141,249 (98.0 %)	133,720	10,102 (7.6 %)	430	4,343,711 bp	207 (.2 %)	7,333 (5.5 %)	115,792 (86.6 %)	0 (.0 %)
8	152,544	150,064 (98.4 %)	142,952	3,142 (2.2 %)	340	1,068,787 bp	911 (.6 %)	12,217 (8.5 %)	126,395 (88.4 %)	0 (.0 %)
TOTAL	**1,219,937**	1,193,986 (97.9 %)	**738,149**	**70,264**		**29,137,841 bp**	**4,083**	**84,387**	**578,000**	**1**

Control Fragment Metrics

ATGC

Region	Raw Wells	Keypass	Percentage of Fragments
			98% Accuracy
1	221814	1494	56.6 % @ 400 bp 94.3 % @ 300 bp 98.1 % @ 200 bp 98.5 % @ 100 bp
2	133455	2754	74.4 % @ 400 bp 95.4 % @ 300 bp 98.9 % @ 200 bp 99.6 % @ 100 bp
3	129602	3559	80.6 % @ 400 bp 97.0 % @ 300 bp 98.9 % @ 200 bp 99.2 % @ 100 bp
4	142151	4135	72.5 % @ 400 bp 91.0 % @ 300 bp 95.5 % @ 200 bp 97.3 % @ 100 bp
5	202049	1714	87.8 % @ 400 bp 97.1 % @ 300 bp 98.0 % @ 200 bp 98.6 % @ 100 bp
6	94253	4393	88.8 % @ 400 bp 97.1 % @ 300 bp 99.0 % @ 200 bp 99.6 % @ 100 bp
7	144069	4478	79.0 % @ 400 bp 96.0 % @ 300 bp 98.9 % @ 200 bp 99.5 % @ 100 bp
8	152544	4360	62.9 % @ 400 bp 90.8 % @ 300 bp 98.2 % @ 200 bp 98.9 % @ 100 bp

CATG

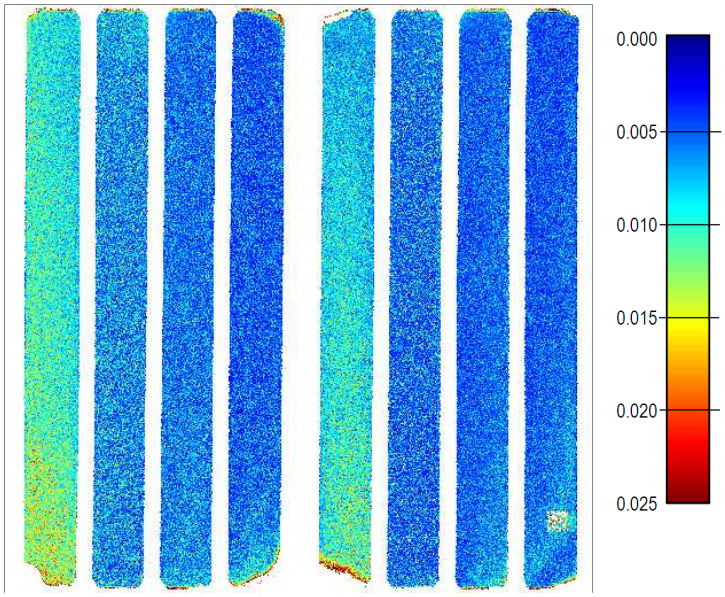
Region	Raw Wells	Keypass	Percentage of Fragments
			98% Accuracy
1	221814	819	13.1 % @ 400 bp 33.0 % @ 300 bp 51.0 % @ 200 bp 59.5 % @ 100 bp
2	133455	1992	32.0 % @ 400 bp 59.8 % @ 300 bp 82.8 % @ 200 bp 90.2 % @ 100 bp
3	129602	2610	35.0 % @ 400 bp 61.5 % @ 300 bp 81.1 % @ 200 bp 87.0 % @ 100 bp
4	142151	3005	30.4 % @ 400 bp 53.7 % @ 300 bp 71.7 % @ 200 bp 77.4 % @ 100 bp
5	202049	1110	24.1 % @ 400 bp 50.5 % @ 300 bp 70.3 % @ 200 bp 79.3 % @ 100 bp
6	94253	3033	48.7 % @ 400 bp 72.8 % @ 300 bp 89.8 % @ 200 bp 93.3 % @ 100 bp
7	144069	3051	31.7 % @ 400 bp 59.8 % @ 300 bp 81.6 % @ 200 bp 88.2 % @ 100 bp
8	152544	2752	19.2 % @ 400 bp 45.5 % @ 300 bp 71.3 % @ 200 bp 81.6 % @ 100 bp

Combined

Region	Raw Wells	Keypass	Percentage of Fragments
			98% Accuracy
1	221814	2313	41.2 % @ 400 bp 72.6 % @ 300 bp 81.5 % @ 200 bp 84.7 % @ 100 bp
2	133455	4746	56.6 % @ 400 bp 80.5 % @ 300 bp 92.2 % @ 200 bp 95.7 % @ 100 bp
3	129602	6169	61.3 % @ 400 bp 82.0 % @ 300 bp 91.4 % @ 200 bp 94.0 % @ 100 bp
4	142151	7140	54.8 % @ 400 bp 75.3 % @ 300 bp 85.5 % @ 200 bp 88.9 % @ 100 bp
5	202049	2824	62.8 % @ 400 bp 78.8 % @ 300 bp 87.1 % @ 200 bp 91.0 % @ 100 bp
6	94253	7426	72.4 % @ 400 bp 87.2 % @ 300 bp 95.2 % @ 200 bp 97.0 % @ 100 bp
7	144069	7529	59.8 % @ 400 bp 81.4 % @ 300 bp 91.9 % @ 200 bp 94.9 % @ 100 bp
8	152544	7112	46.0 % @ 400 bp 73.3 % @ 300 bp 87.8 % @ 200 bp 92.2 % @ 100 bp

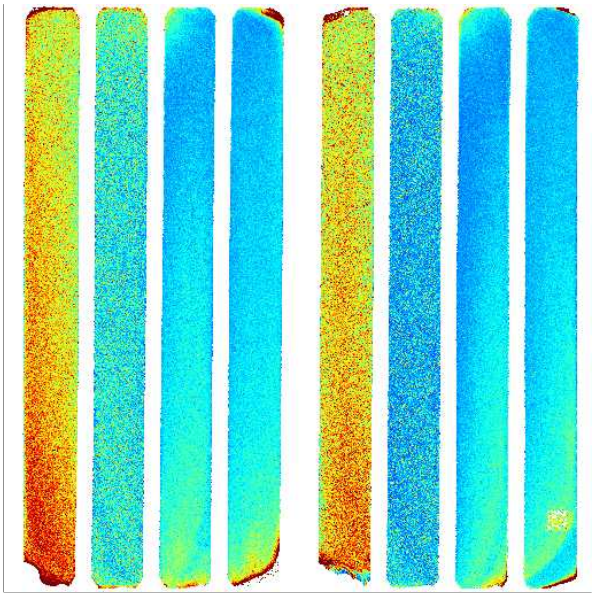
Carry Forward / Incomplete Extension Corrections

Carry Forward



Region	Median	Mean	Untrainable % Keypass
1	.00782	.01044 ± .00738	39793 18.3 %
2	.00425	.00713 ± .00641	30215 23.1 %
3	.00371	.00622 ± .00572	1958 1.5 %
4	.00316	.00569 ± .00592	3258 2.4 %
5	.00631	.00906 ± .00690	45030 22.8 %
6	.00412	.00622 ± .00542	14128 15.4 %
7	.00357	.00574 ± .00517	1948 1.4 %
8	.00343	.00564 ± .00535	3005 2.0 %

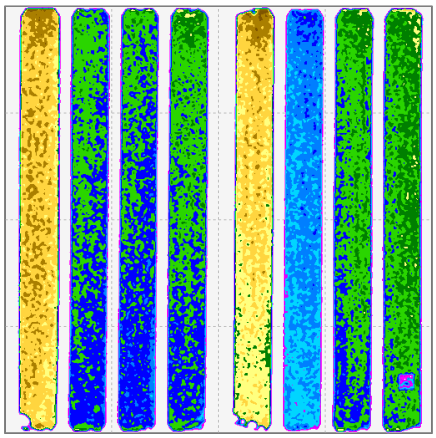
Incomplete Extension



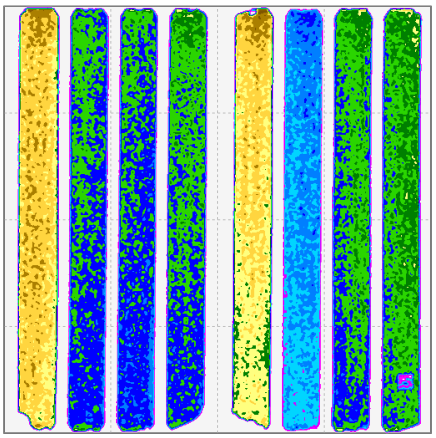
Region	Median	Mean	Untrainable % Keypass
1	.00294	.00335 ± .00158	2639 1.2 %
2	.00145	.00197 ± .00132	370 .3 %
3	.00165	.00174 ± .00054	346 .3 %
4	.00165	.00176 ± .00068	359 .3 %
5	.00267	.00324 ± .00180	3559 1.8 %
6	.00125	.00162 ± .00114	820 .9 %
7	.00153	.00162 ± .00053	368 .3 %
8	.00161	.00170 ± .00058	355 .2 %

Bead Loading Density

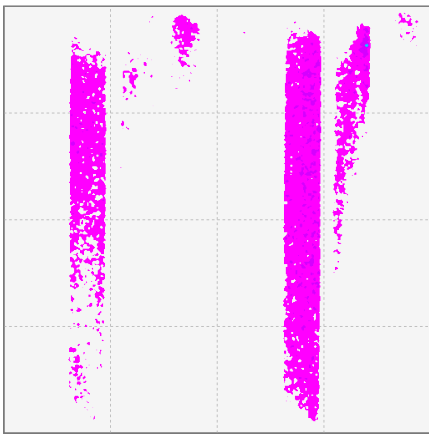
Raw Wells Only

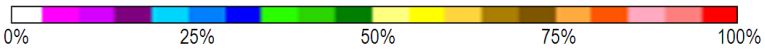


Keypass Wells Only



High Quality Wells





Signal Per Base / ATP Values

Control

ATGC

Region	Raw Wells	Keypass	Sig / Base	ATP 1	ATP 2	ATP 3
1	221814	1494	185 ± 39	154 ± 27	90 ± 12	73 ± 11
2	133455	2754	182 ± 48	171 ± 30	101 ± 15	82 ± 14
3	129602	3559	164 ± 43	175 ± 37	107 ± 15	87 ± 13
4	142151	4135	160 ± 42	155 ± 40	99 ± 19	81 ± 16
5	202049	1714	220 ± 51	195 ± 43	98 ± 16	79 ± 13
6	94253	4393	186 ± 60	212 ± 43	117 ± 18	96 ± 16
7	144069	4478	175 ± 48	196 ± 42	106 ± 18	86 ± 15
8	152544	4360	163 ± 44	185 ± 40	100 ± 19	81 ± 16

CATG

Region	Raw Wells	Keypass	Sig / Base	ATP 1	ATP 2	ATP 3
1	221814	819	144 ± 33	154 ± 27	93 ± 12	78 ± 11
2	133455	1992	137 ± 38	174 ± 31	105 ± 15	87 ± 14
3	129602	2610	126 ± 36	180 ± 37	112 ± 16	94 ± 14
4	142151	3005	124 ± 33	159 ± 41	104 ± 20	87 ± 17
5	202049	1110	165 ± 37	196 ± 42	102 ± 17	85 ± 14
6	94253	3033	142 ± 40	217 ± 44	121 ± 19	102 ± 17
7	144069	3051	134 ± 38	201 ± 42	110 ± 18	92 ± 16
8	152544	2752	124 ± 34	190 ± 41	105 ± 20	87 ± 17

Library

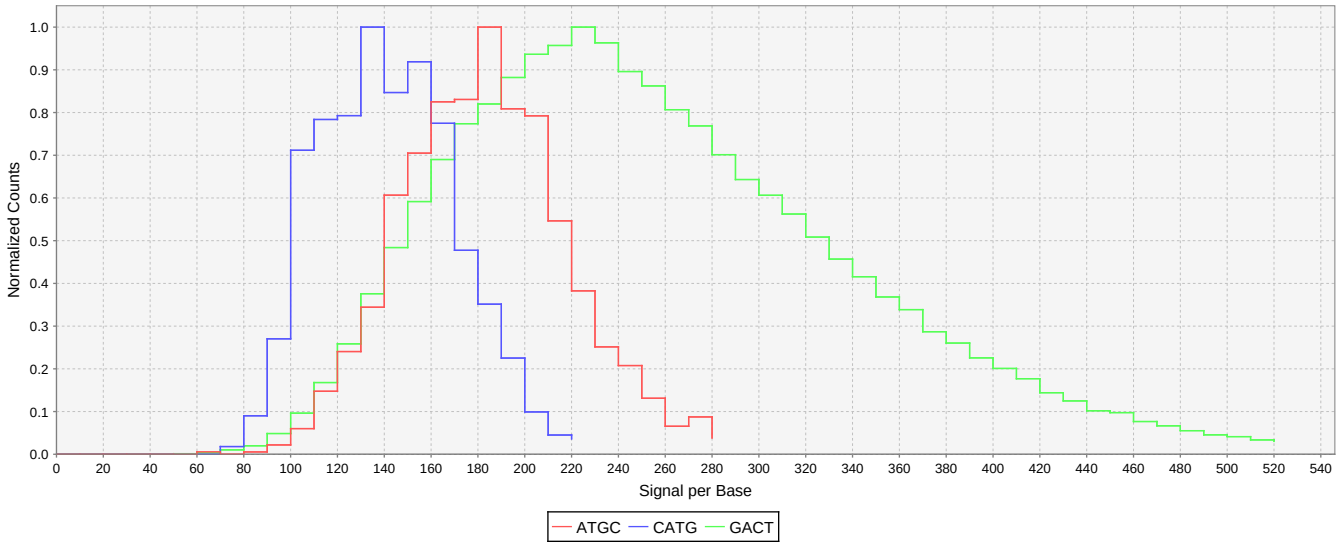
GACT

Region	Raw Wells	Keypass	Sig / Base	ATP 1	ATP 2	ATP 3
1	221814	215489	260 ± 91	148 ± 27	82 ± 12	68 ± 11
5	202049	195089	299 ± 111	187 ± 40	89 ± 15	75 ± 13

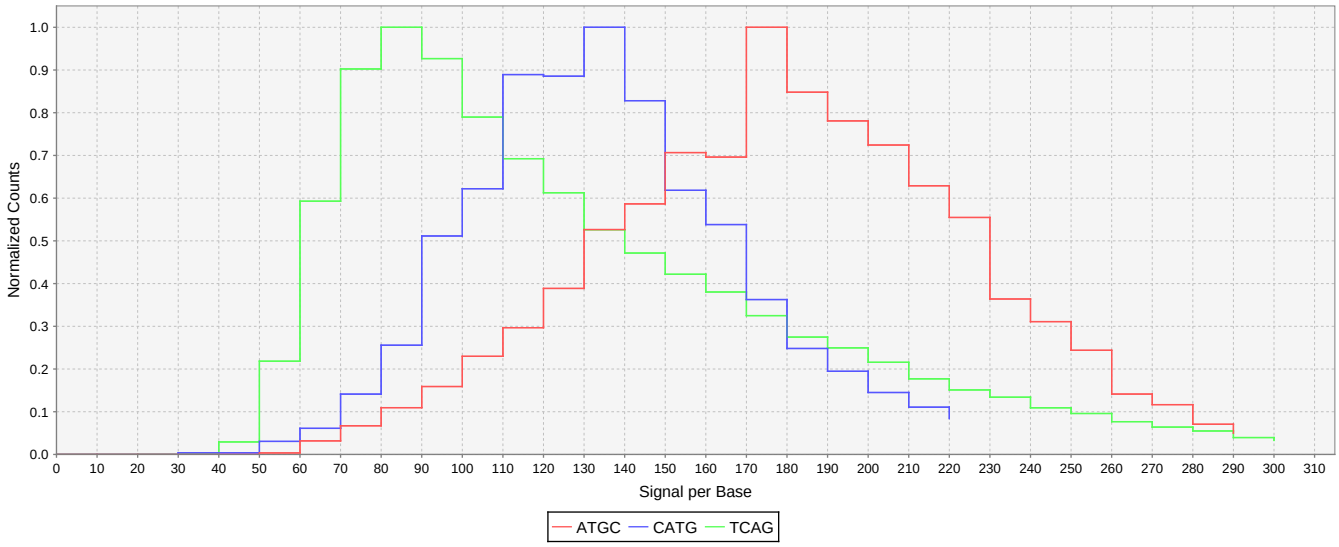
TCAG

Region	Raw Wells	Keypass	Sig / Base	ATP 1	ATP 2	ATP 3
2	133455	125935	131 ± 59	168 ± 29	99 ± 15	82 ± 14
3	129602	120733	116 ± 51	175 ± 36	105 ± 15	88 ± 14
4	142151	131071	138 ± 60	155 ± 39	97 ± 19	81 ± 16
6	94253	83738	121 ± 64	214 ± 42	117 ± 18	99 ± 17
7	144069	133720	126 ± 54	195 ± 42	104 ± 18	86 ± 16
8	152544	142952	146 ± 61	183 ± 40	97 ± 19	79 ± 17

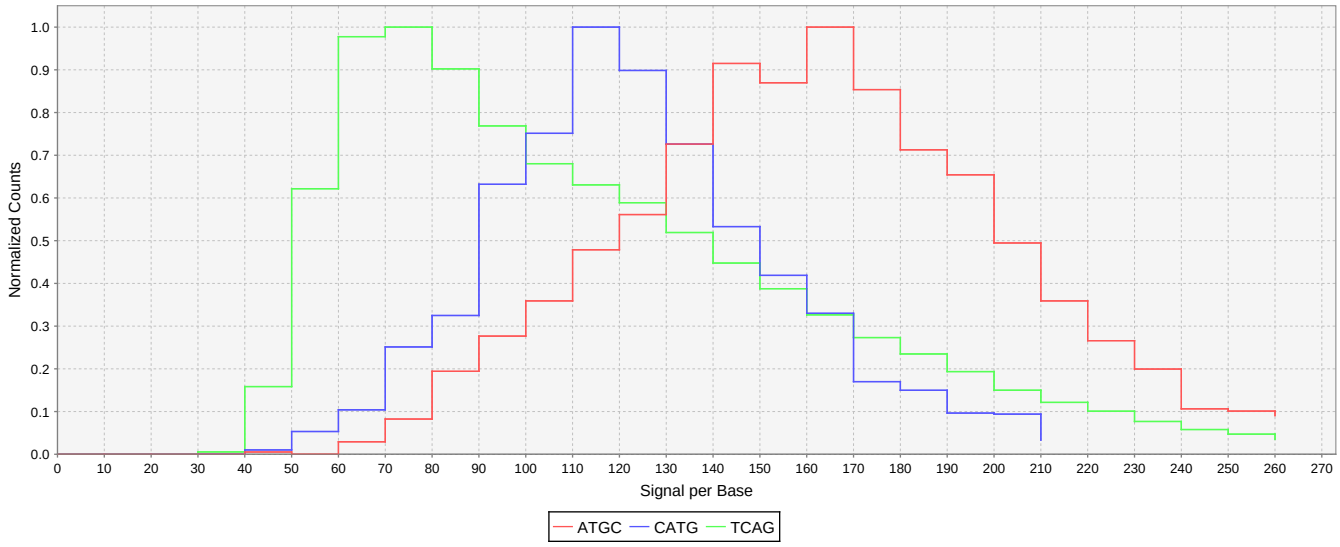
Region 1



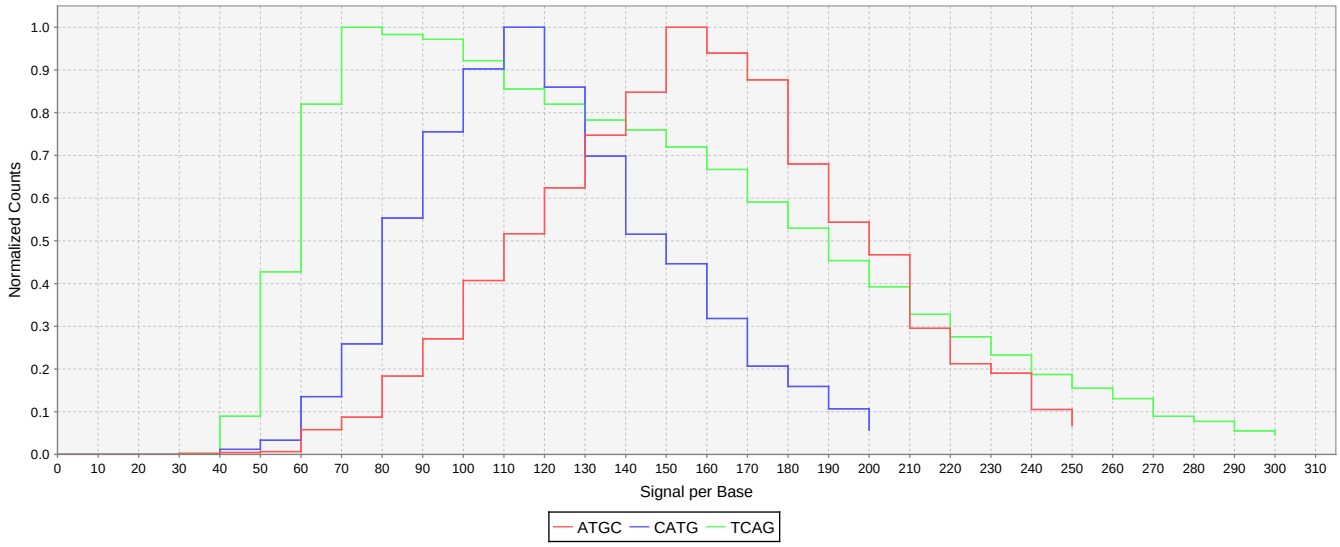
Region 2



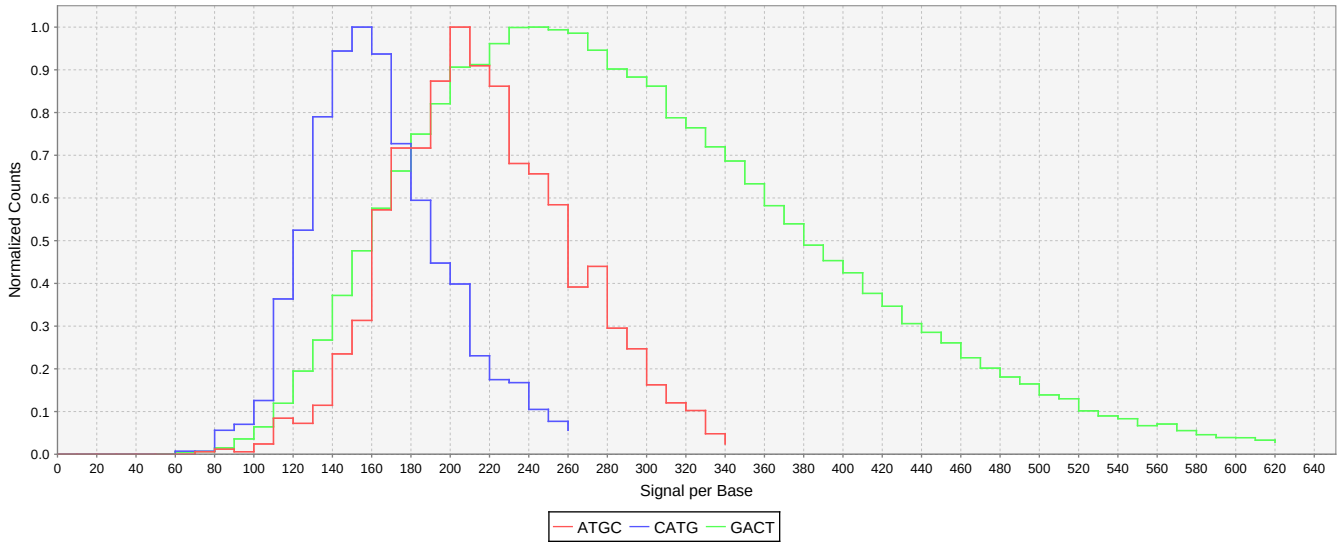
Region 3



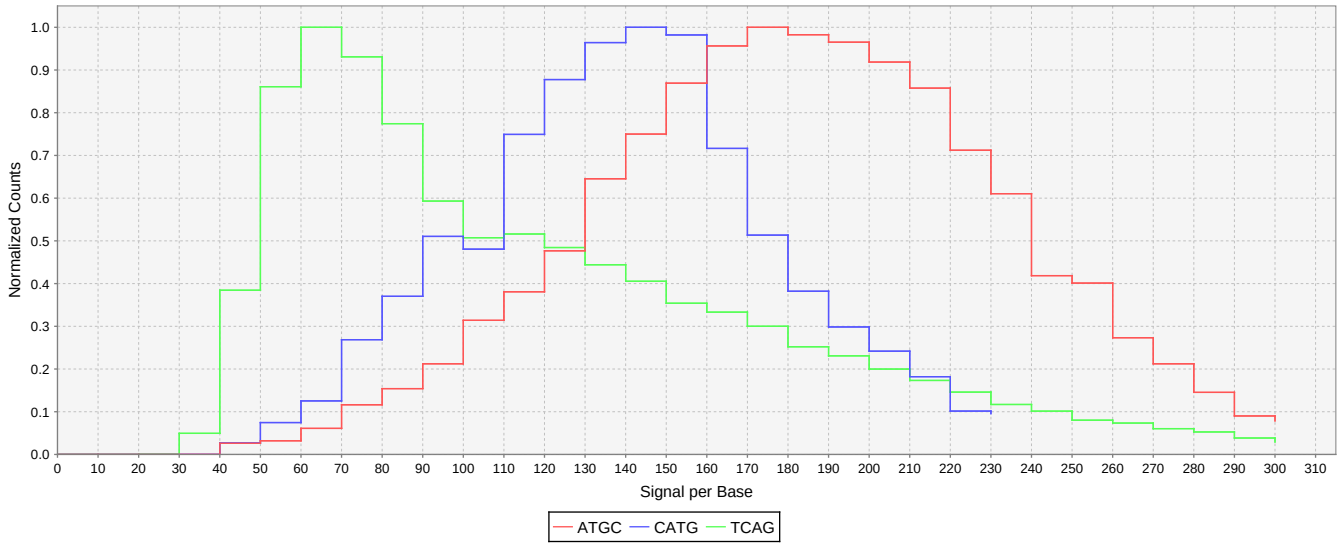
Region 4



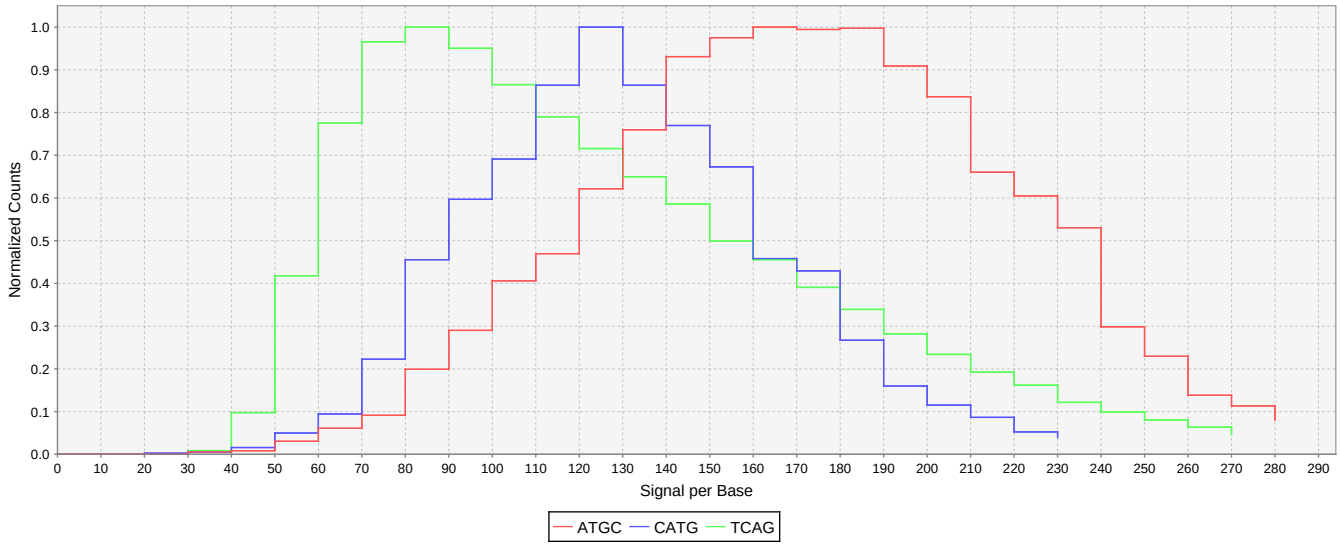
Region 5



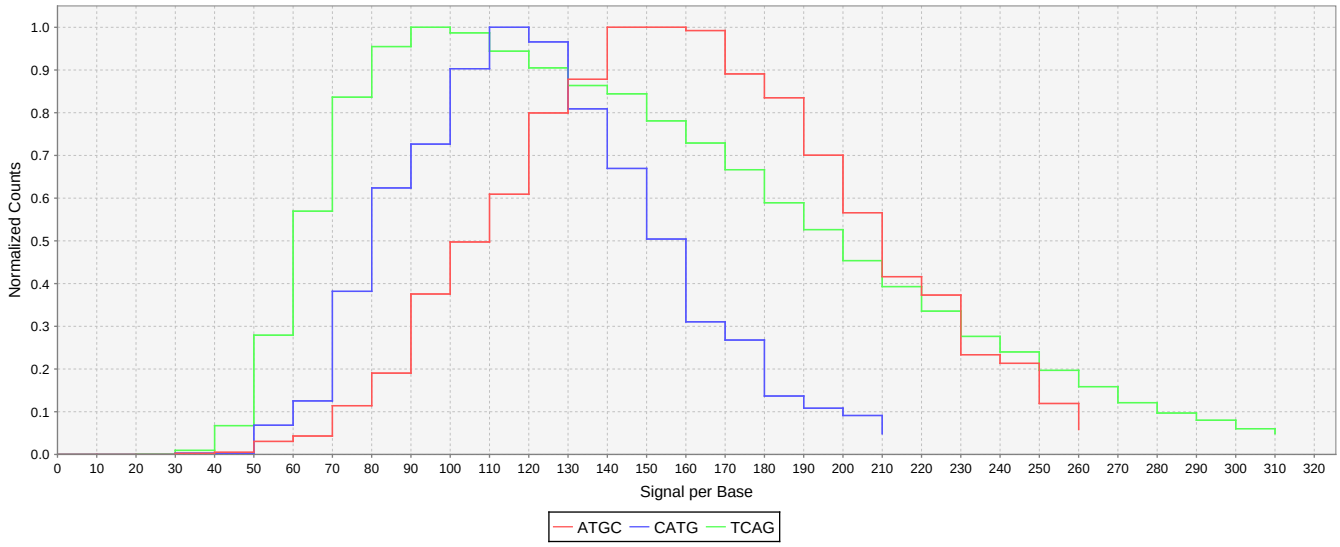
Region 6



Region 7



Region 8



Test Fragment Flowgrams

Summary

ECTF

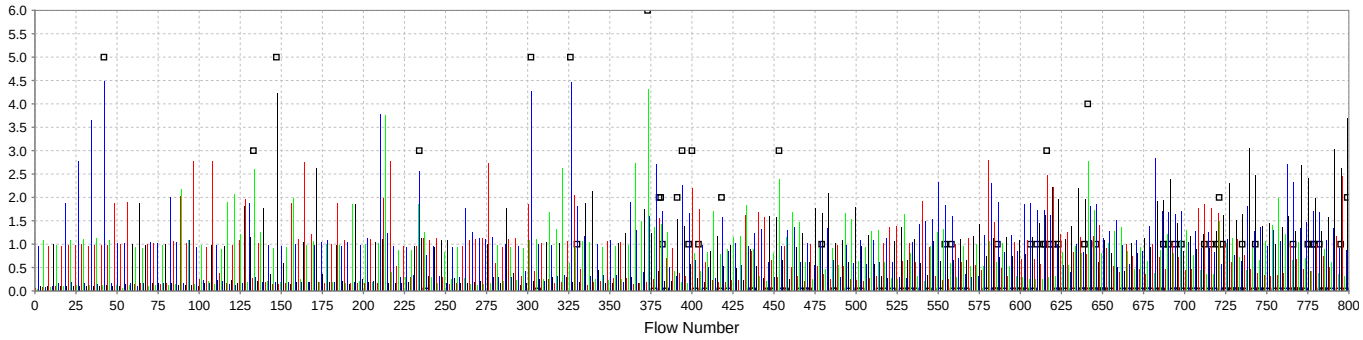
Region	ECTF301		ECTF302		ECTF303		ECTF304	
	# Matches	Error	# Matches	Error	# Matches	Error	# Matches	Error
1	309	0.35%	471	1.33%	353	0.64%	370	0.3%
2	717	0.07%	687	0.18%	704	0.06%	648	0.02%
3	934	0.03%	931	0.25%	910	0.1%	780	0.07%
4	1088	0.22%	1046	1.04%	1048	0.27%	957	0.09%
5	377	0.4%	530	0.6%	433	0.46%	378	0.21%
6	1107	0.08%	1166	0.15%	1150	0.08%	975	0.16%
7	1166	0.08%	1182	0.2%	1126	0.13%	1008	0.09%
8	1139	0.08%	1167	0.42%	1070	0.27%	987	0.16%
Total	6837	0.12%	7180	0.46%	6794	0.2%	6103	0.12%

AVTF

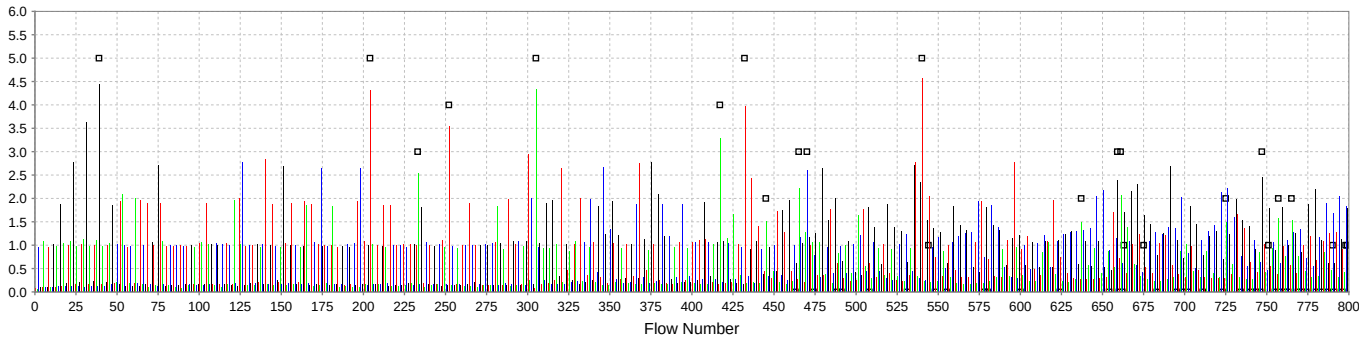
Region	AVTF100		AVTF120		AVTF150		AVTF2		AVTF7		AVTF90	
	# Matches	Error	# Matches	Error	# Matches	Error	# Matches	Error	# Matches	Error	# Matches	Error
1	117	9.6%	89	6.82%	63	7.48%	120	5.09%	177	3.81%	232	18.43%
2	335	4.17%	300	2.79%	359	2.84%	352	1.31%	343	2.33%	301	9.18%
3	460	2.72%	396	1.54%	434	2.21%	458	1.15%	428	2.05%	424	10.71%
4	523	4.25%	460	2.14%	440	2.84%	530	2.12%	494	2.2%	431	13.87%
5	189	7.61%	155	4.54%	98	6.87%	173	2.95%	219	2.62%	285	9.95%
6	528	2.28%	479	1.38%	519	1.55%	506	0.92%	512	2.0%	488	3.78%
7	565	3.09%	502	1.56%	467	2.27%	522	1.22%	495	2.13%	494	7.27%
8	457	4.81%	473	2.65%	449	3.19%	429	1.83%	440	2.75%	469	11.49%
Total	3174	3.96%	2854	2.26%	2829	2.71%	3090	1.66%	3108	2.35%	3124	9.99%

All Regions

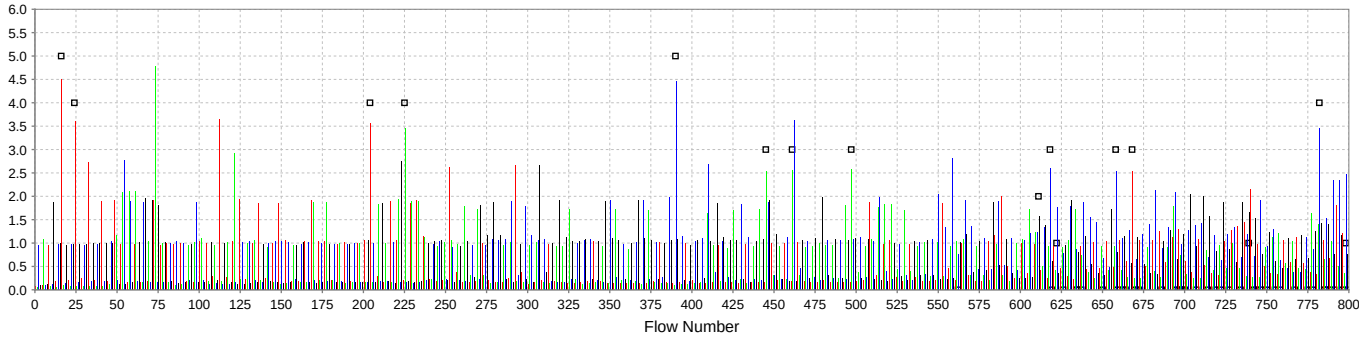
AVTF100



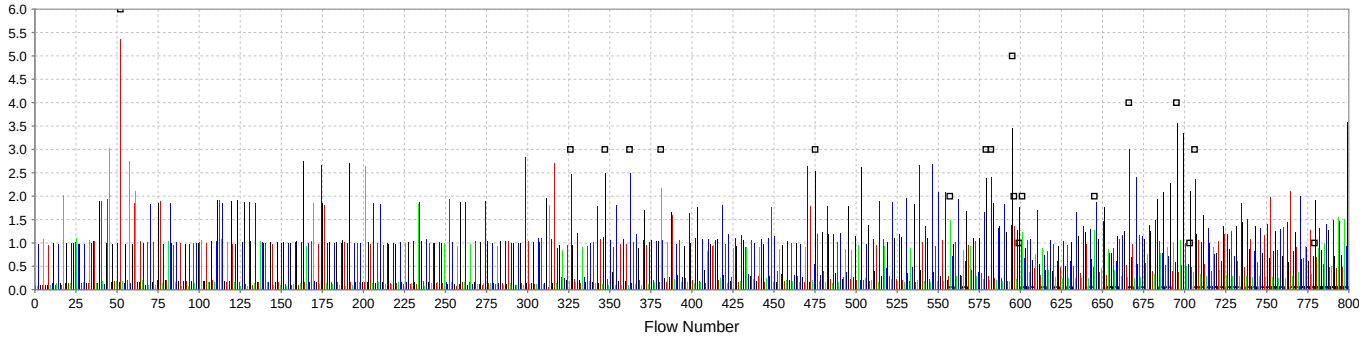
AVTF120



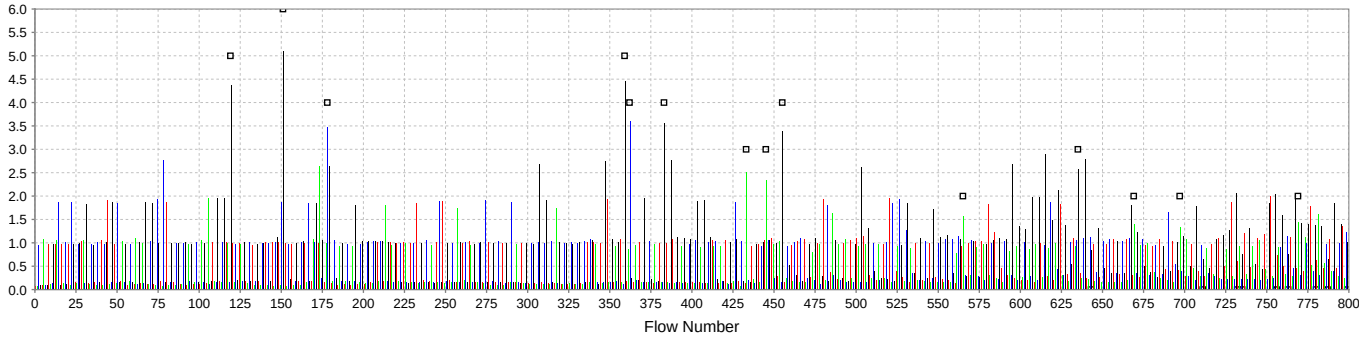
AVTF150



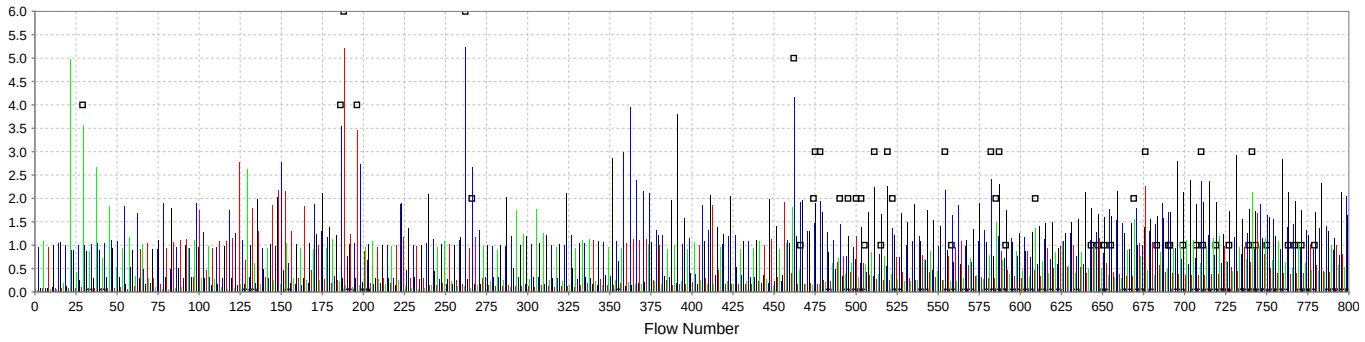
AVTF2



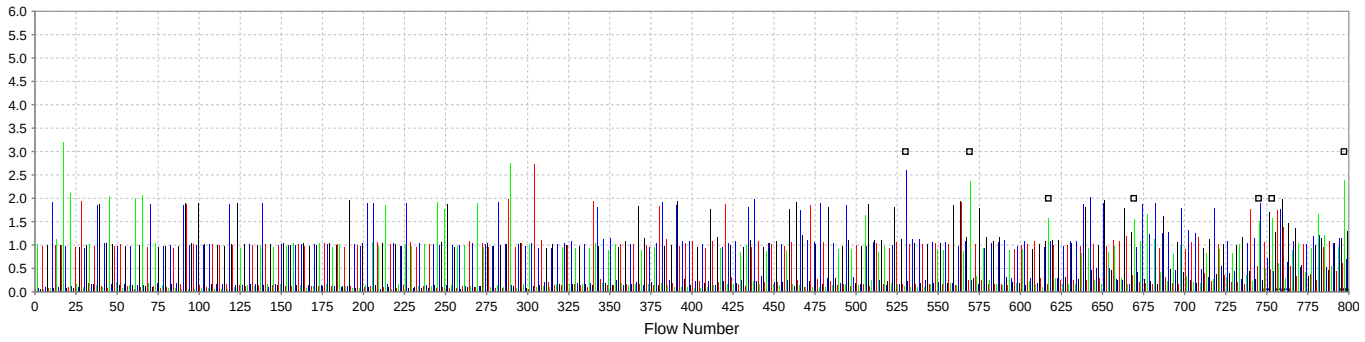
AVTF7



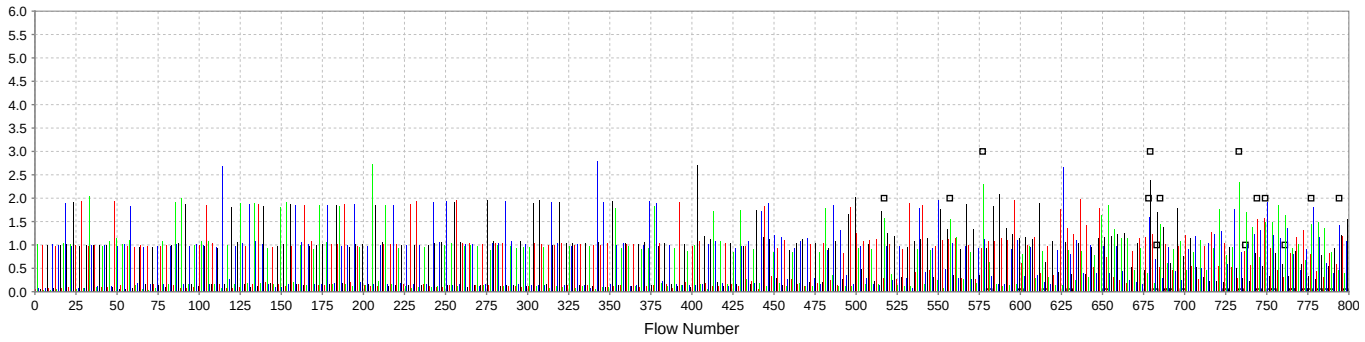
AVTF90



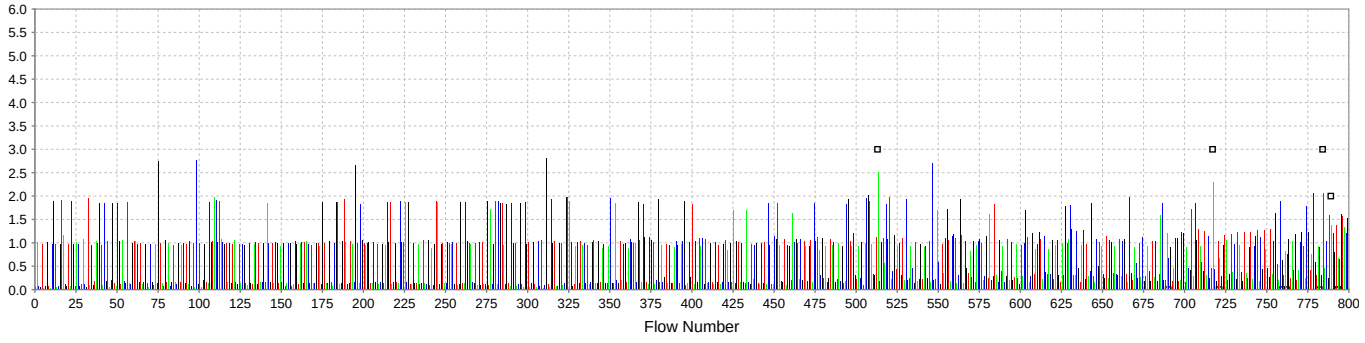
ECTF301



ECTF302



ECTF303



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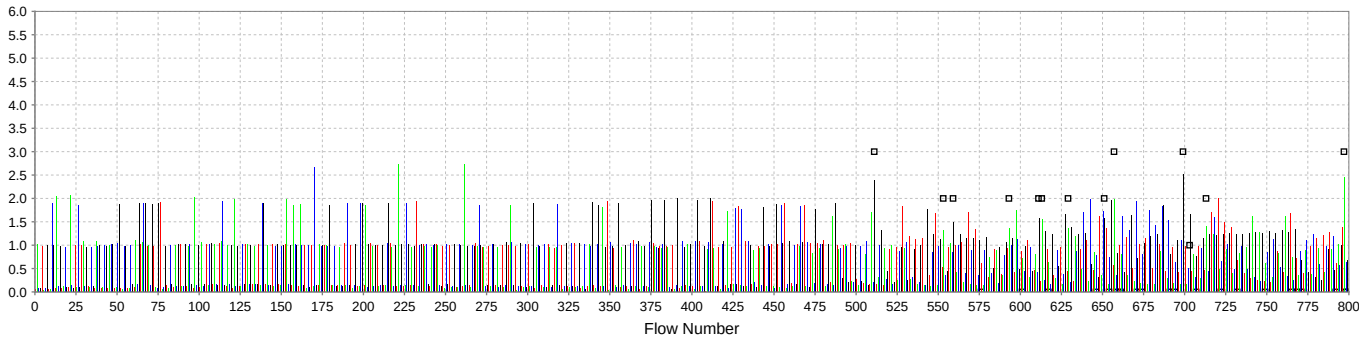
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Flow Number

Region 1

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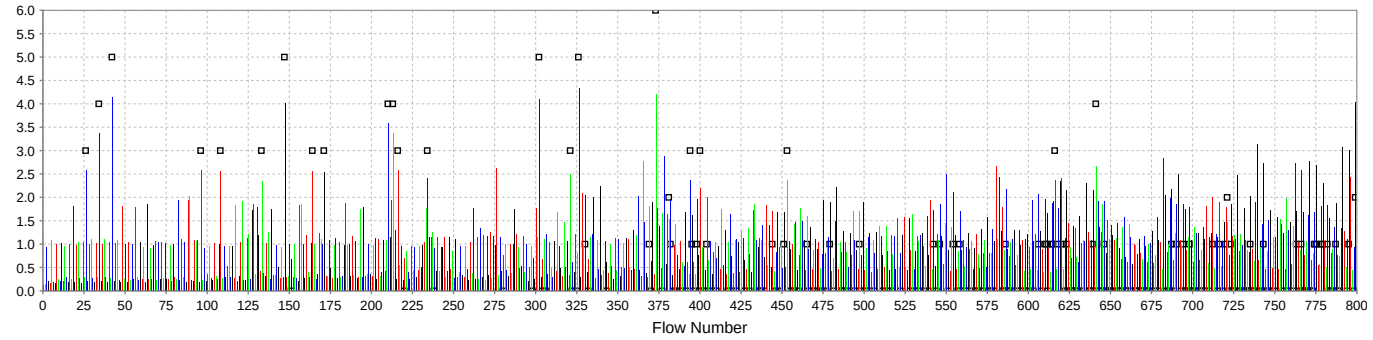

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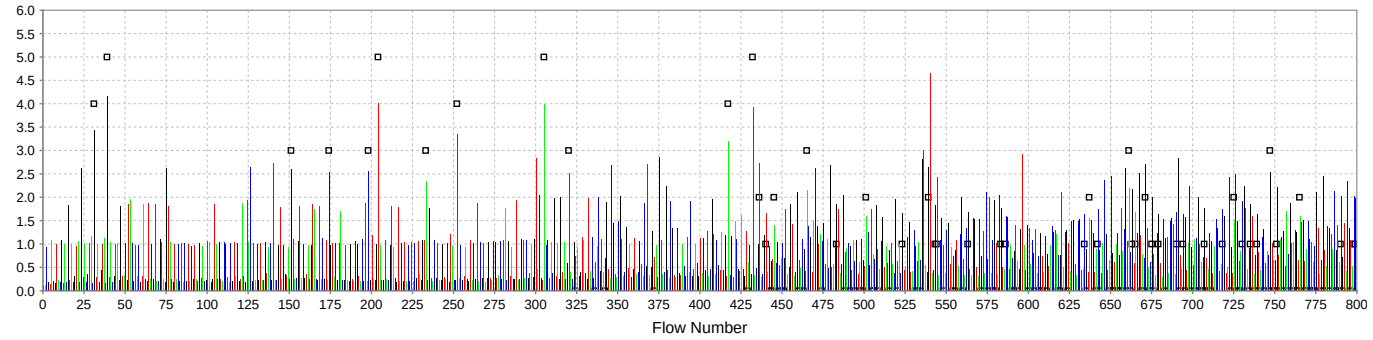

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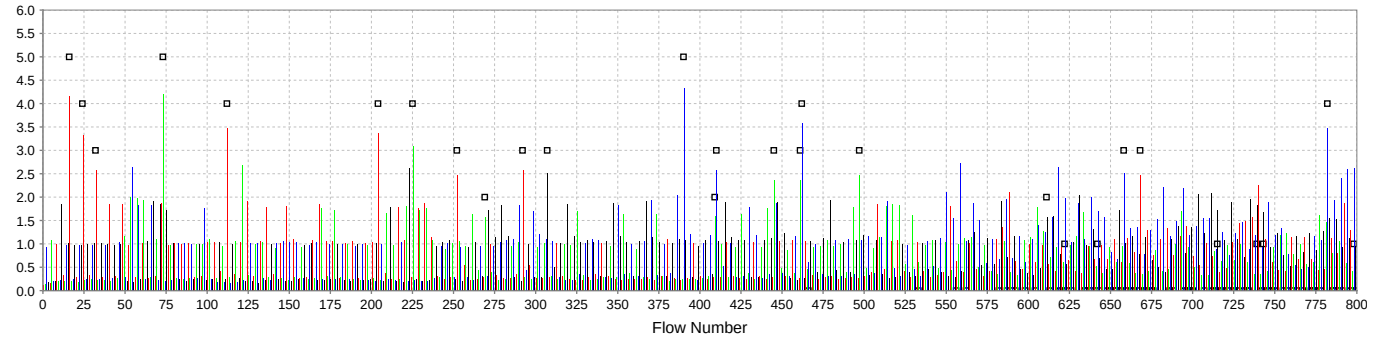

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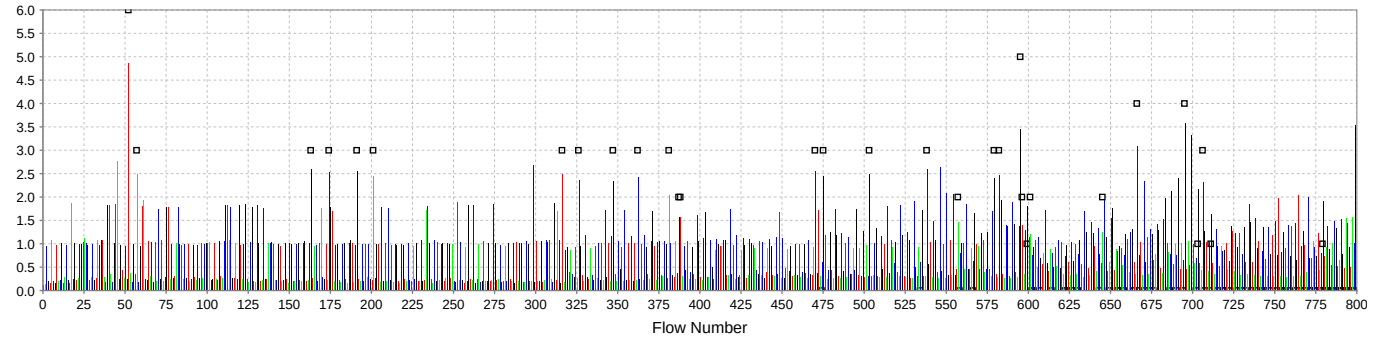

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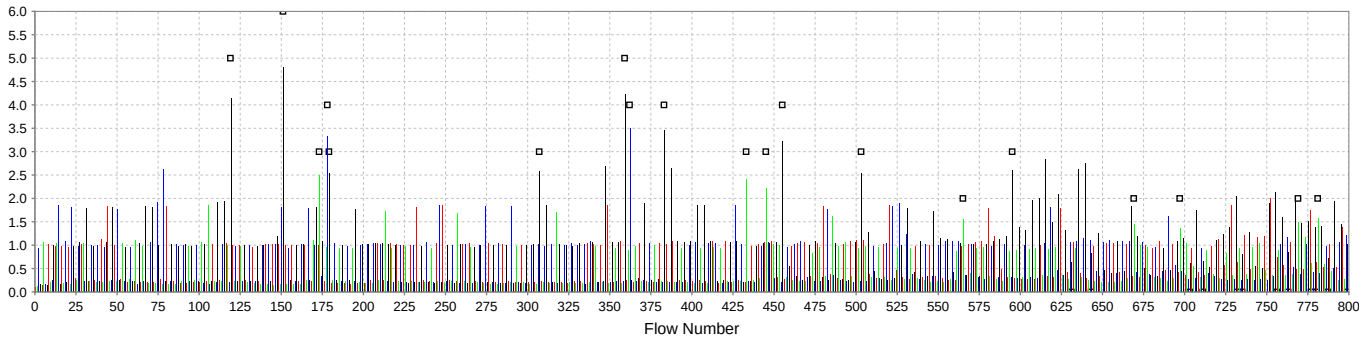

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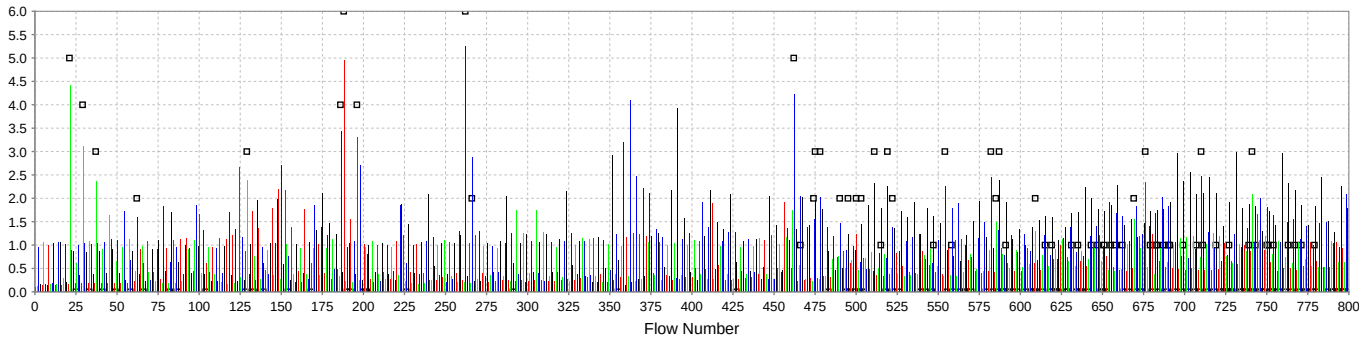

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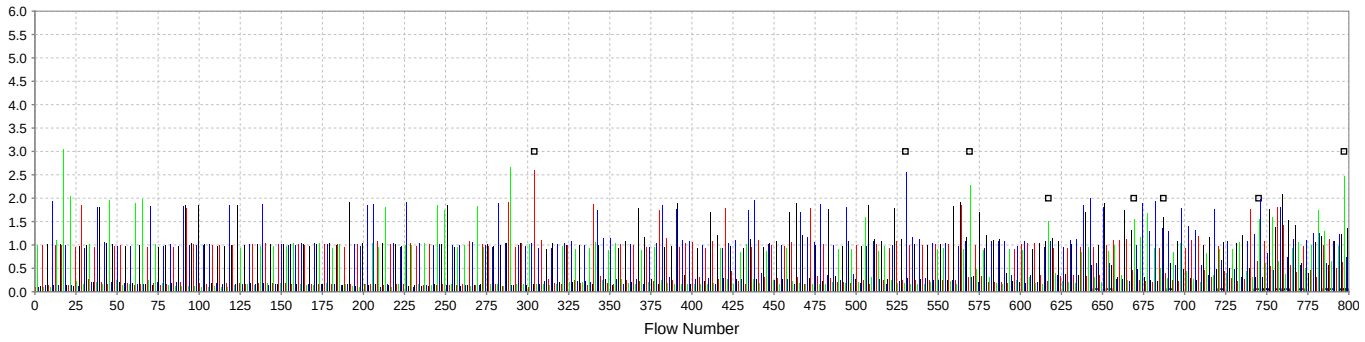

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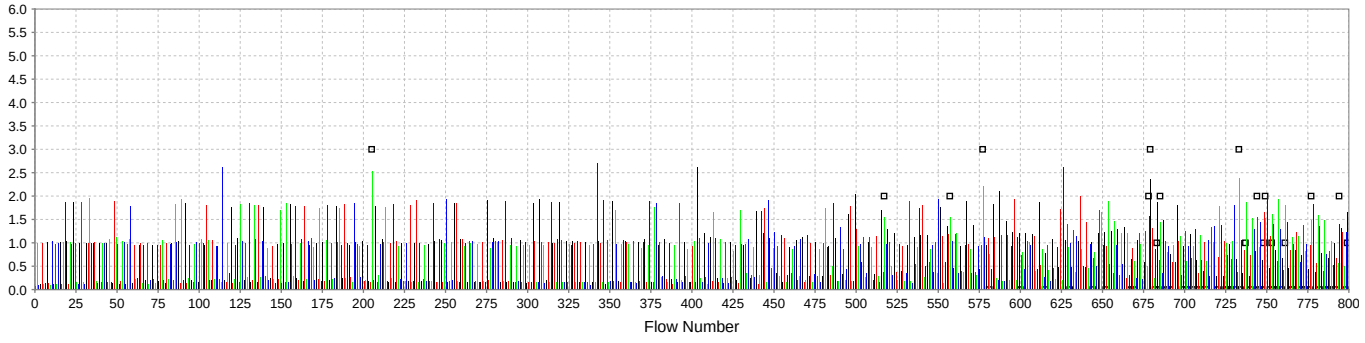

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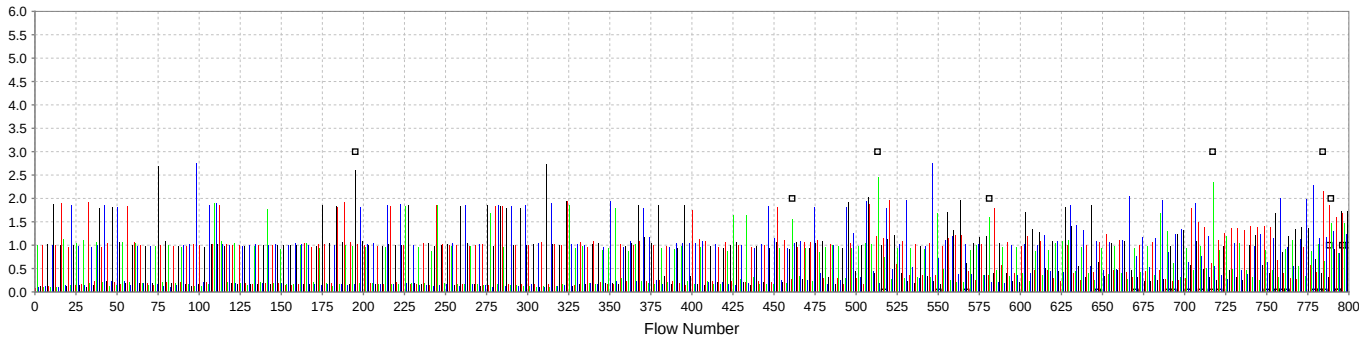

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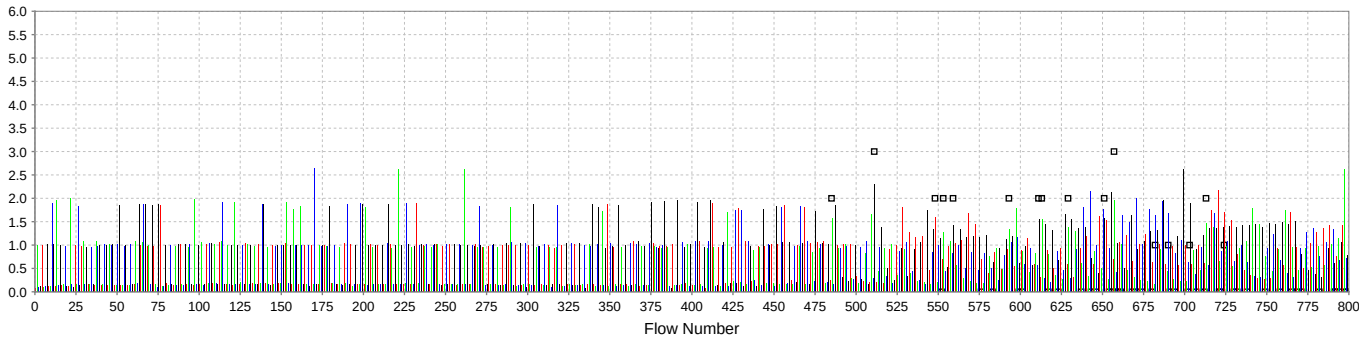

Region 2

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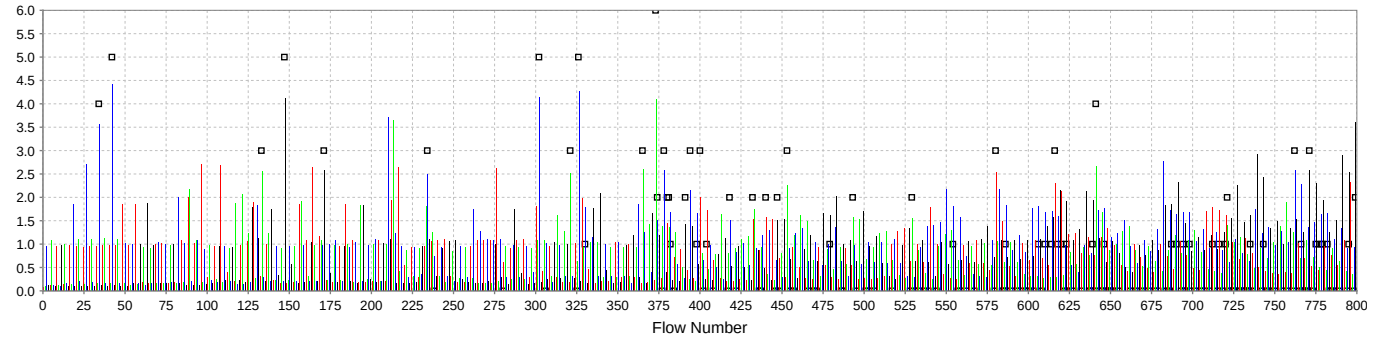

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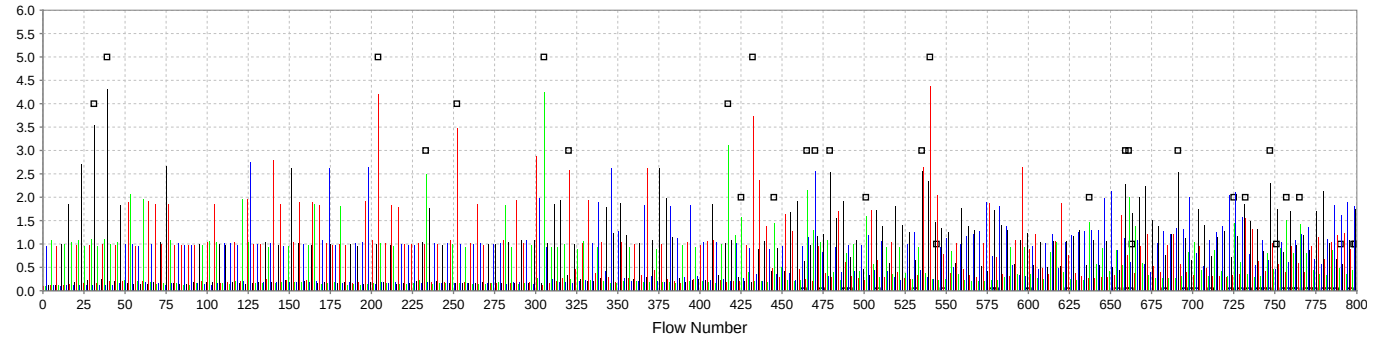

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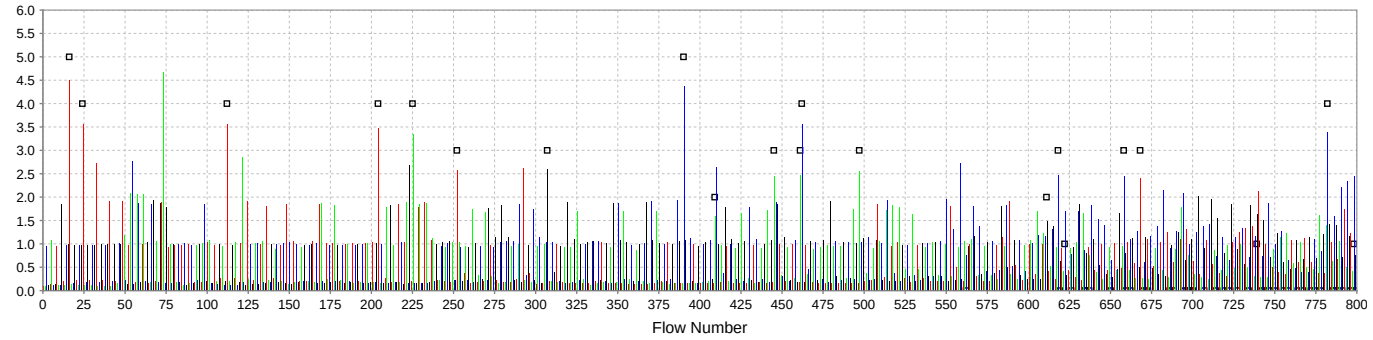

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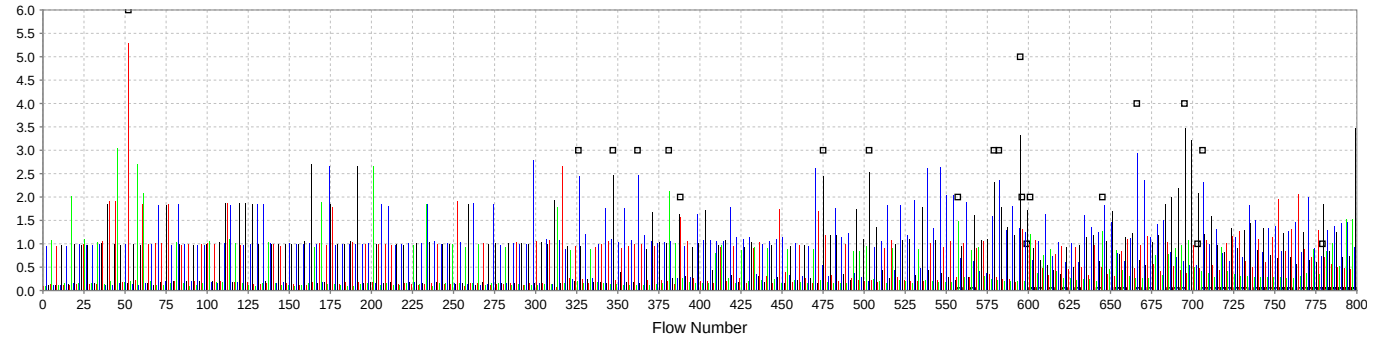

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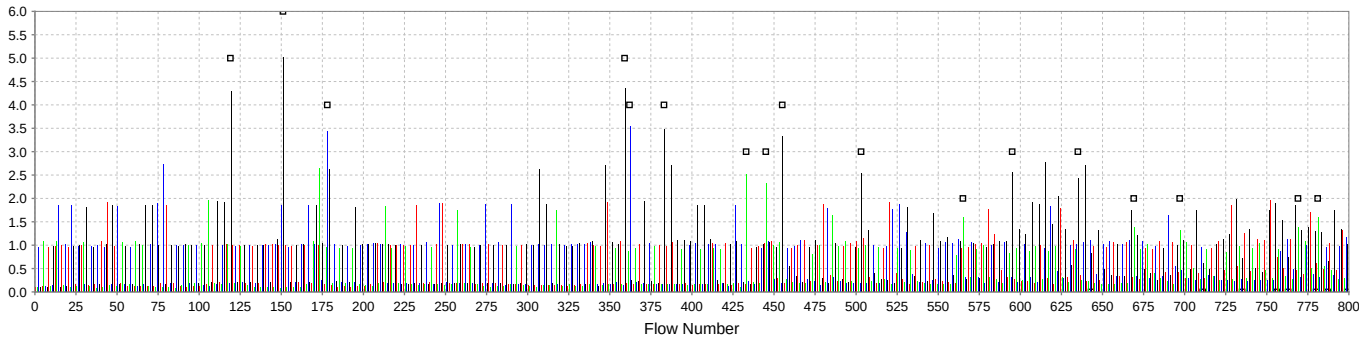

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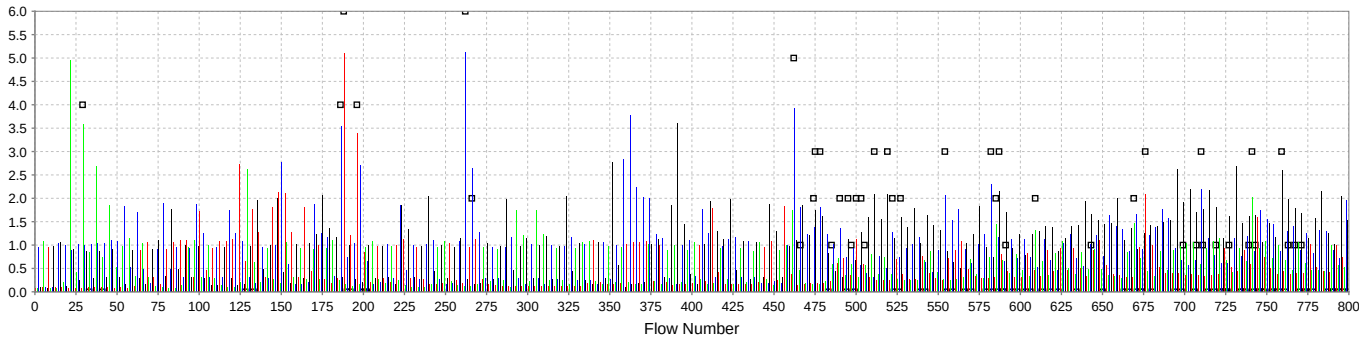

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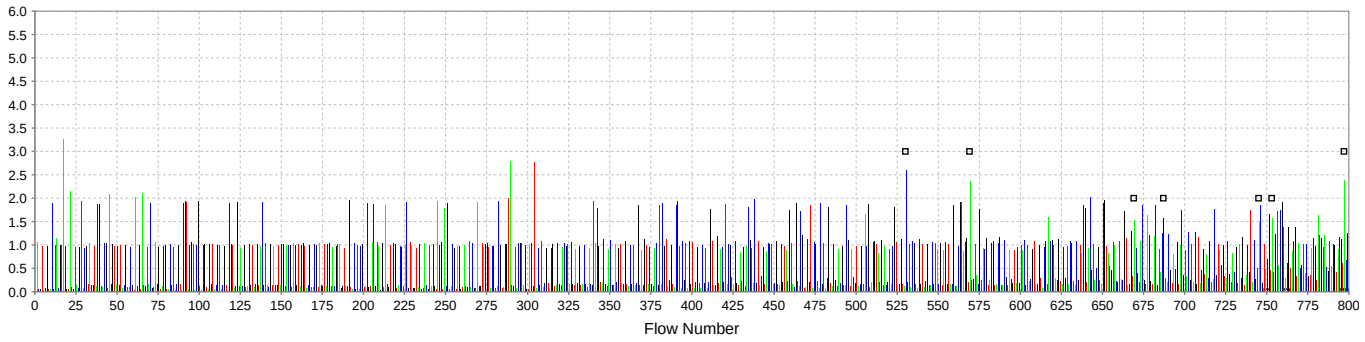

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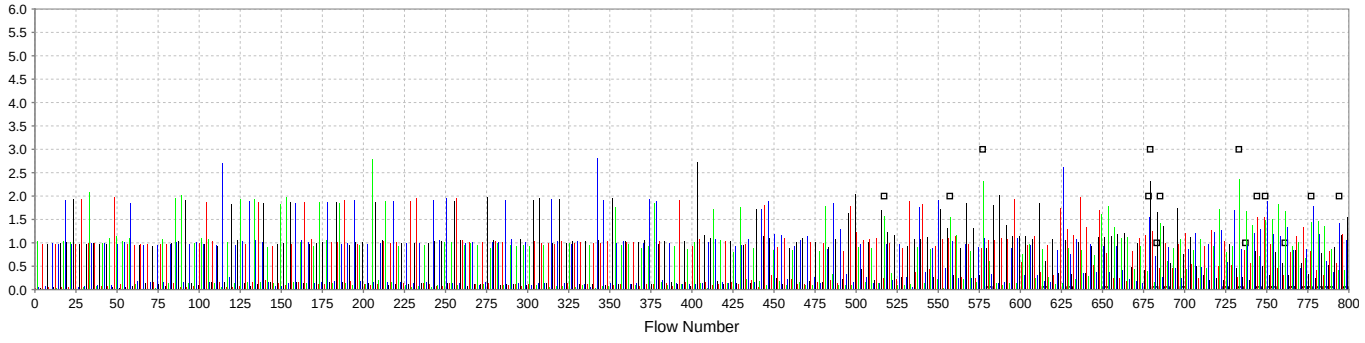

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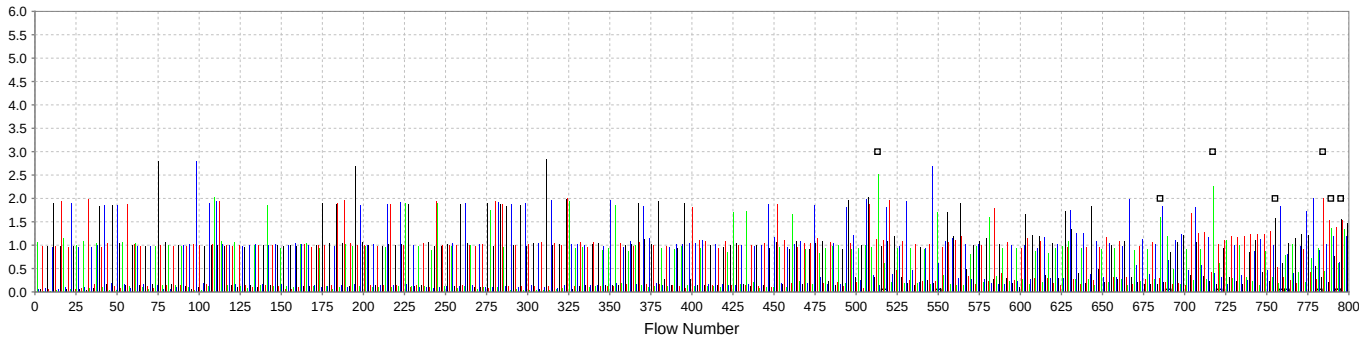

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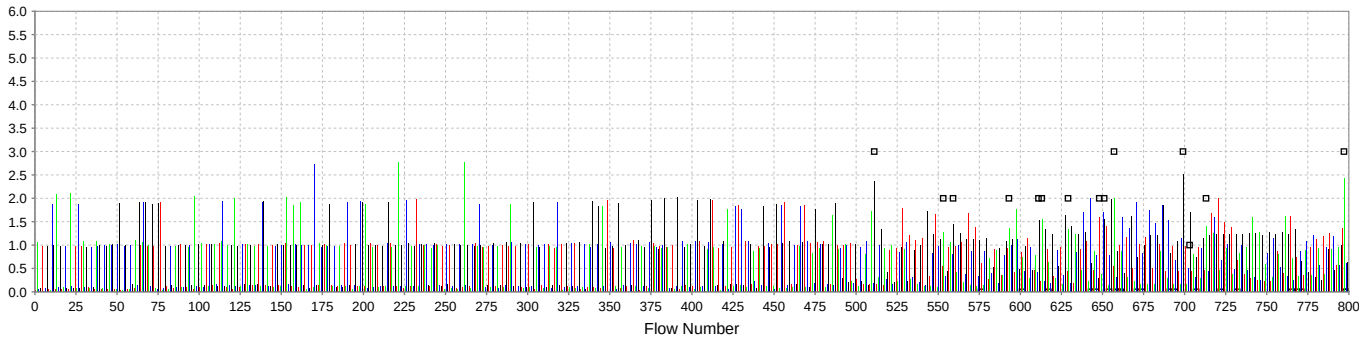

Region 3

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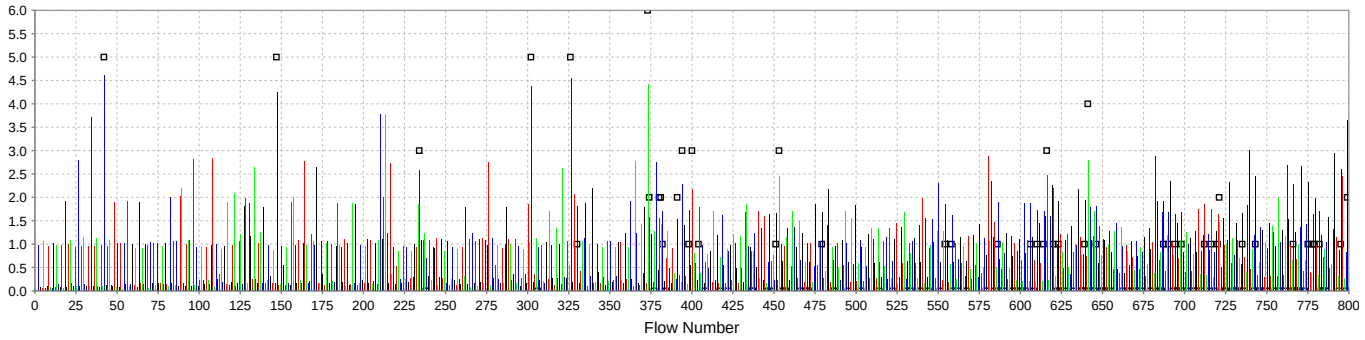

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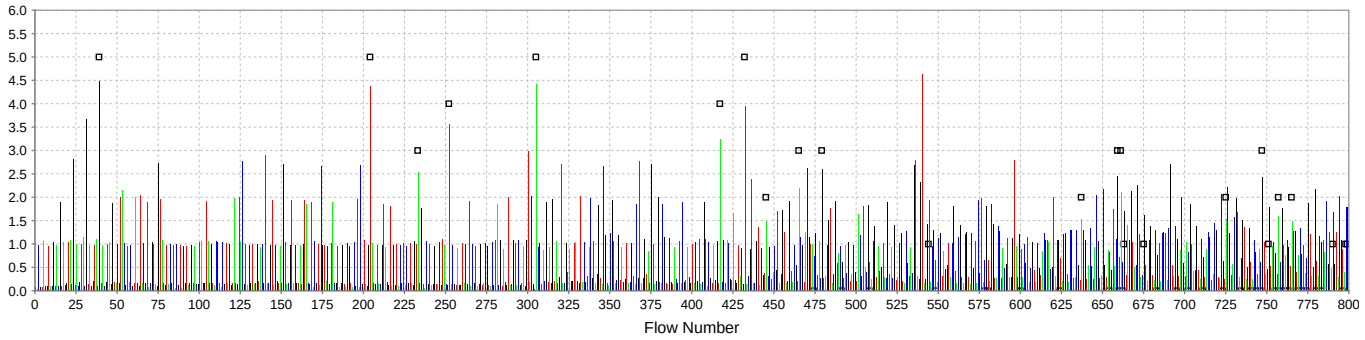

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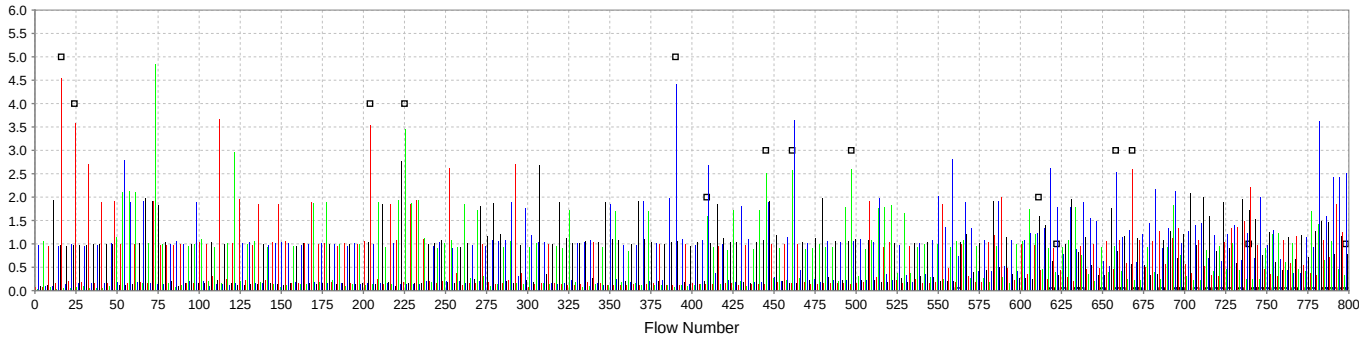

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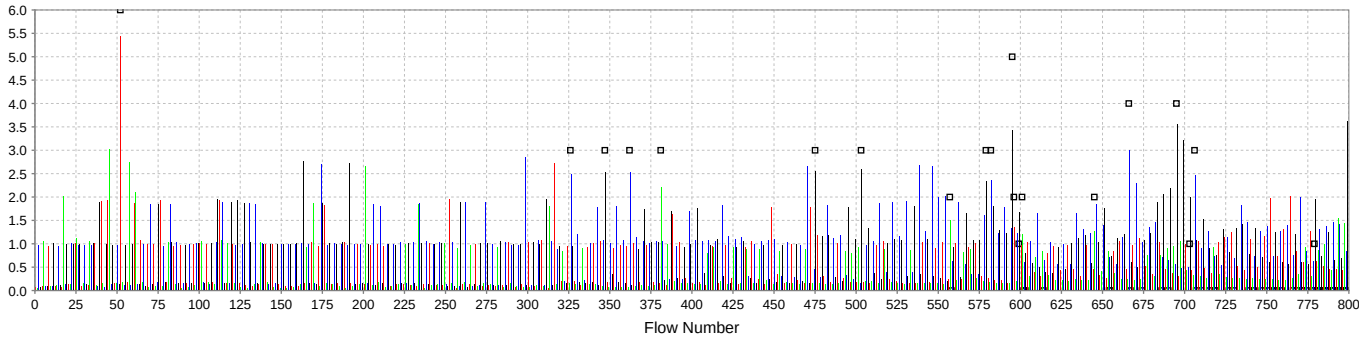

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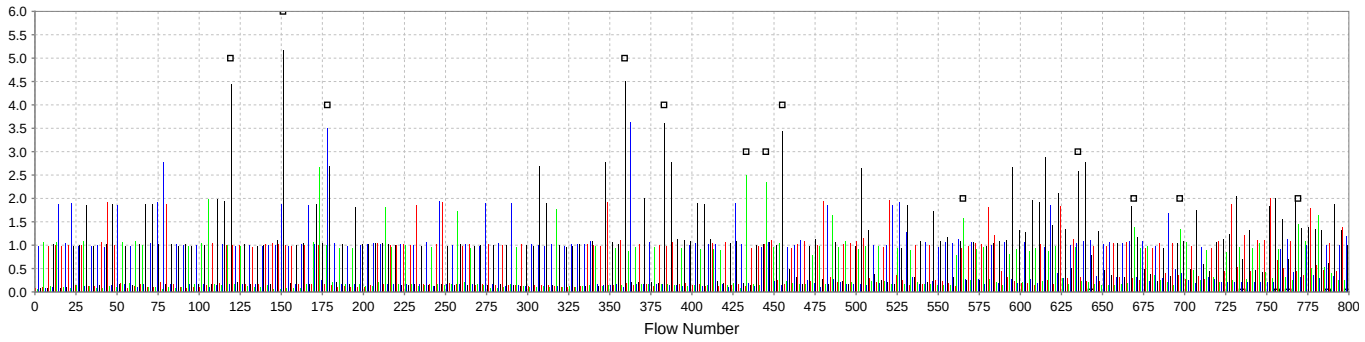

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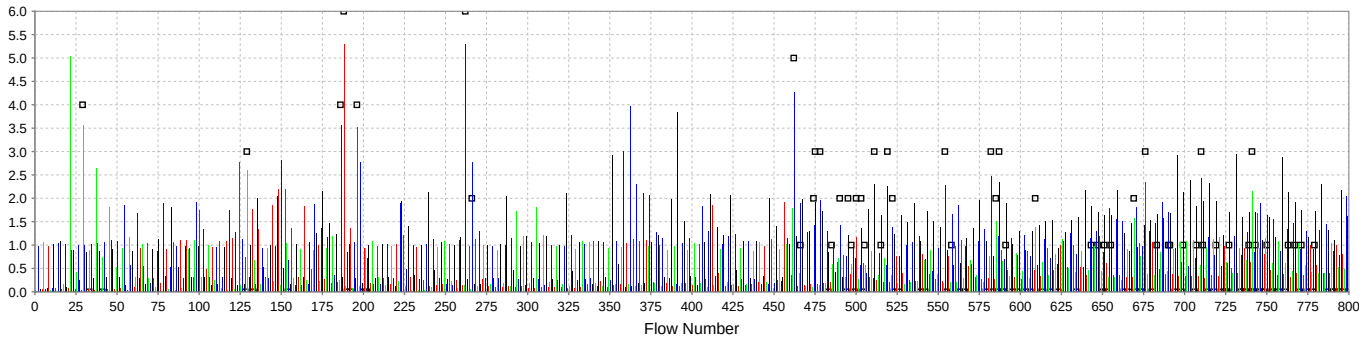

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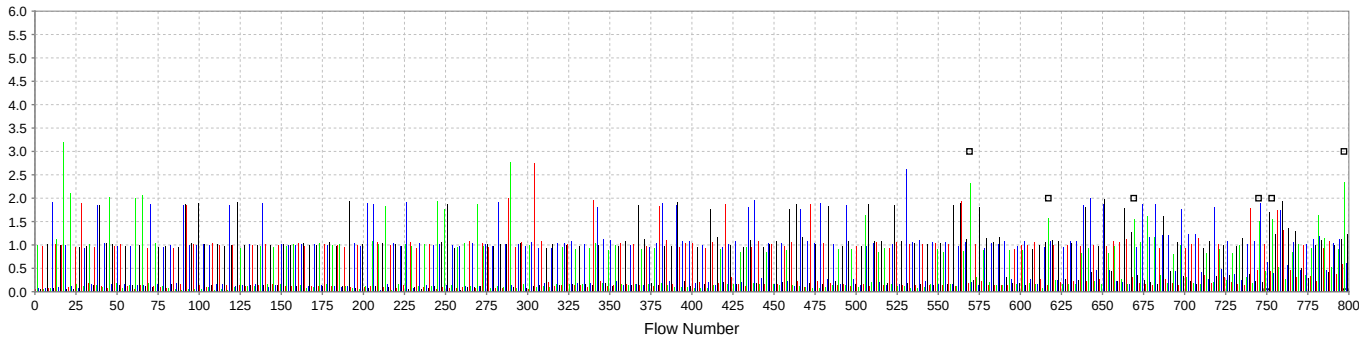

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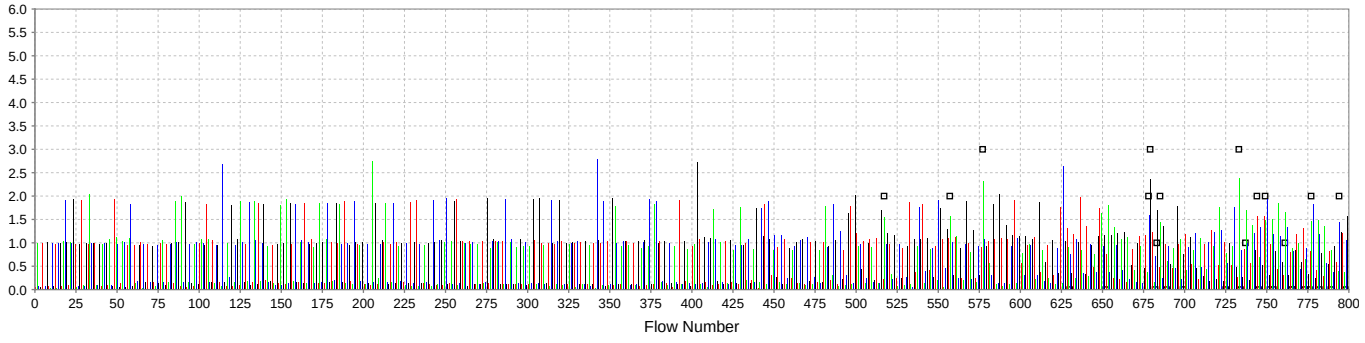

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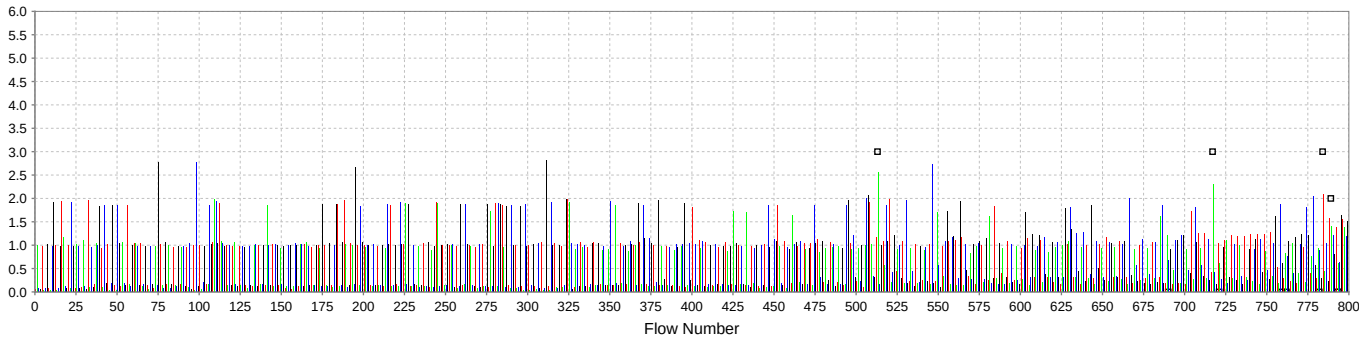

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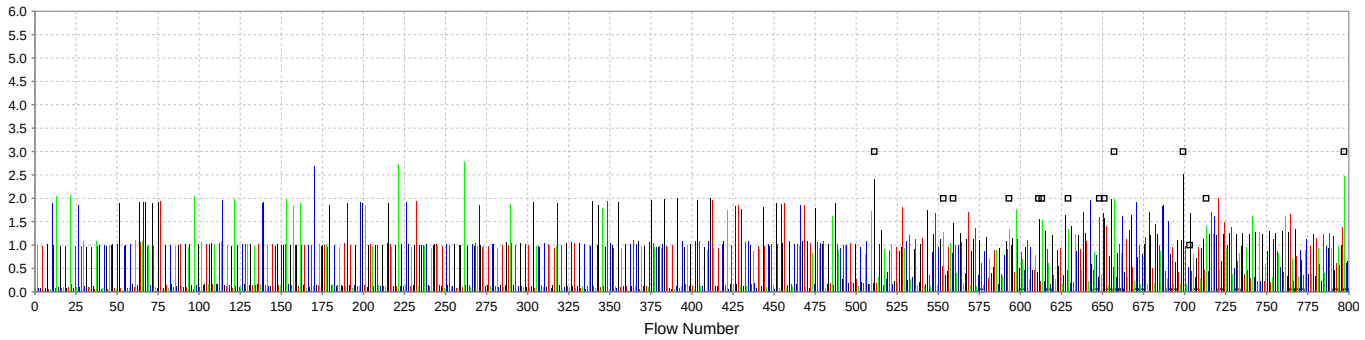

Region 4

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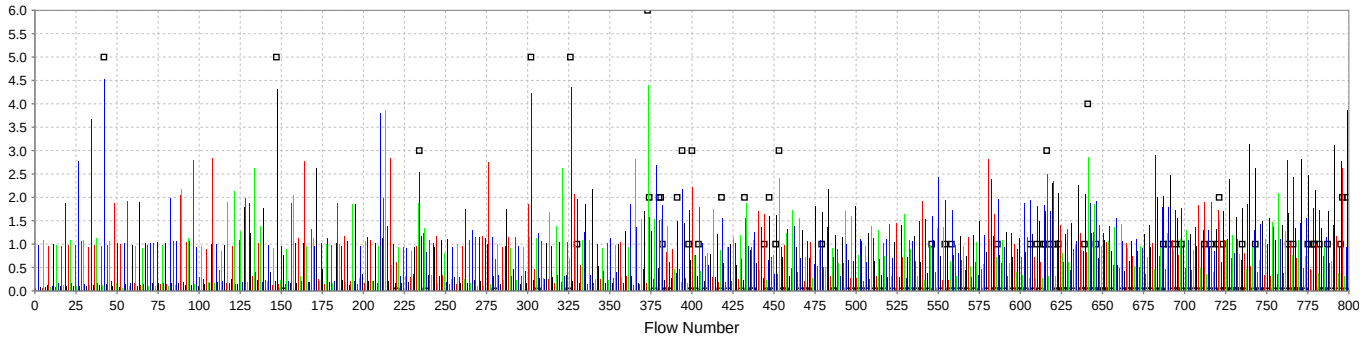

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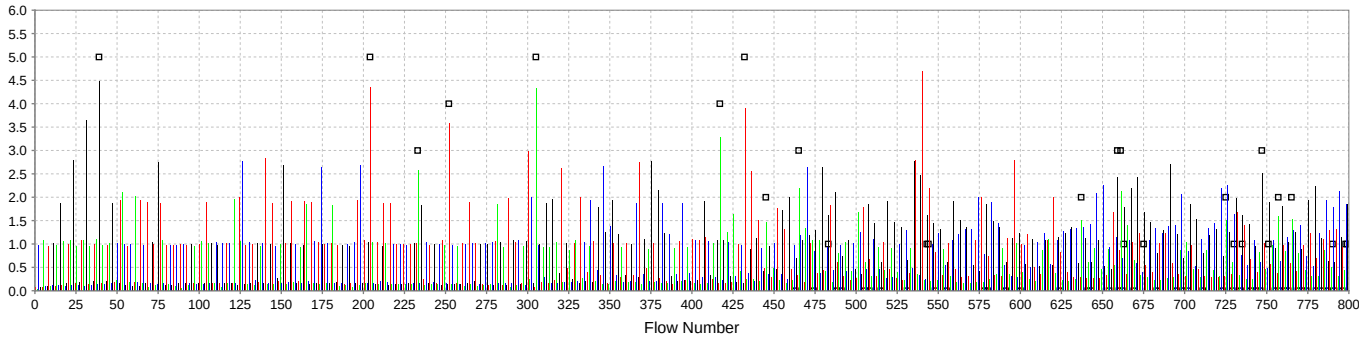

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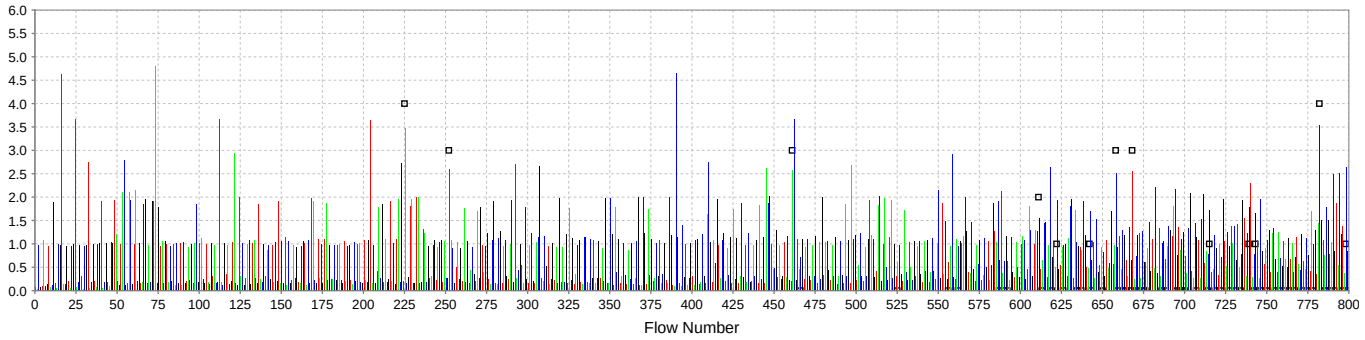

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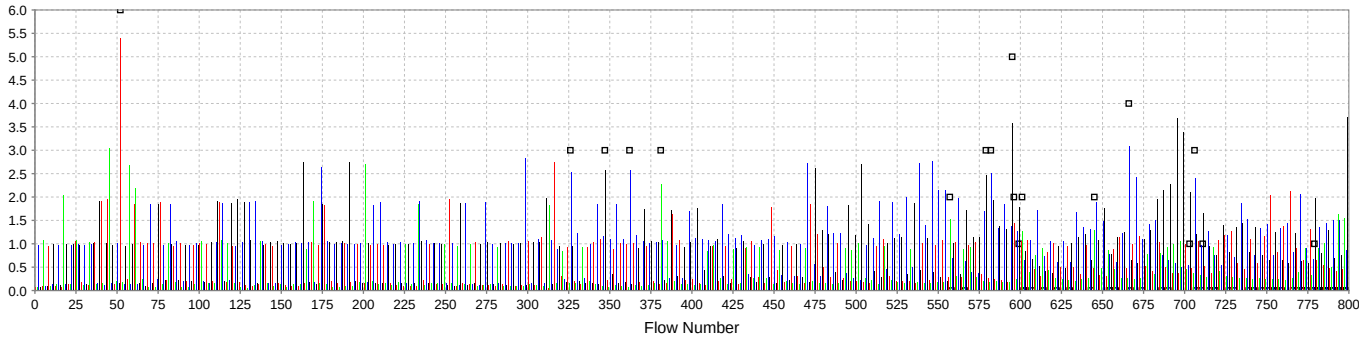

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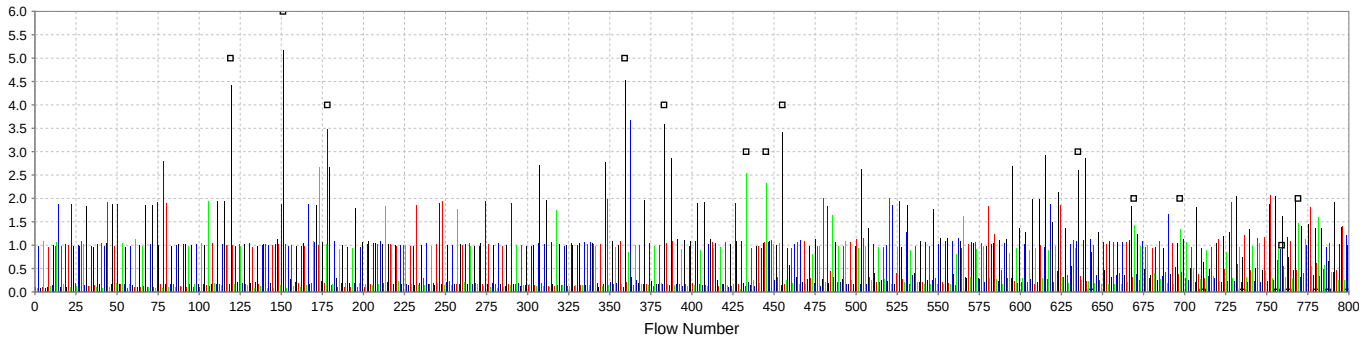

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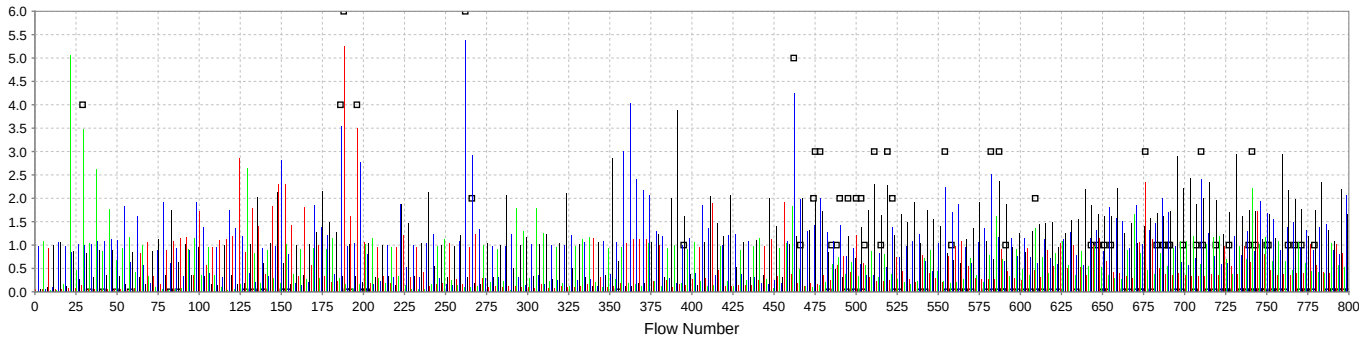

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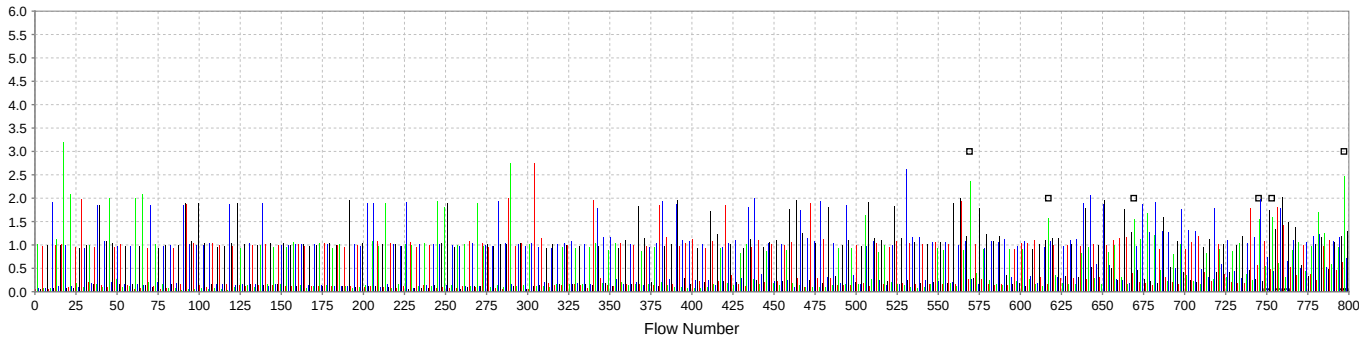

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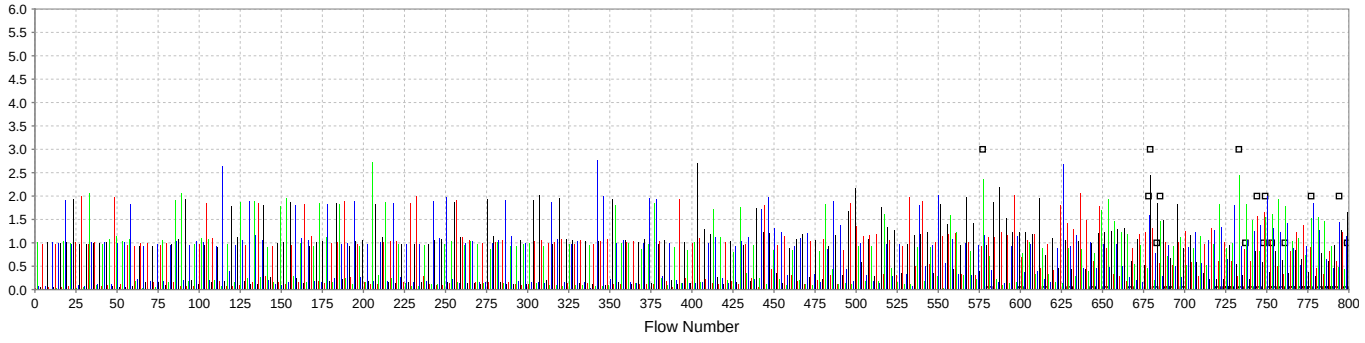

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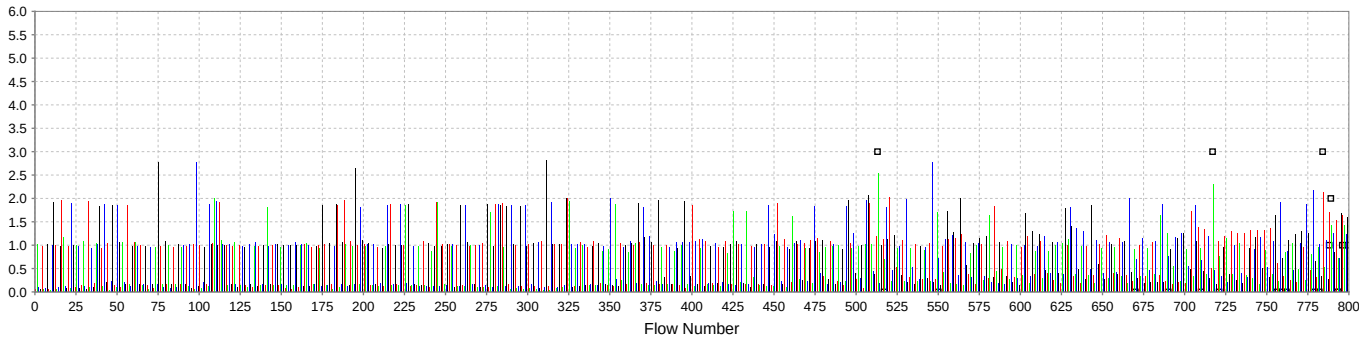

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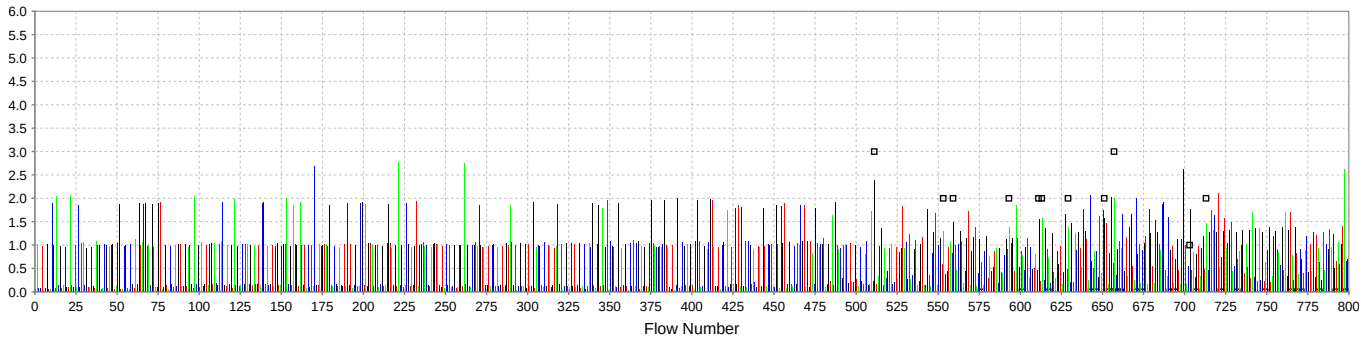

Region 5

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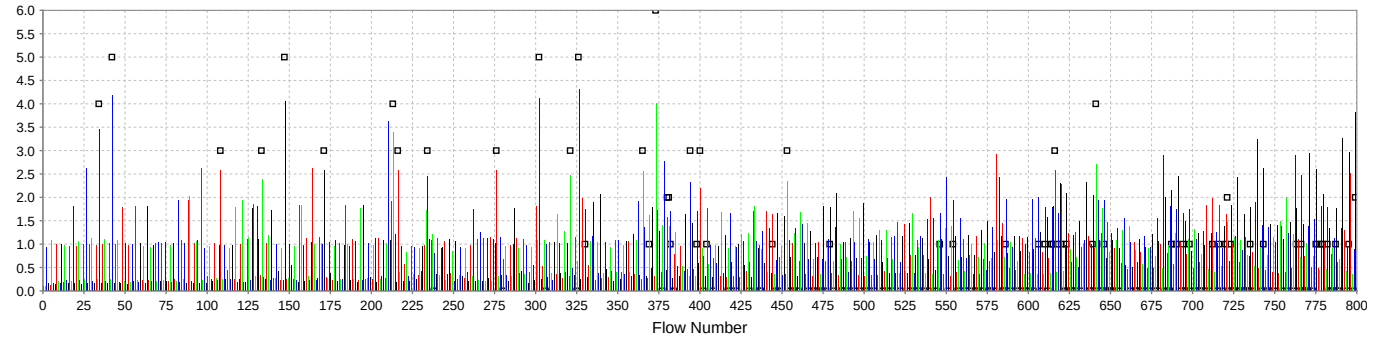

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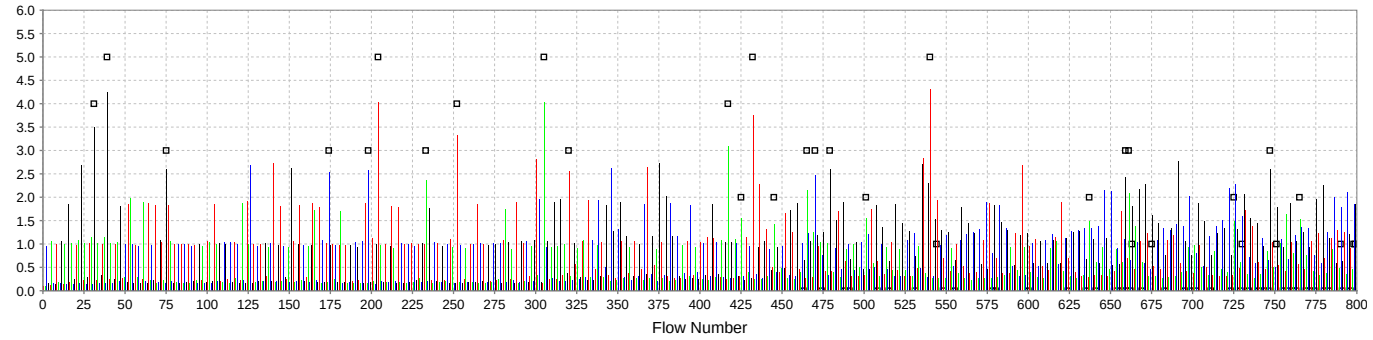

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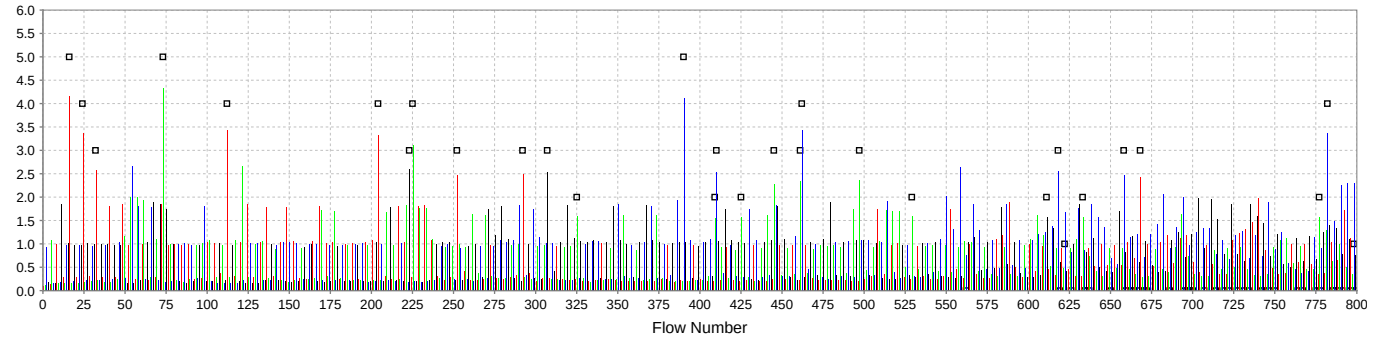

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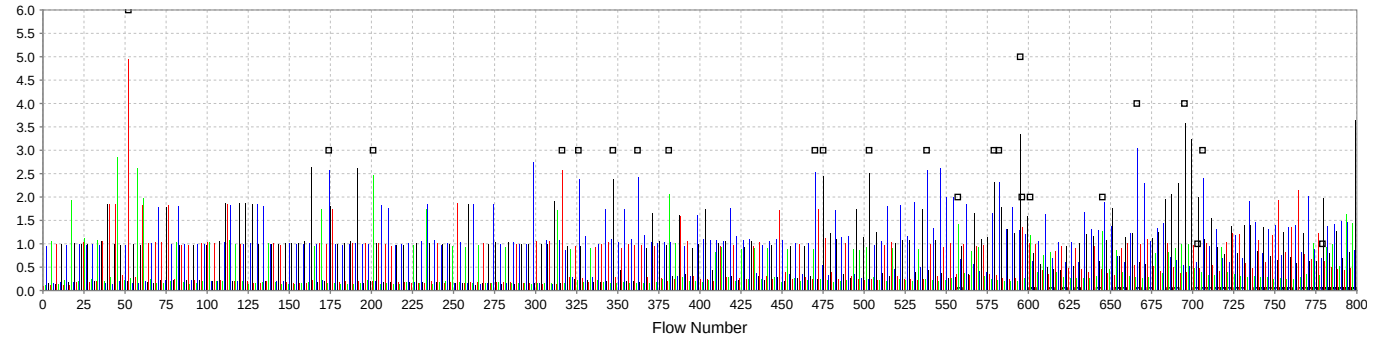
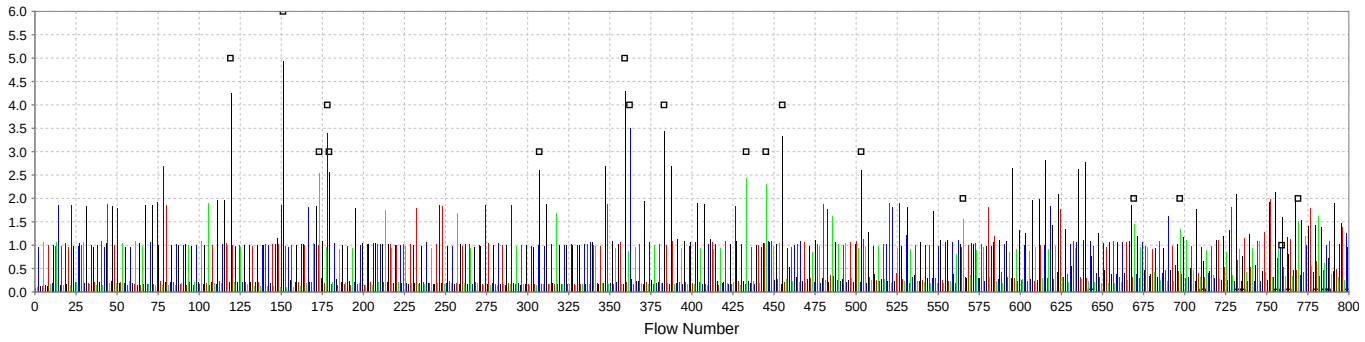


AVTF7



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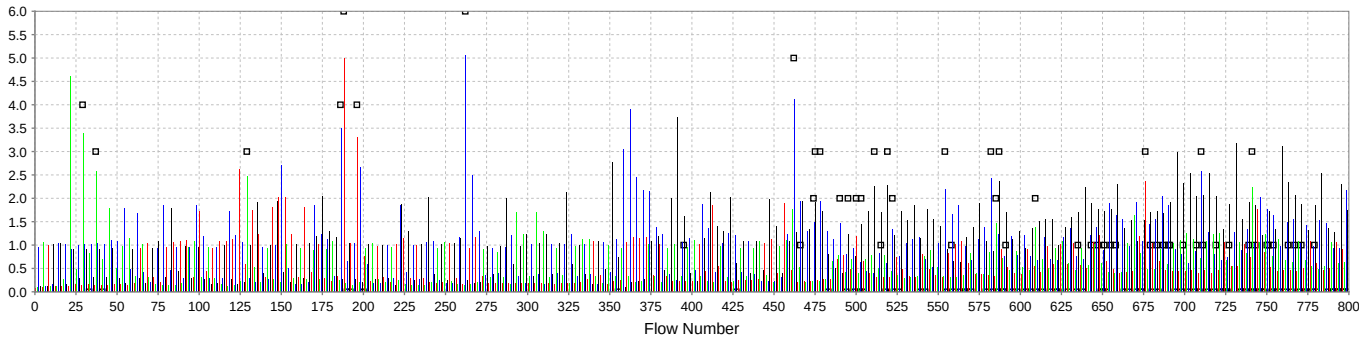

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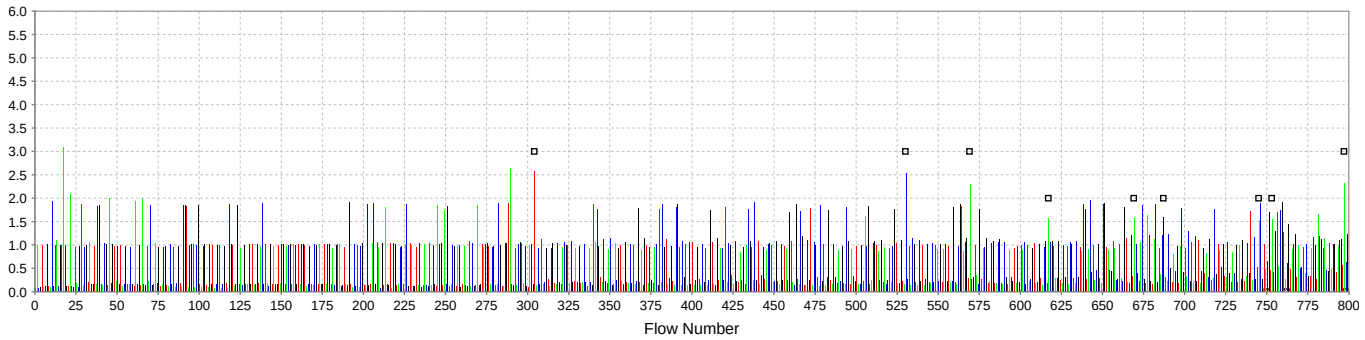

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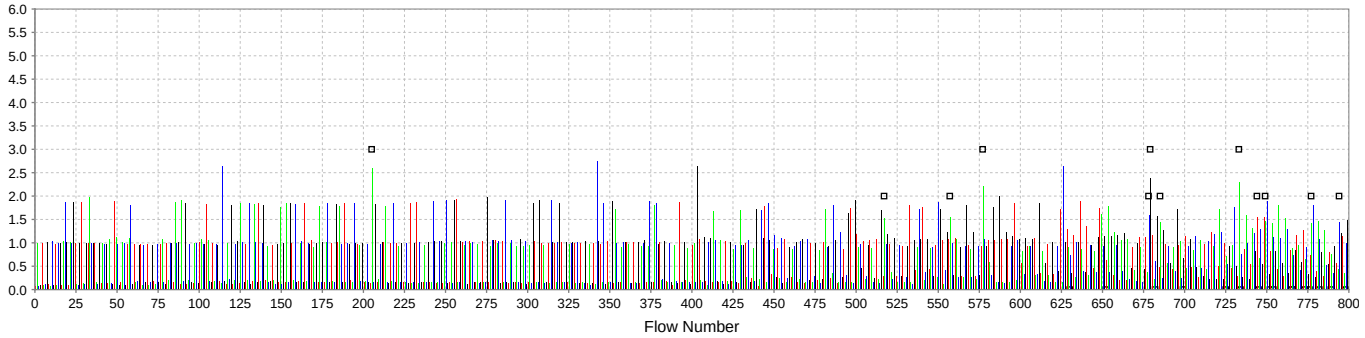

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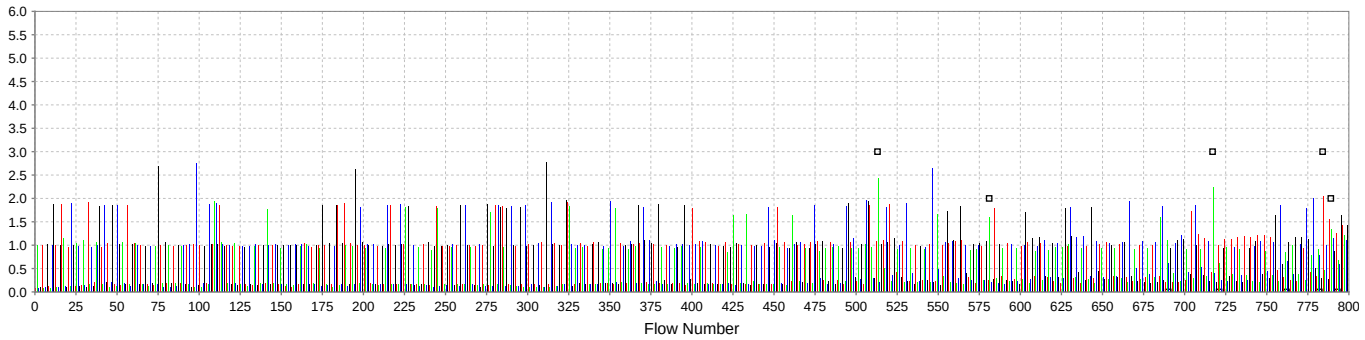

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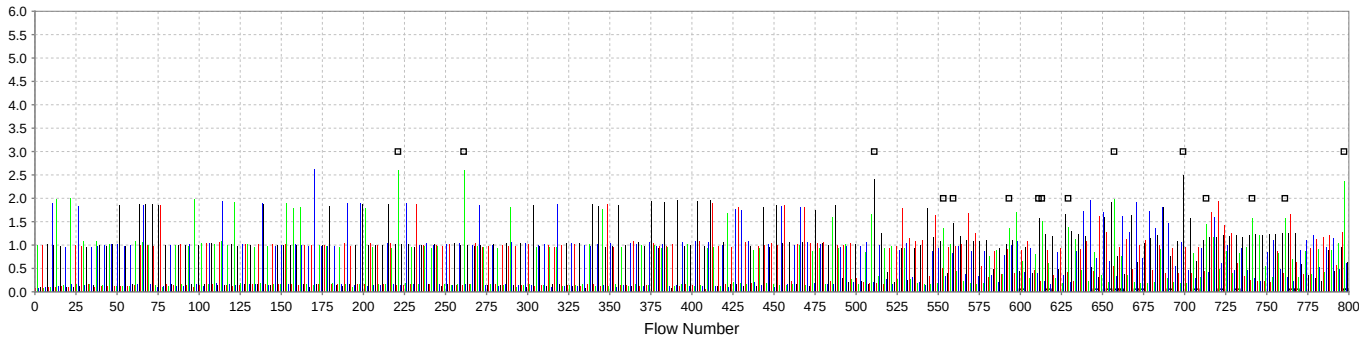
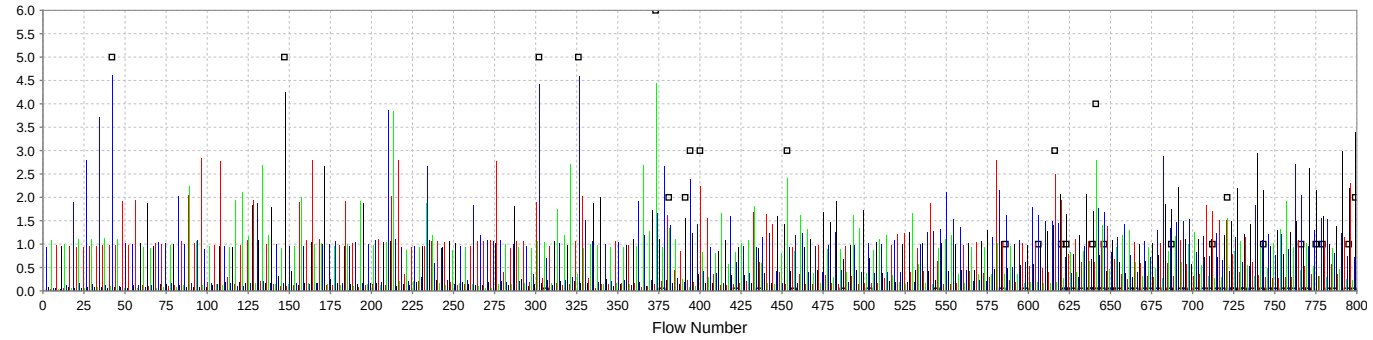


Region 6

AVTF100



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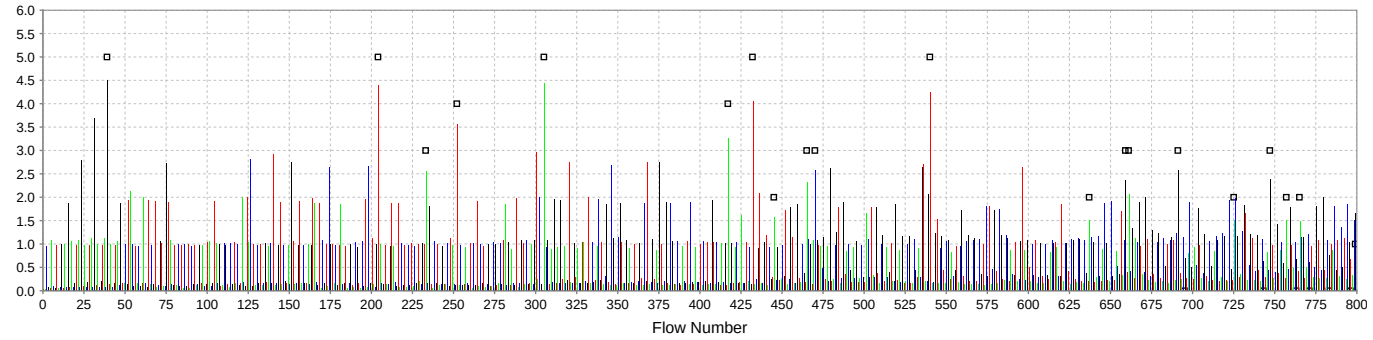

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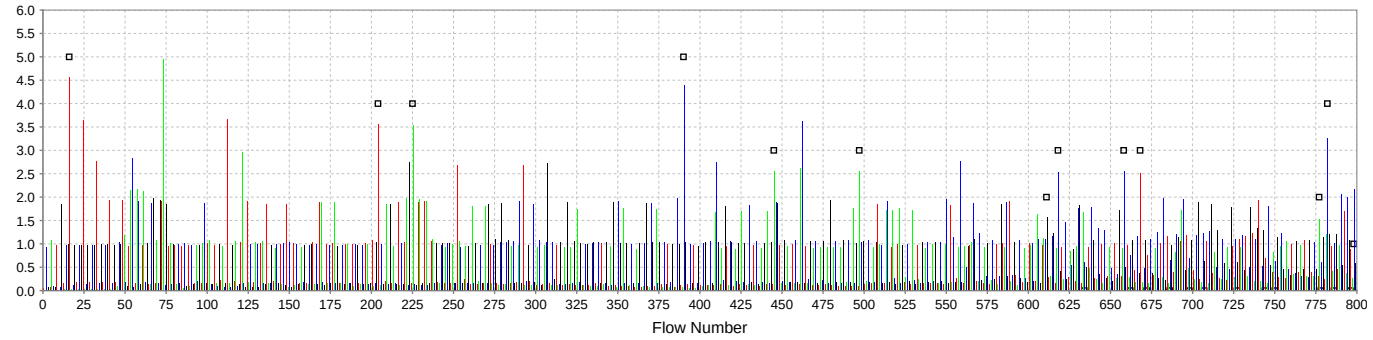

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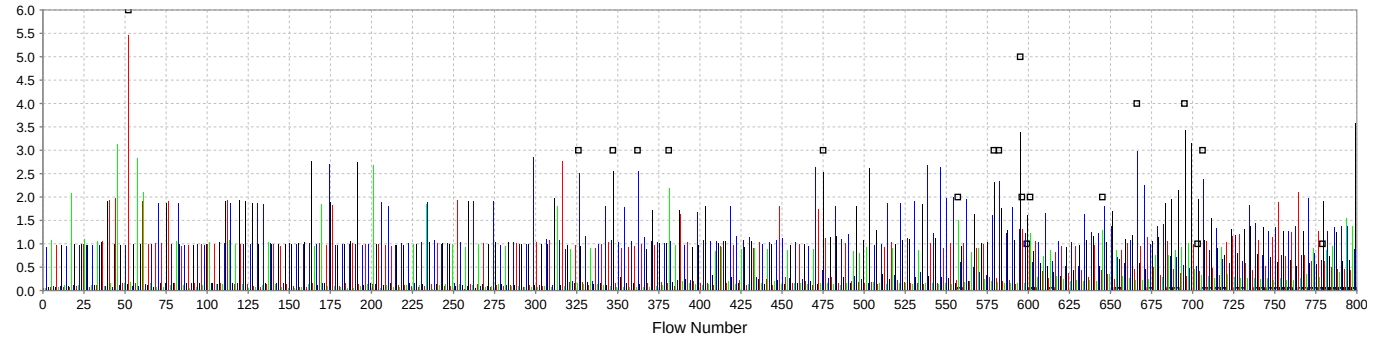
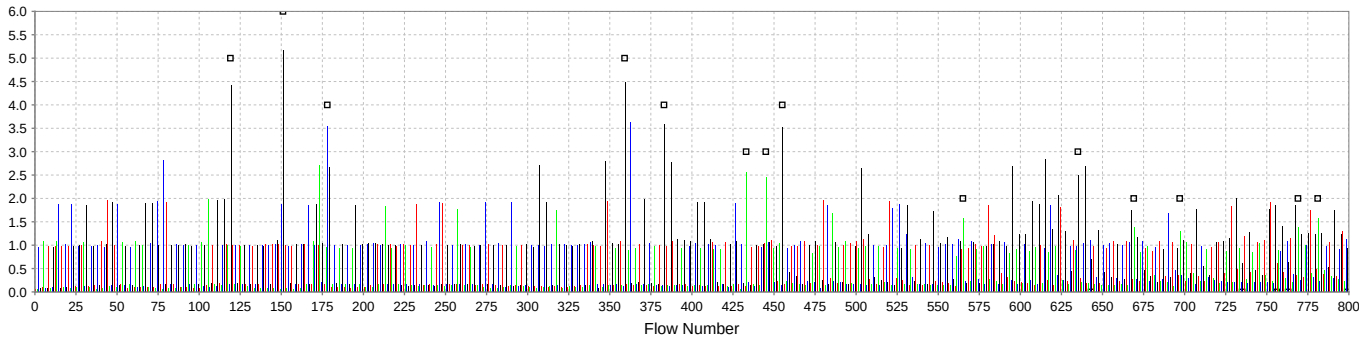


AVTF7



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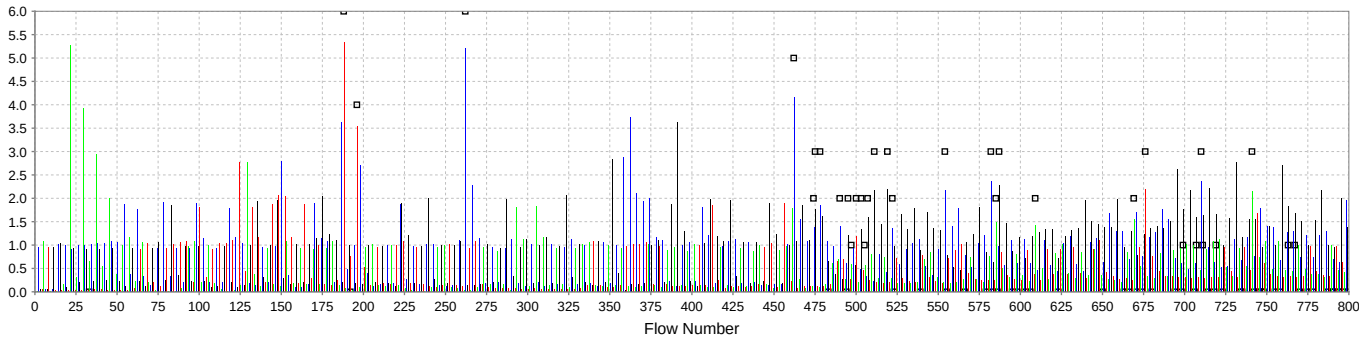

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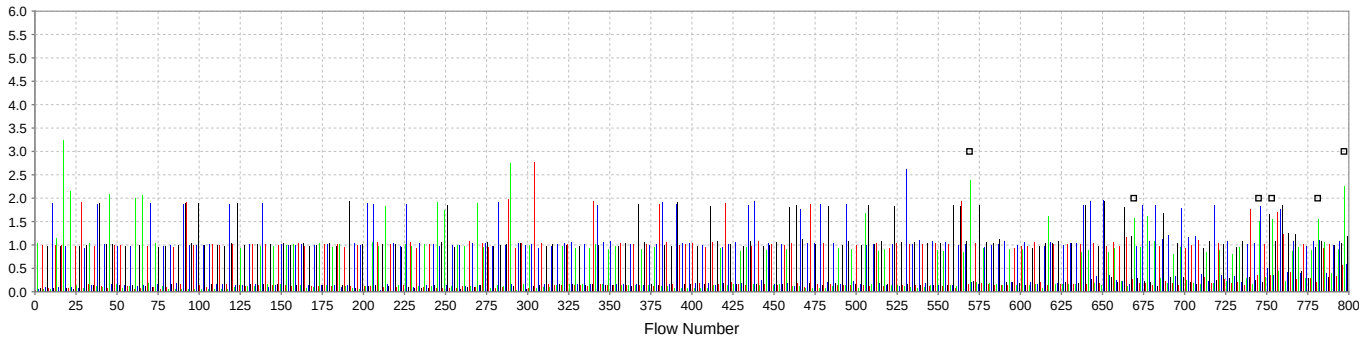

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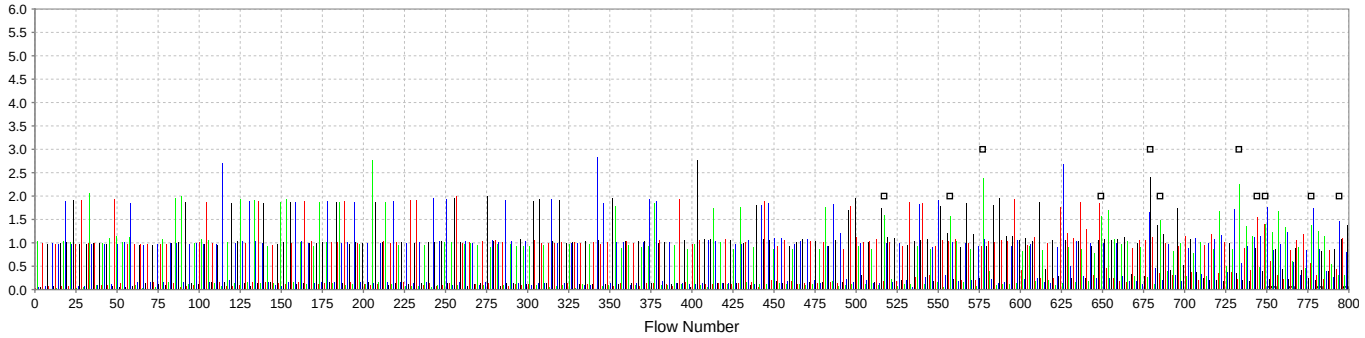

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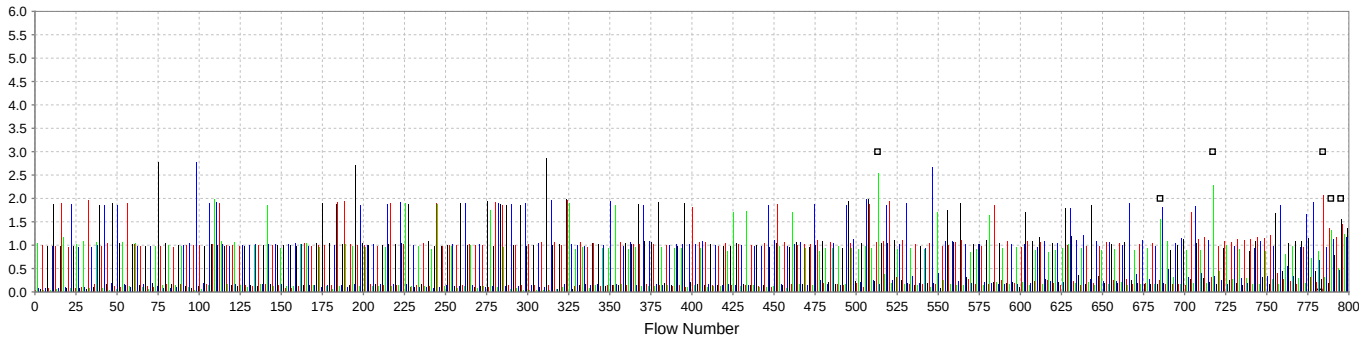
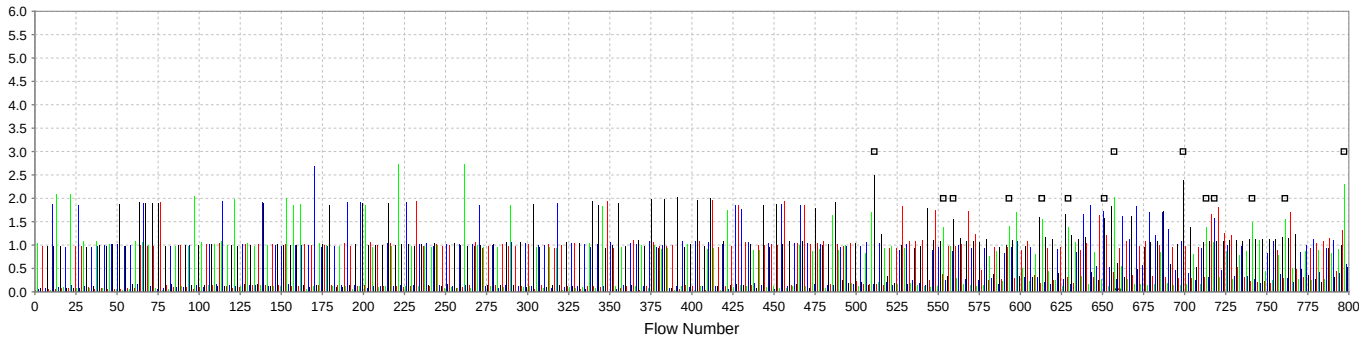
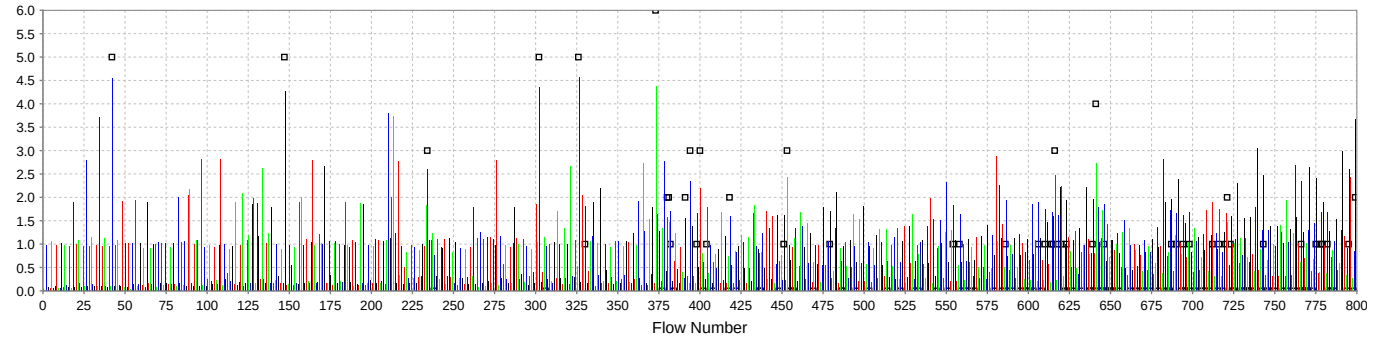


ECTF304

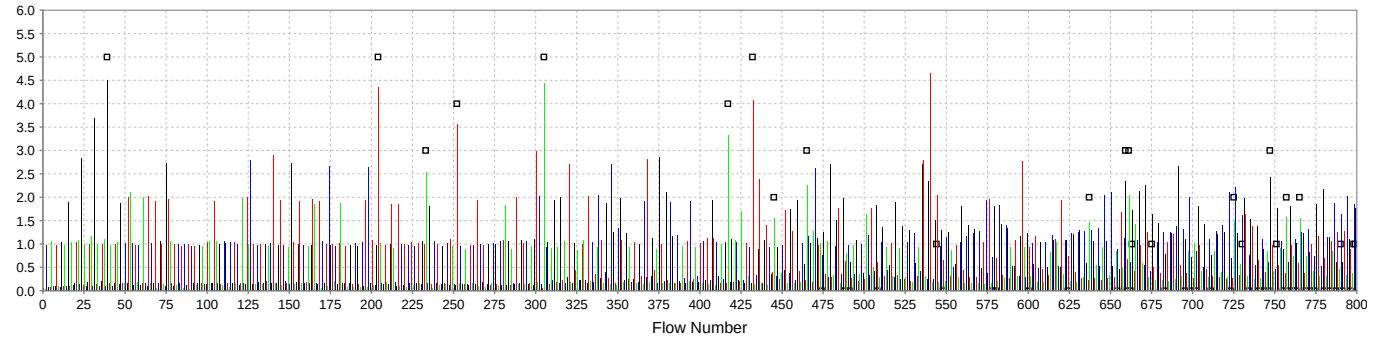


Region 7

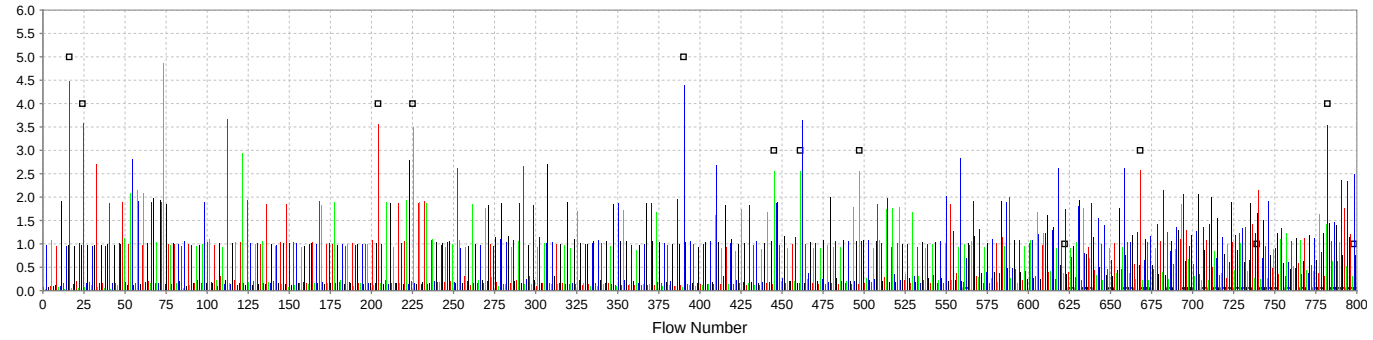
AVTF100



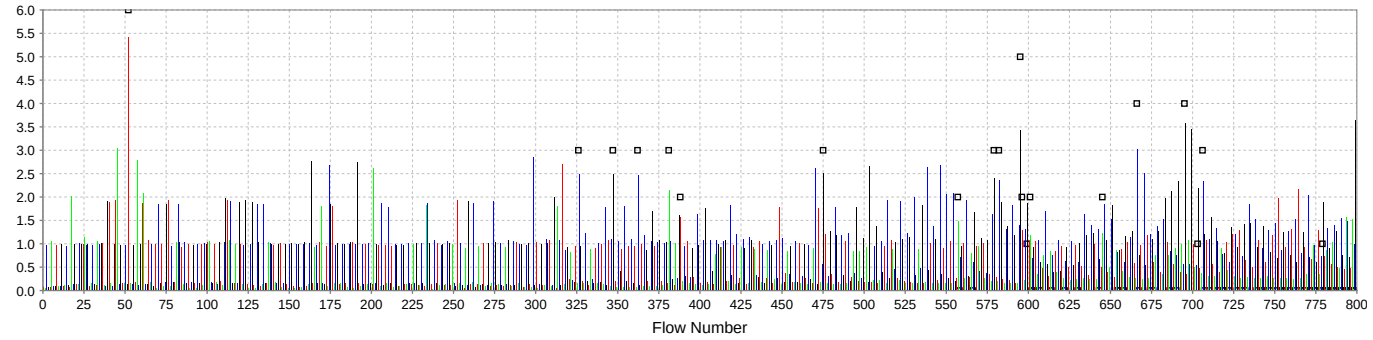
AVTF120



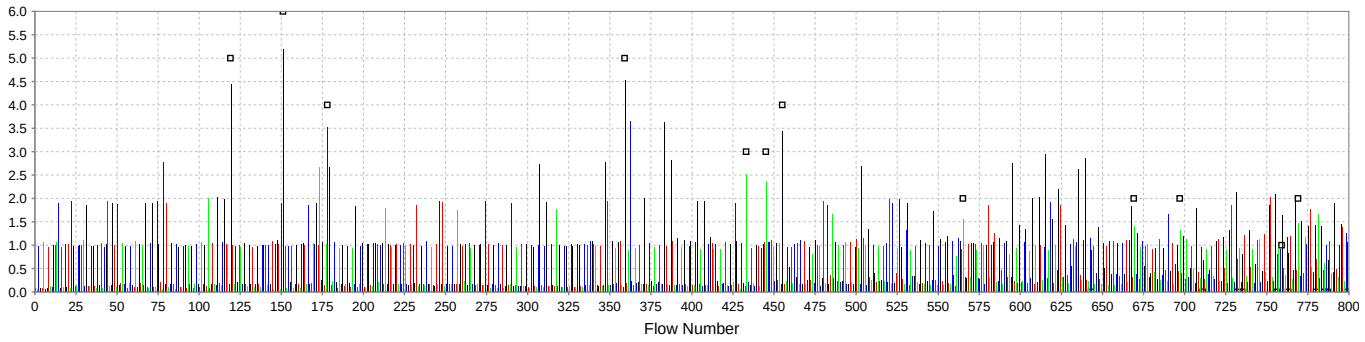
AVTF150



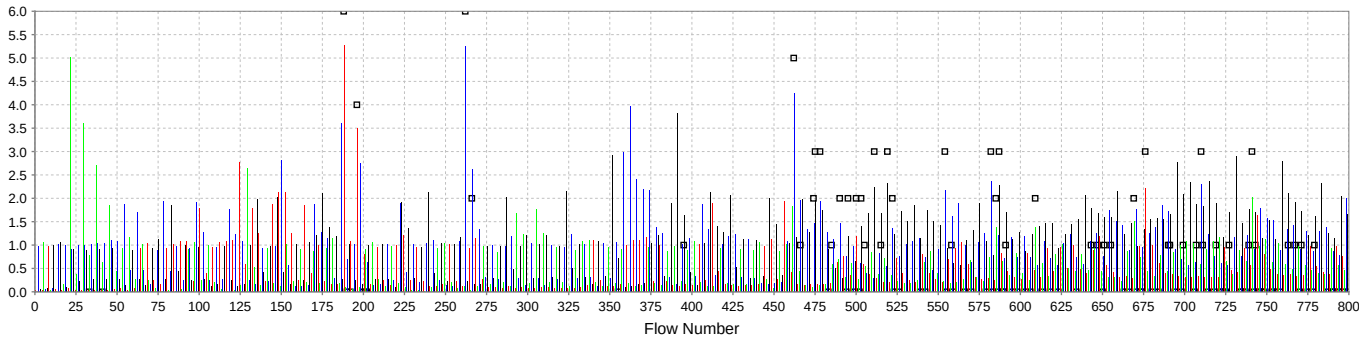
AVTF2



AVTF7



AVTF90



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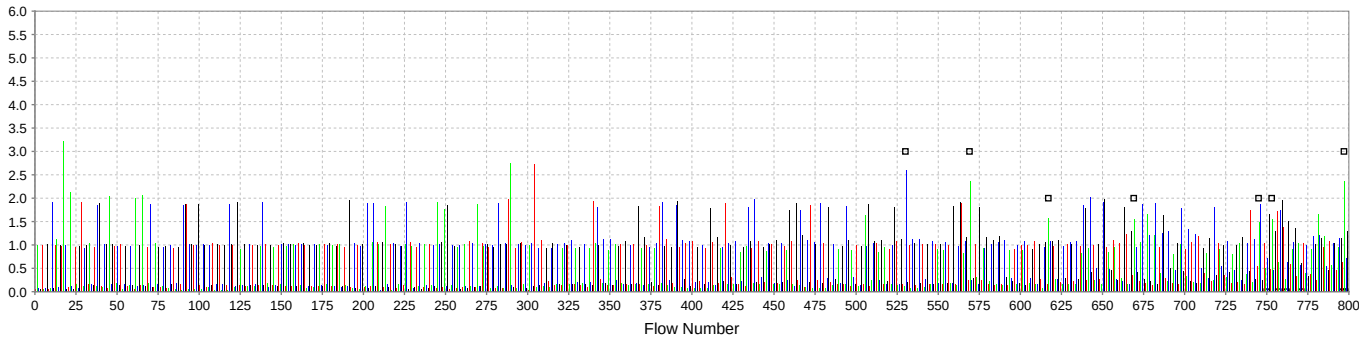
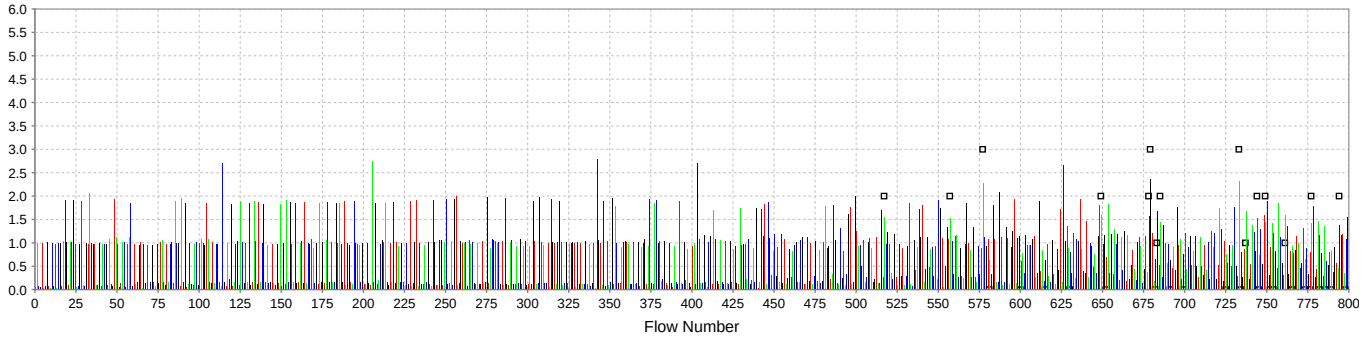


ECTF302



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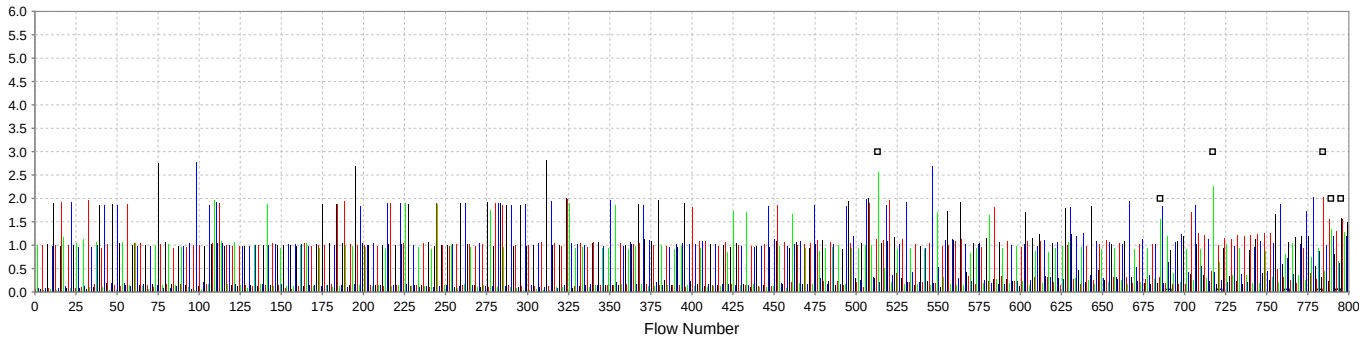

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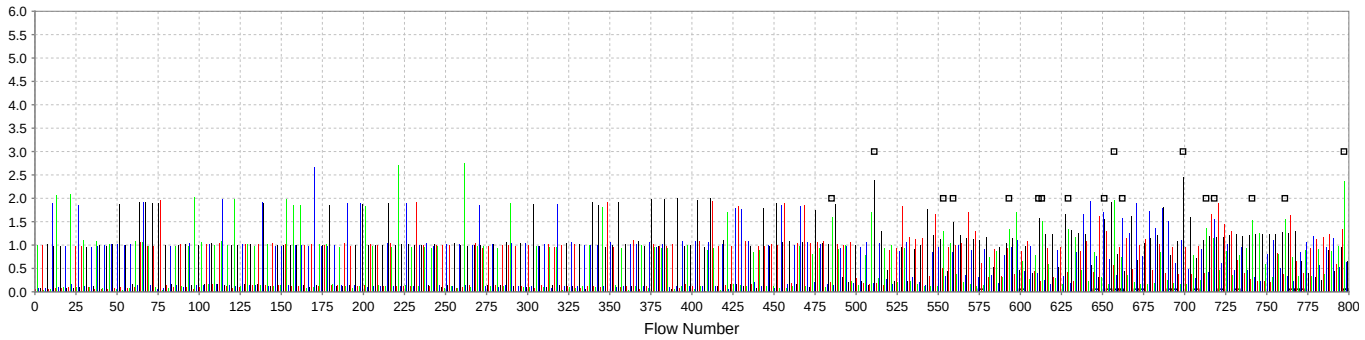
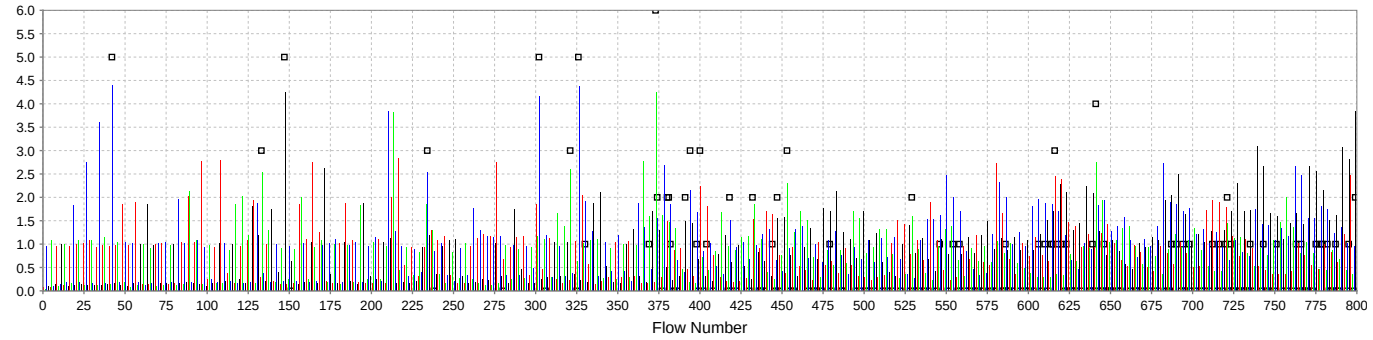


Region 8

AVTF100



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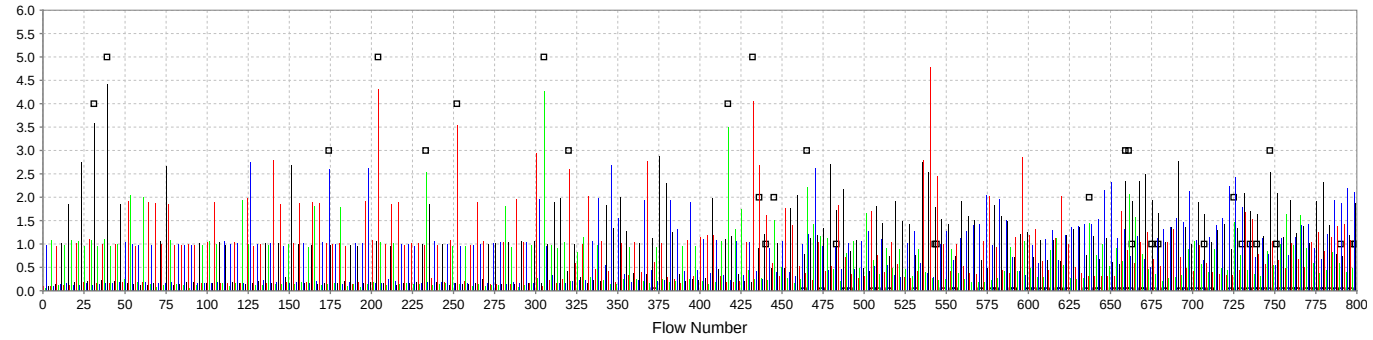

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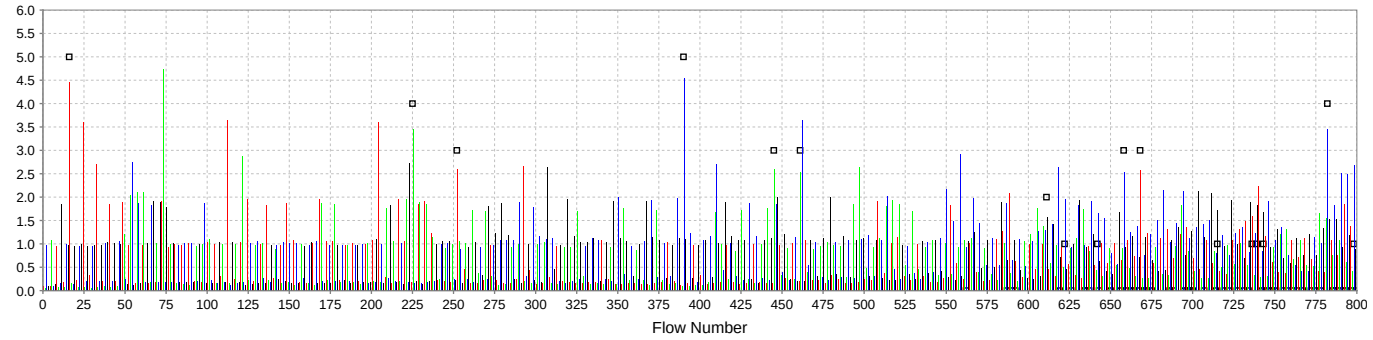
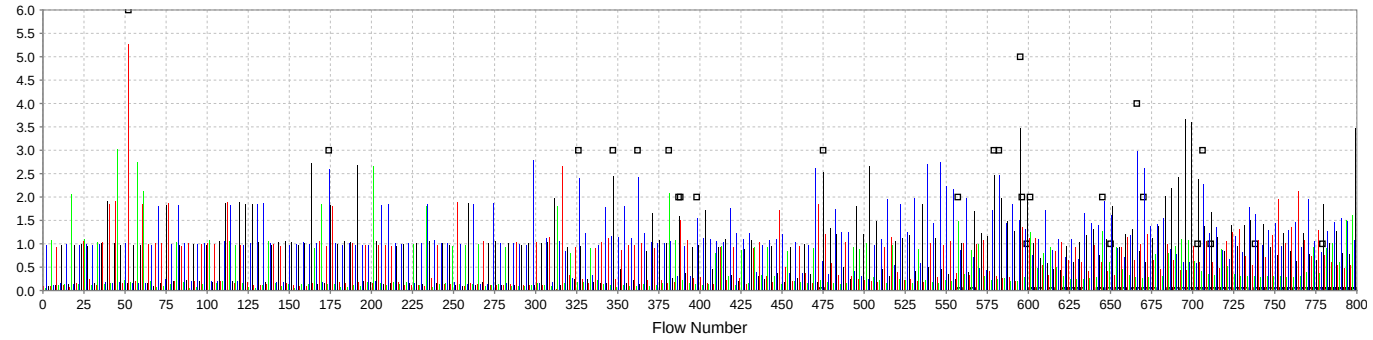
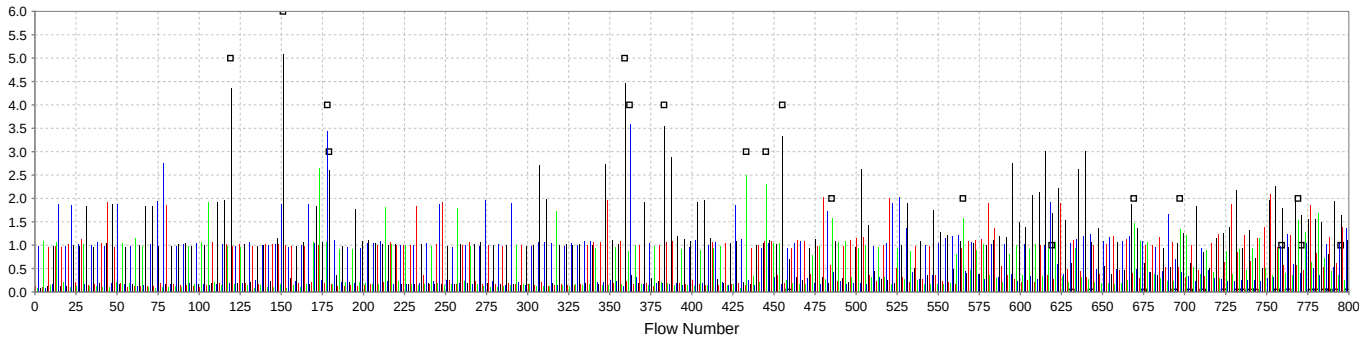


AVTF2



AVTF7



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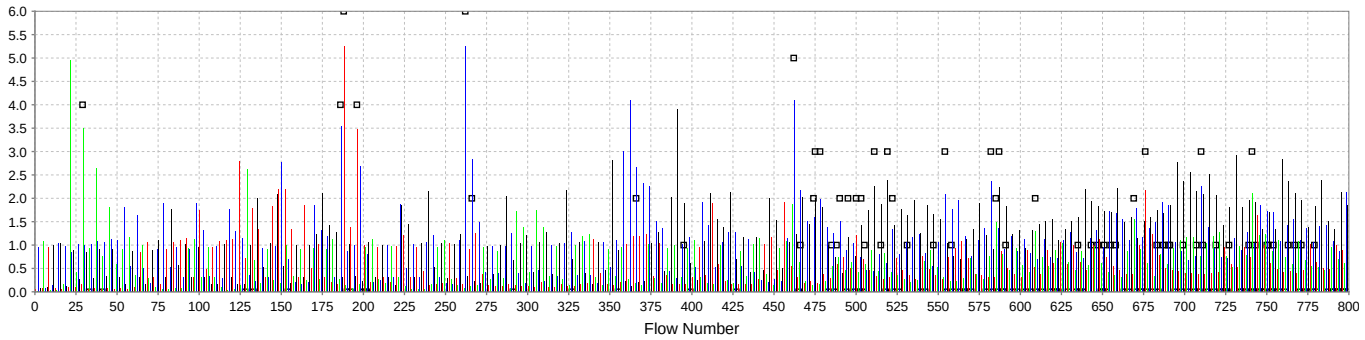

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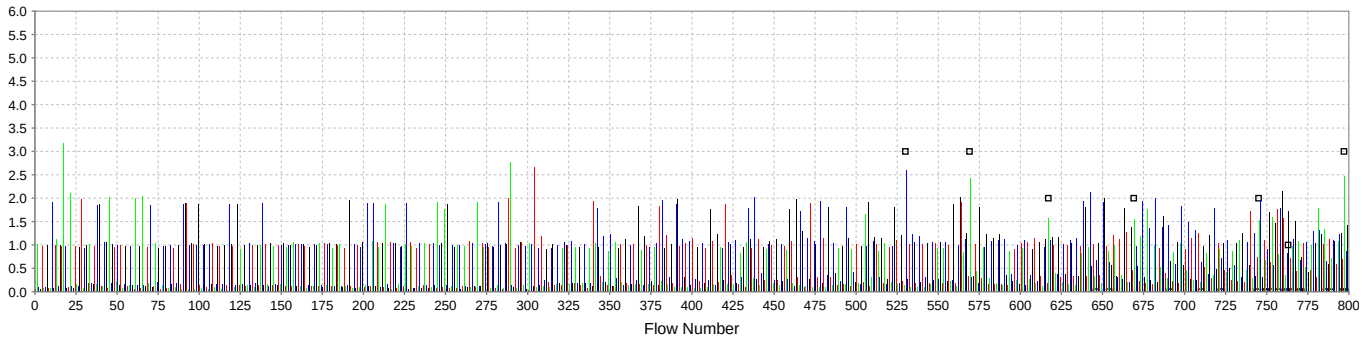

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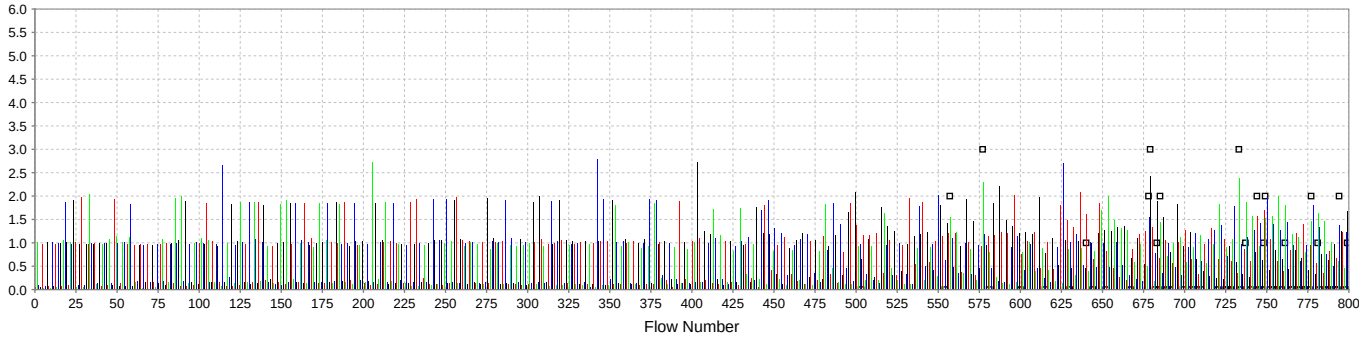

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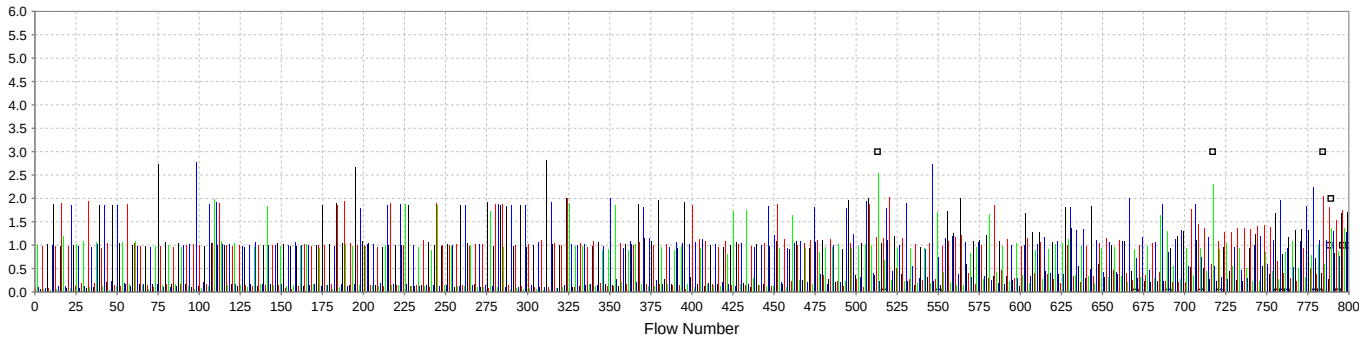

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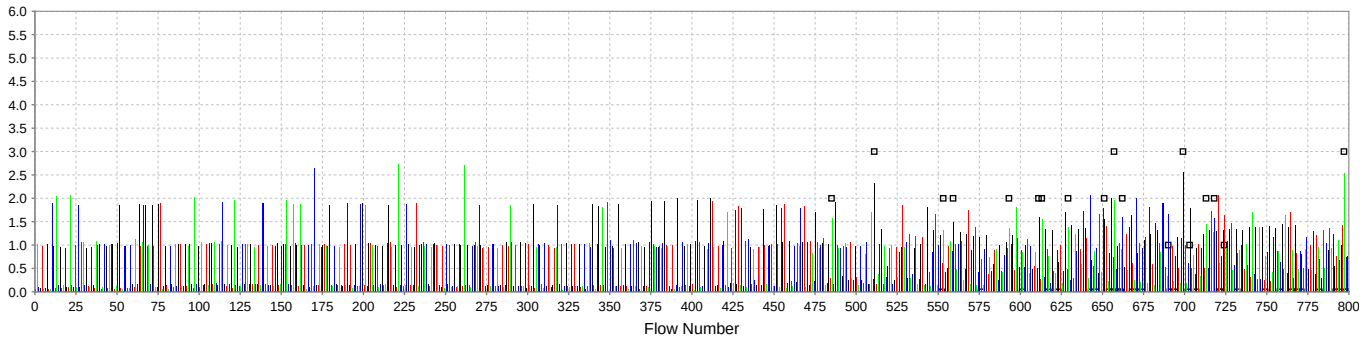
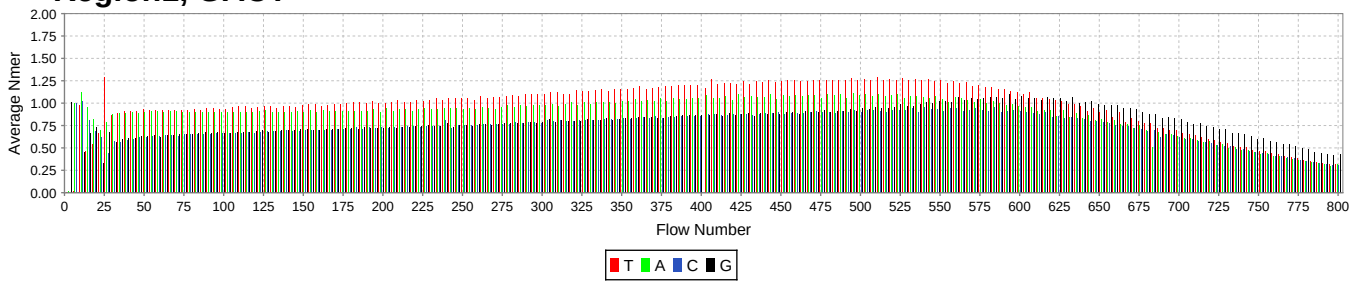
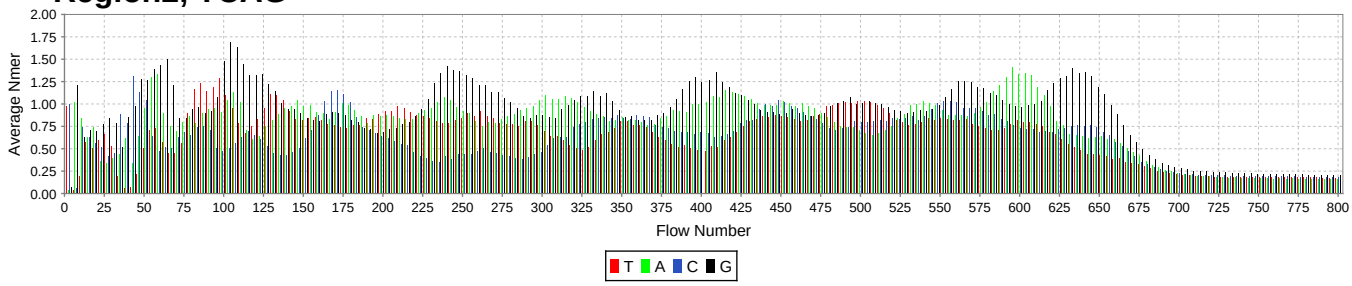


Library Consensus Flowgrams

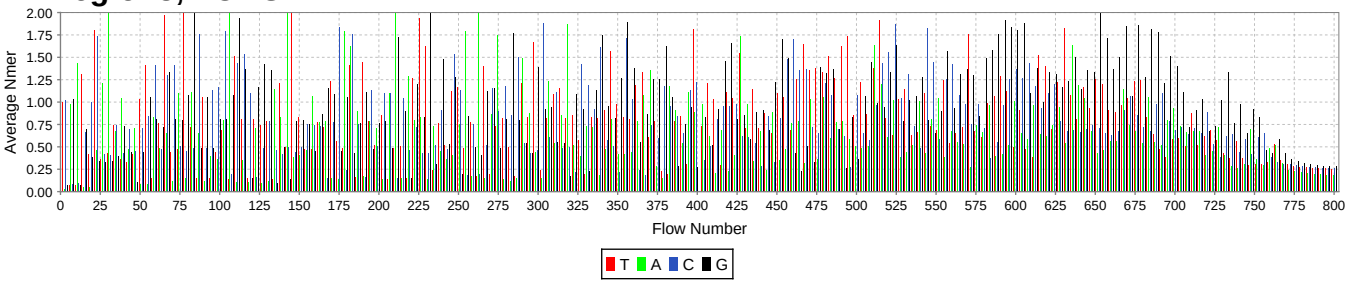
Region1, GACT



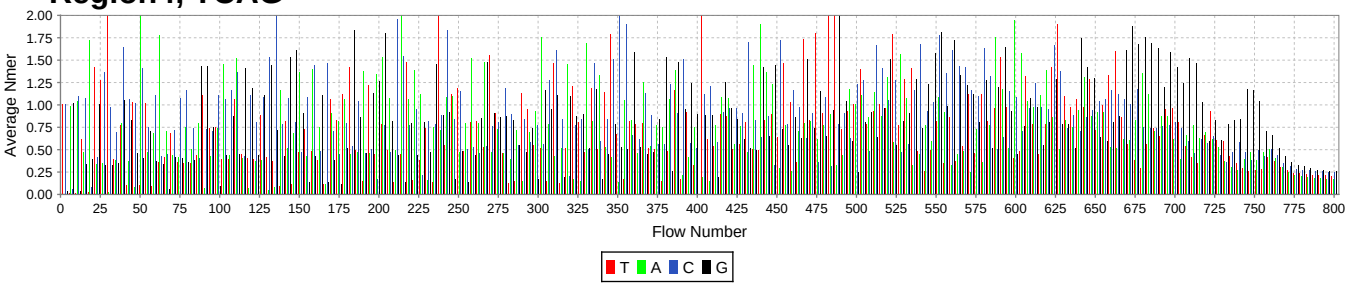
Region2, TCAG



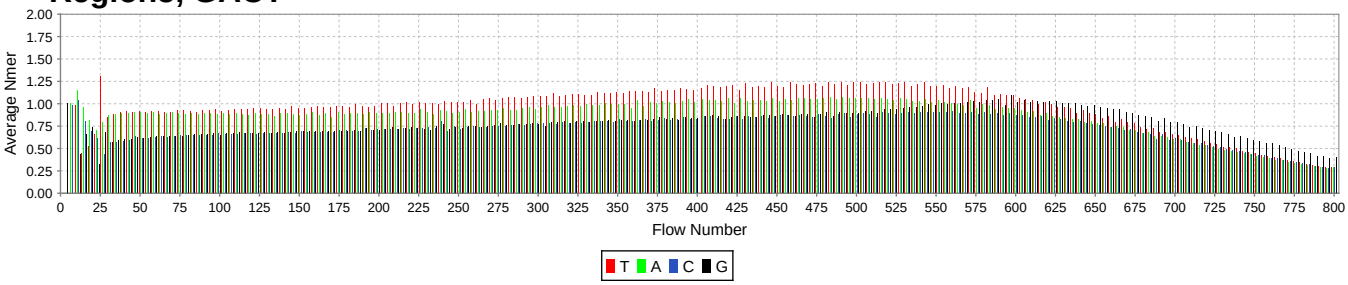
Region3, TCAG



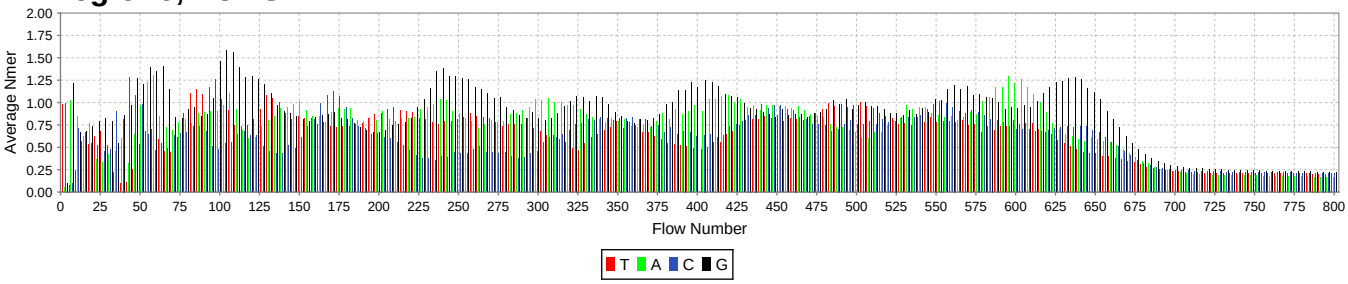
Region4, TCAG



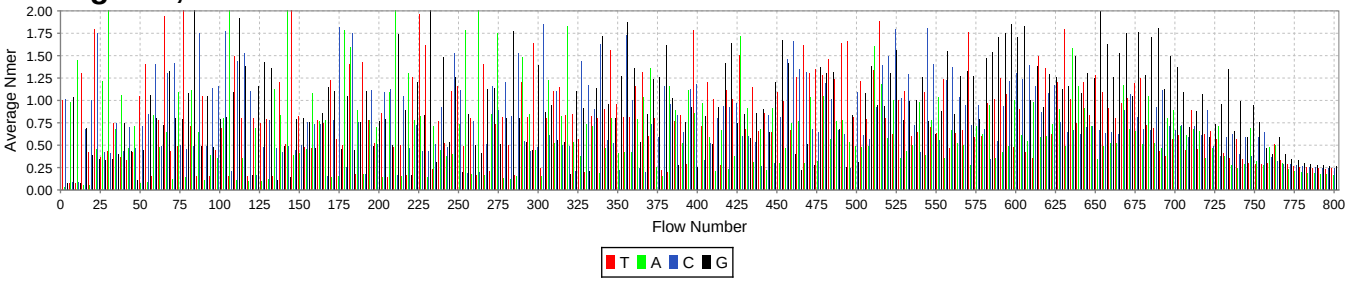
Region5, GACT



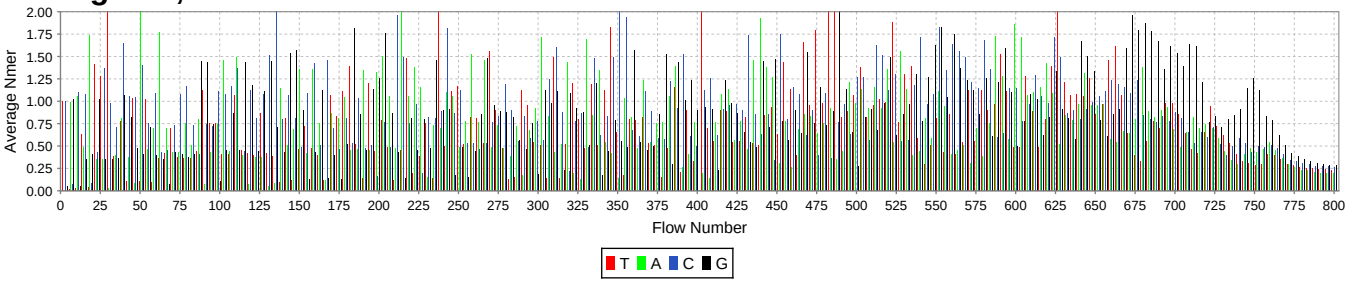
Region6, TCAG



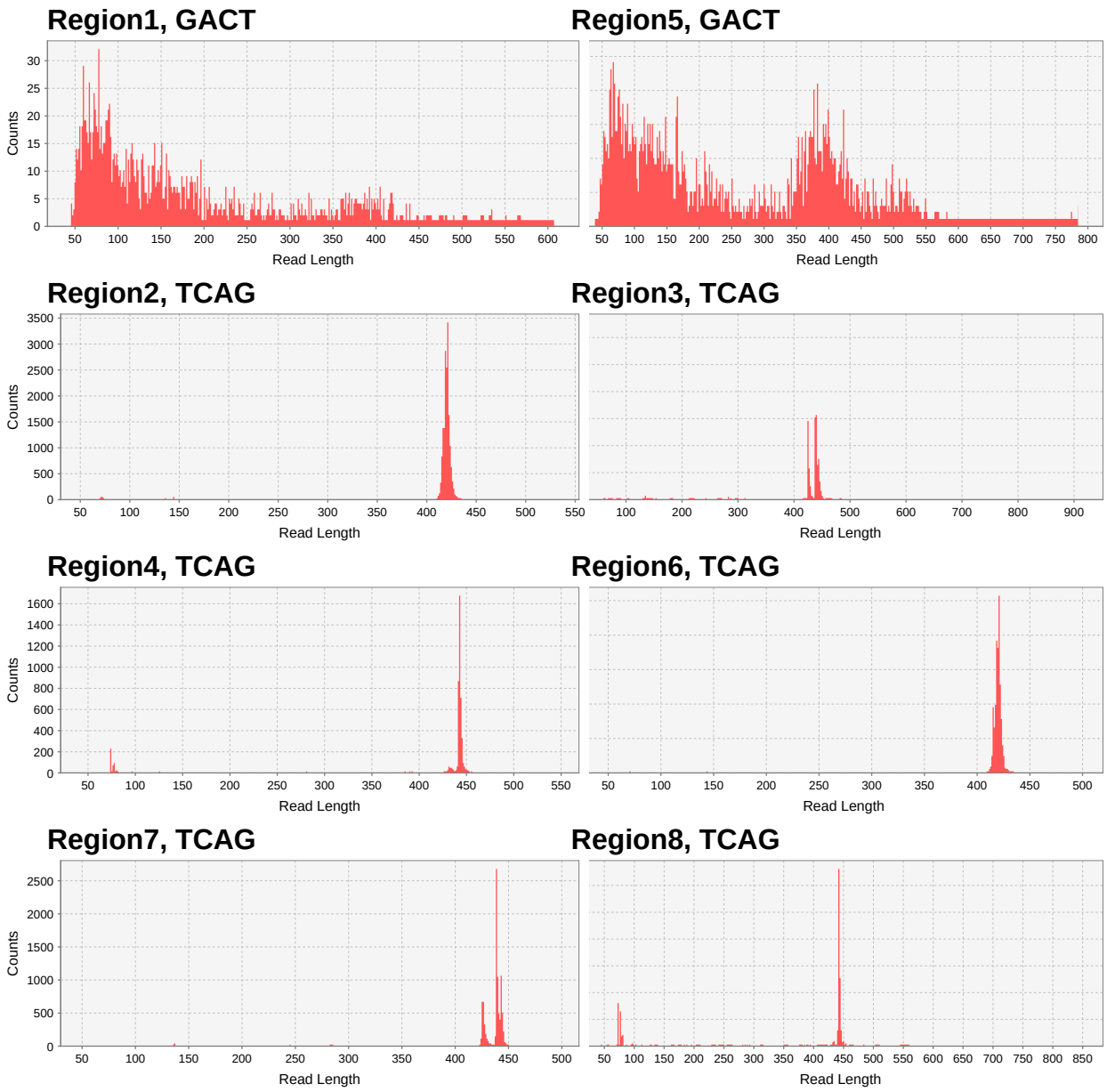
Region7, TCAG



Region8, TCAG



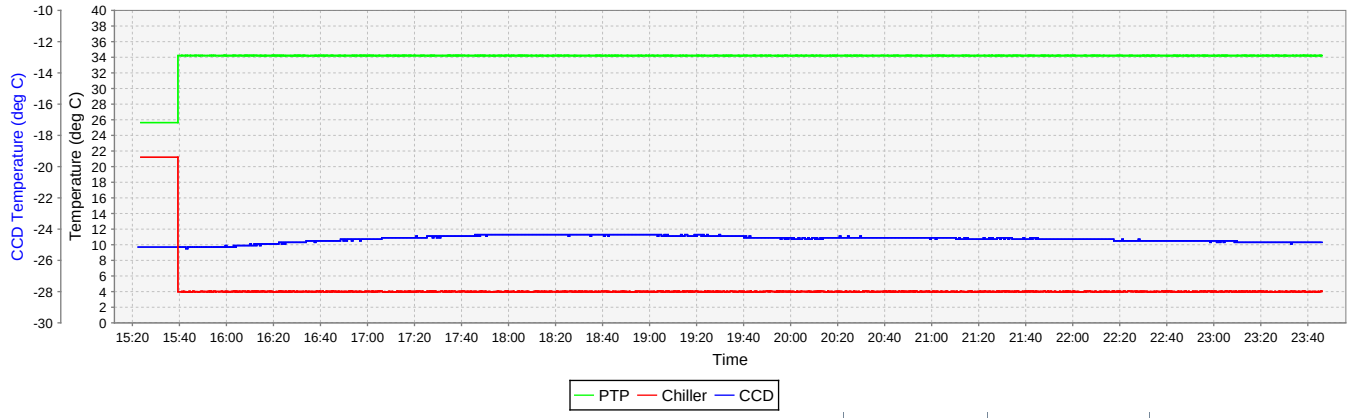
Read Length Distributions



Library Sequence Composition By Region

Region	Filter Pass Reads	Trimmed Primer		Quality Primer		No Trimming	
		Reads (% Total)	Average Length (bp)	Reads (% Total)	Average Length (bp)	Reads (% Total)	Average Length (bp)
1	2365	500 (21.14 %)	316	1805 (76.32 %)	149	-	-
2	20658	805 (3.90 %)	433	19831 (96.00 %)	420	-	-
3	3458	19 (.55 %)	449	3418 (98.84 %)	429	-	-
4	5351	4 (.07 %)	516	5341 (99.81 %)	408	-	-
5	3600	1332 (37.00 %)	380	2210 (61.39 %)	202	43 (1.19 %)	556
6	27553	1674 (6.08 %)	430	25764 (93.51 %)	421	1 (.00 %)	587
7	10102	36 (.36 %)	435	9850 (97.51 %)	434	-	-
8	3142	2 (.06 %)	772	3132 (99.68 %)	343	-	-

Rig State

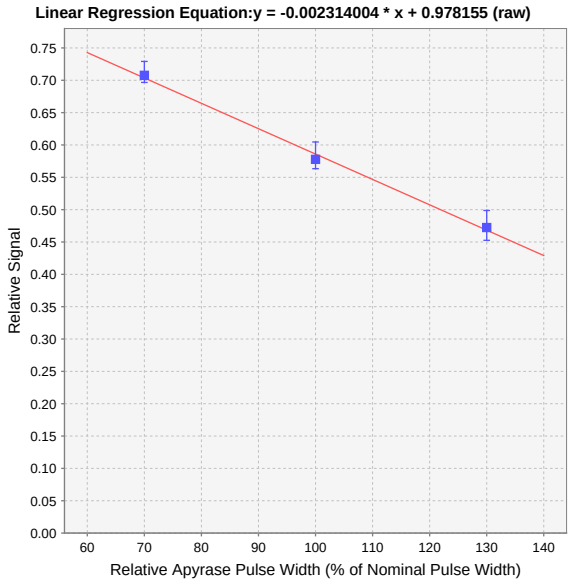


Parameter	Min	Max	Mean
Camera Pressure (Torr)	.92	.92	.92

Calibration Status

No calibration information is found

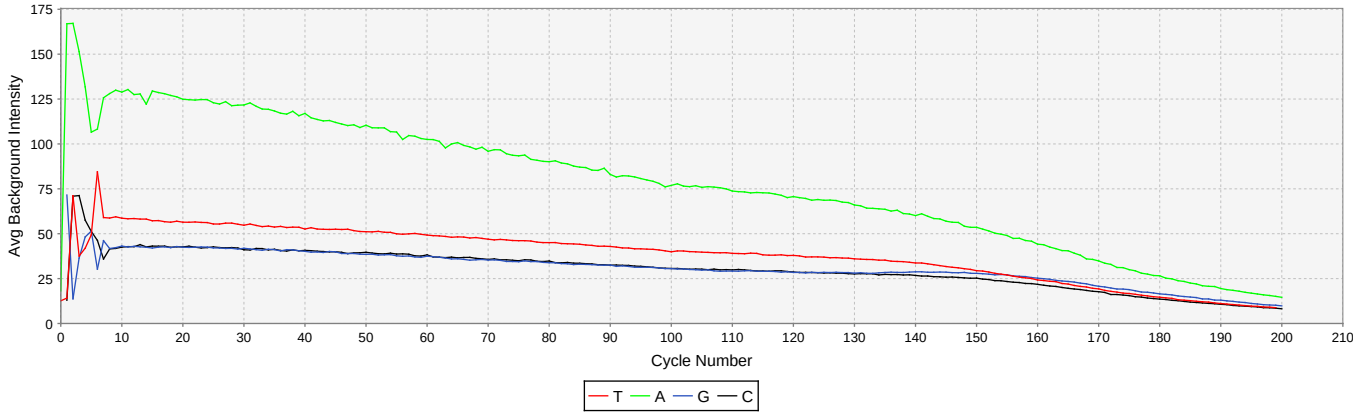
Adaptive Apyrase



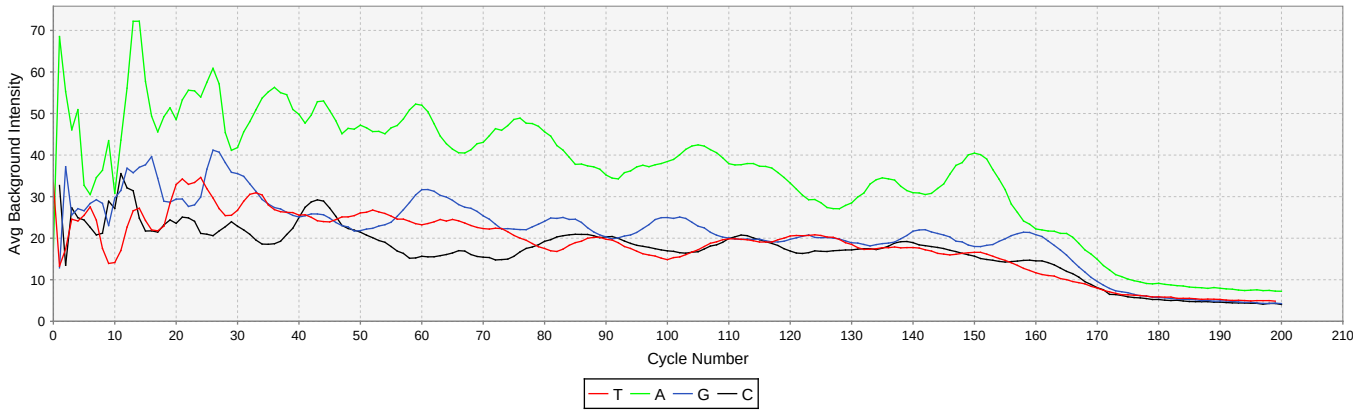
Parameter	Pulse Width in msec	Concentration
Nominal	236.7	1.0X
Calibrated	101.9	0.43 X
Actual	189.0	0.799 X

Background Levels

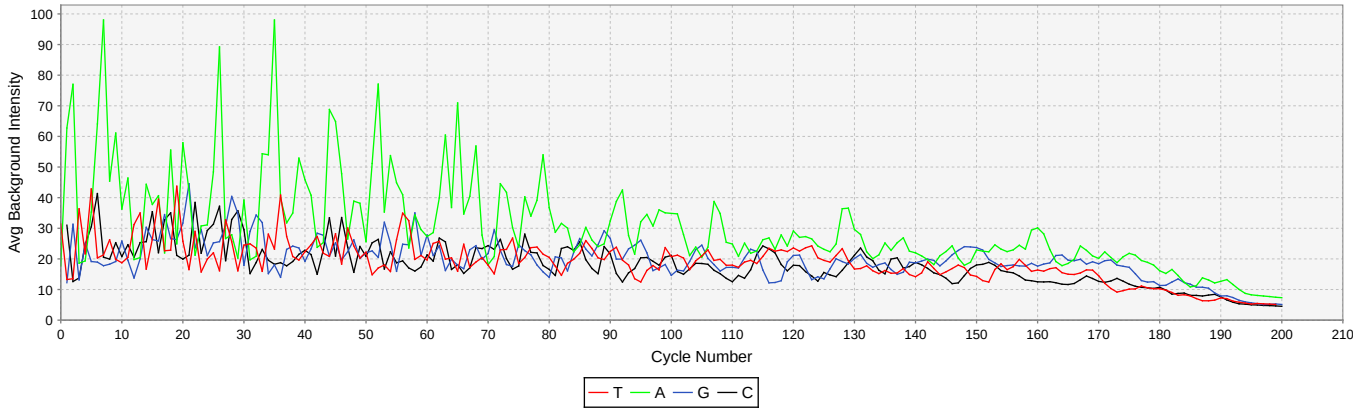
Region 1



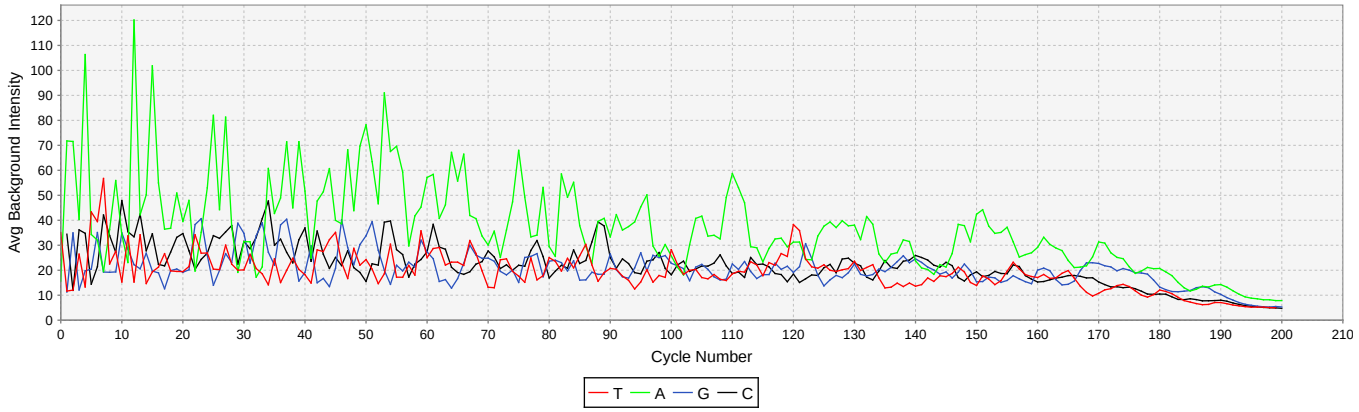
Region 2



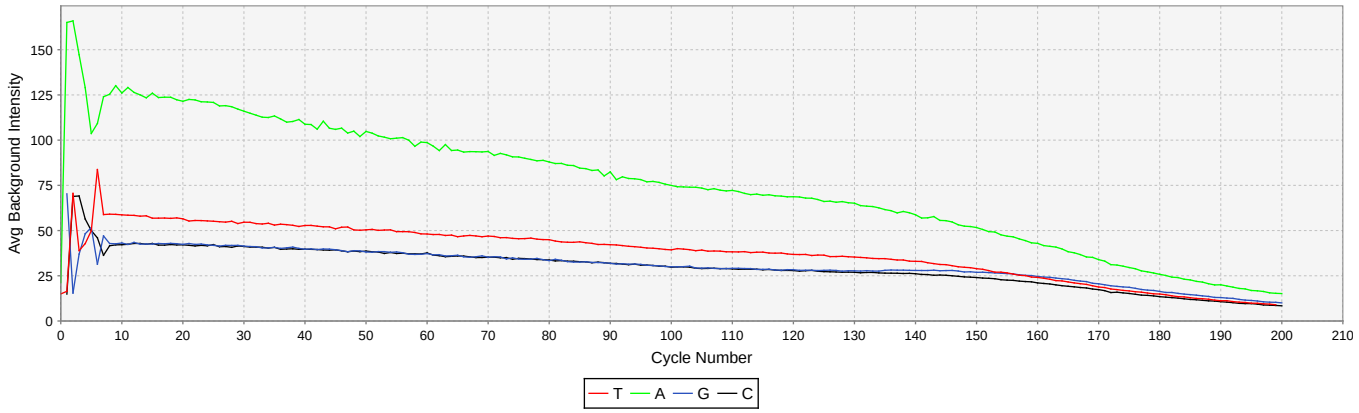
Region 3



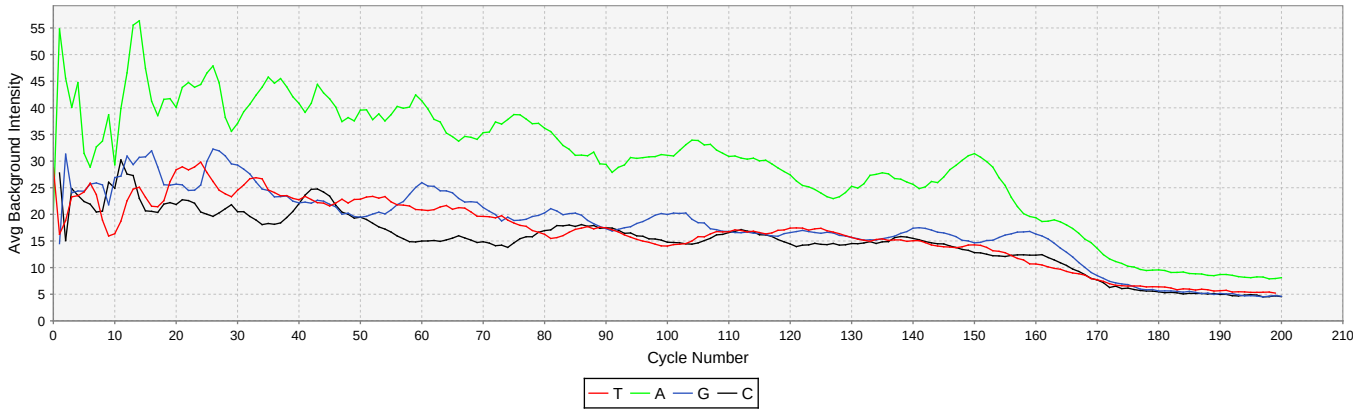
Region 4



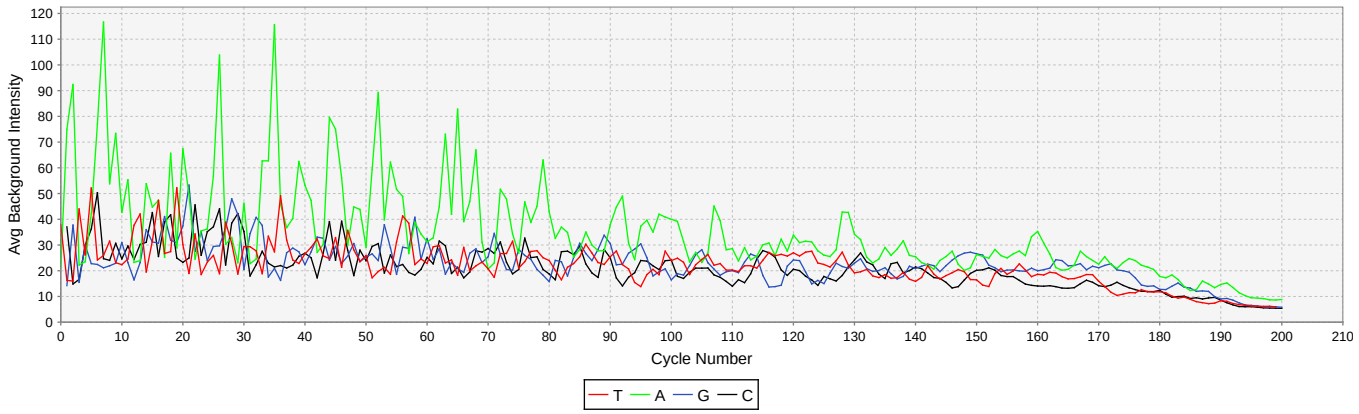
Region 5



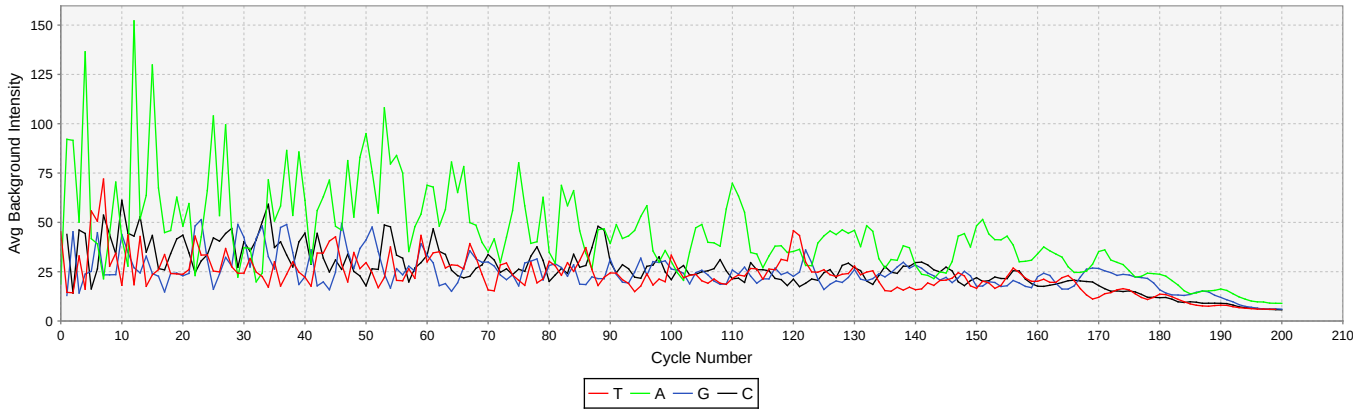
Region 6



Region 7



Region 8



Background Intensity: Mean Value

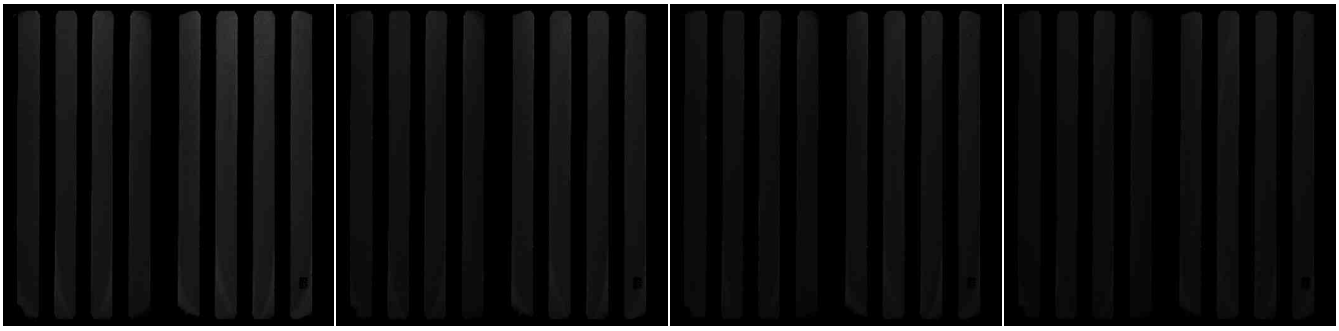
Nucleotide	Region 1	Region 2	Region 3	Region 4	Region 5	Region 6	Region 7	Region 8
T	39	18	19	19	38	17	22	23
A	78	36	30	37	76	30	35	44
C	30	17	18	22	30	15	21	26
G	31	21	20	20	31	18	23	24

Background Intensity Values at First Five Cycles

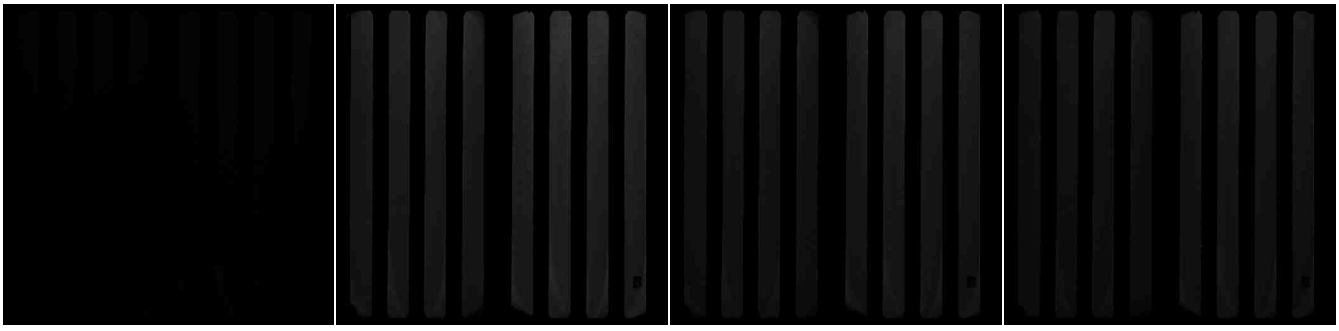
Nucleotide	Flow Number	Region 1	Region 2	Region 3	Region 4	Region 5	Region 6	Region 7	Region 8
T	2	13	33	31	35	15	29	38	45
A	3	18	17	18	16	21	21	21	19
C	4	13	33	31	34	15	28	37	44
G	5	72	13	12	11	70	15	14	13
T	6	14	13	13	12	16	16	16	15
A	7	167	69	63	72	165	55	75	92
C	8	71	13	13	12	69	15	15	14
G	9	14	37	31	35	15	31	38	45
T	10	71	17	14	12	71	19	16	14
A	11	167	55	77	72	166	46	92	92
C	12	71	27	14	36	69	25	17	46
G	13	36	26	13	12	37	24	15	14
T	14	38	25	36	26	39	23	44	33
A	15	151	46	19	40	147	40	22	50
C	16	58	25	25	35	56	24	30	44
G	17	48	27	25	20	48	24	30	24
T	18	42	24	22	13	43	24	26	16
A	19	132	51	19	106	129	45	23	136
C	20	51	24	30	14	50	22	36	16
G	21	51	27	19	20	52	24	23	25

Raw Images

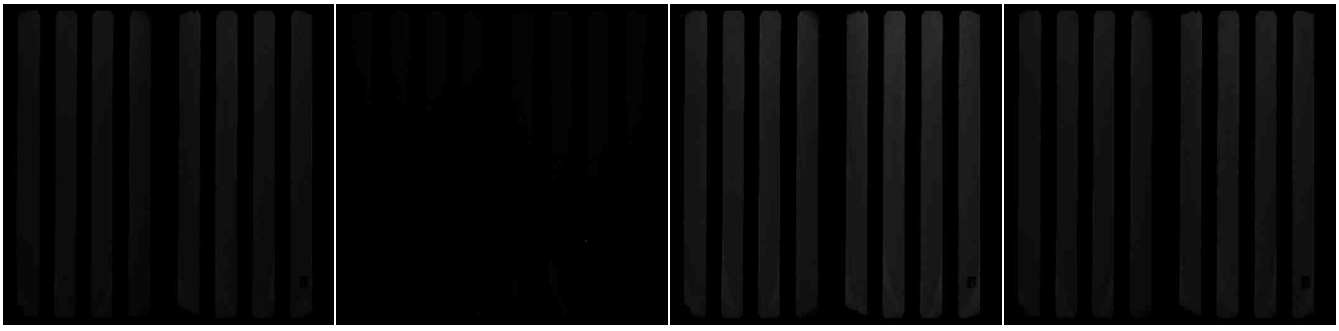
00000: Substrate (--) 00001: Substrate (--) 00002: Substrate (--) 00003: Substrate (--)



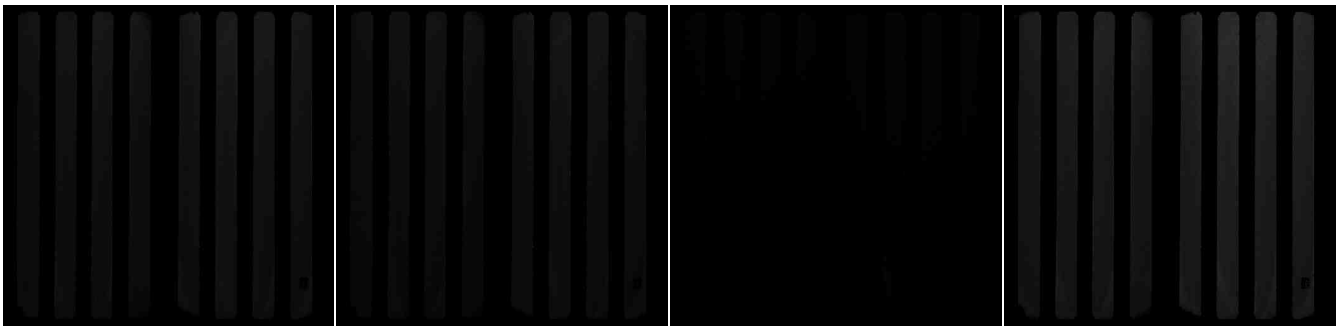
00004: Other (--) 00005: Other (--) 00006: Other (--) 00007: Other (--)



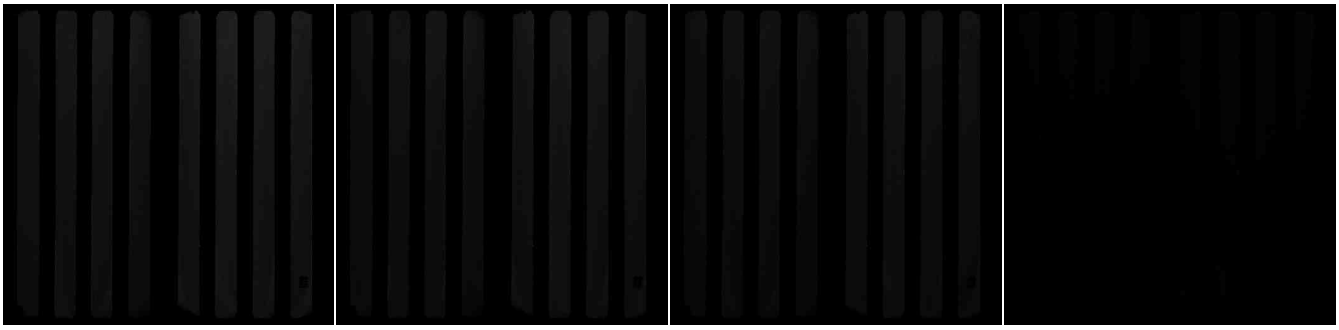
00008: Substrate (--) 00009: Other (--) 00010: Other (--) 00011: Other (--)



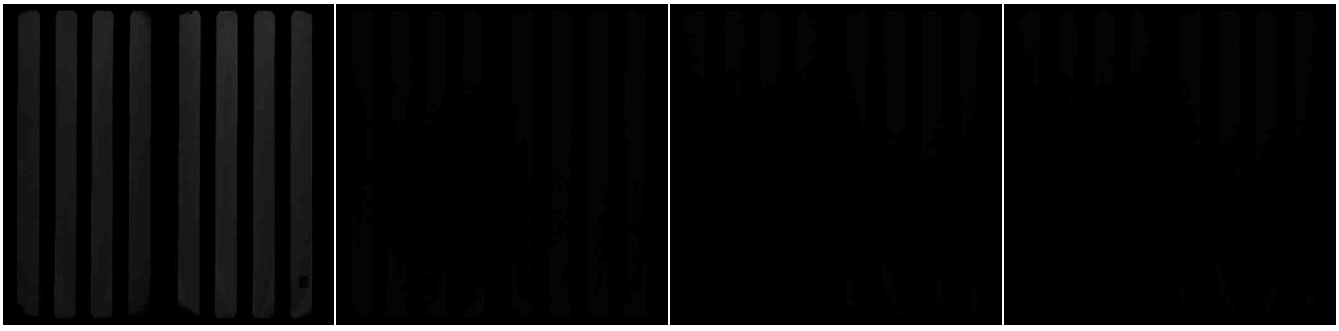
00012: Other (--) 00013: Substrate (--) 00014: Other (--) 00015: Other (--)



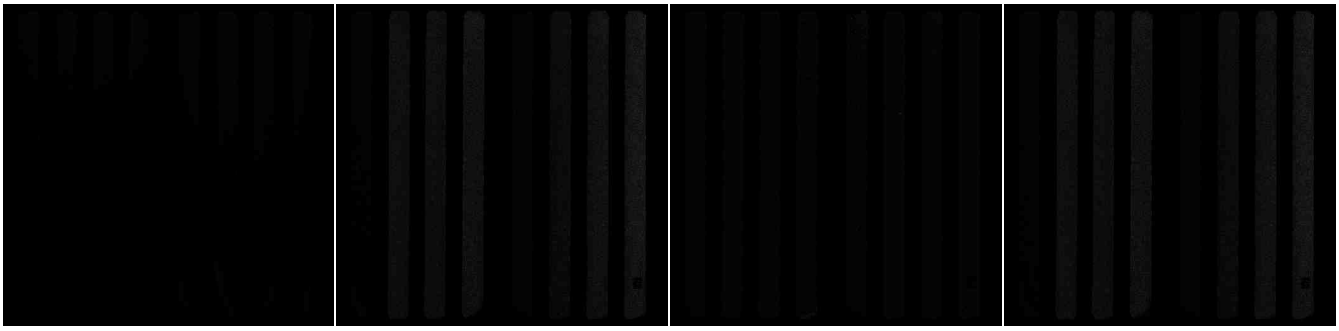
00016: Other (--) 00017: Other (--) 00018: Substrate (--) 00019: Other (--)



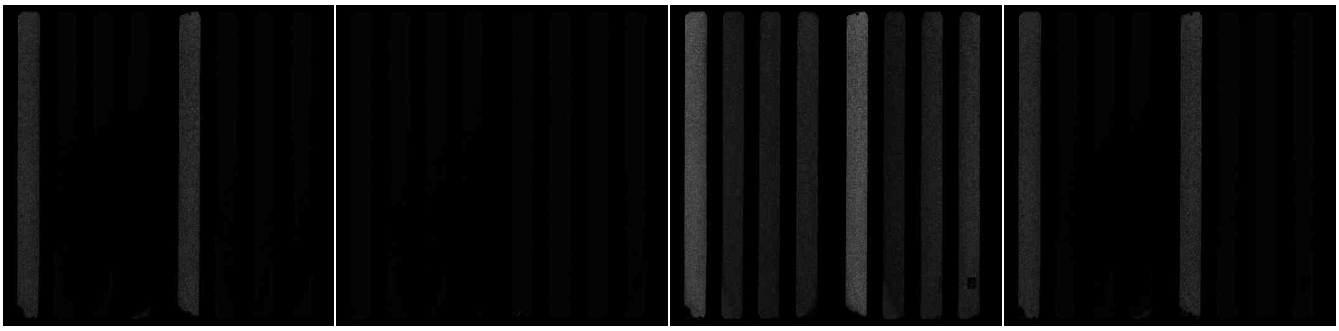
00020: Other (--) 00021: Other (--) 00022: Other (--) 00023: Substrate (--)



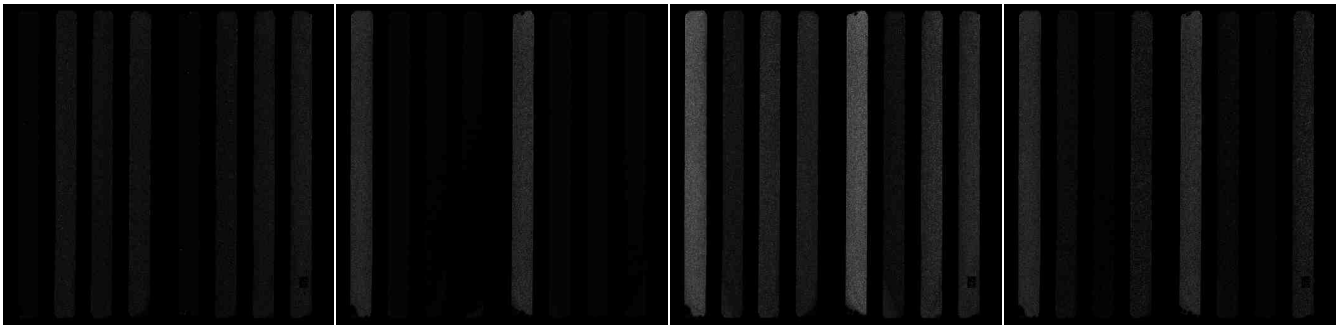
00024: Normalization (0) 00025: Substrate (--) 00026: Substrate (--) 00027: Substrate (--)



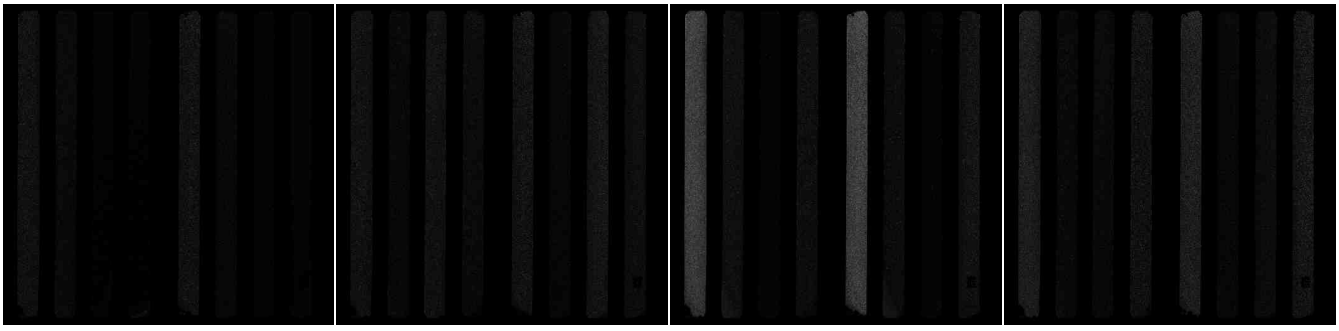
00028: Substrate (--) 00029: T (1) 00030: A (2) 00031: C (3)



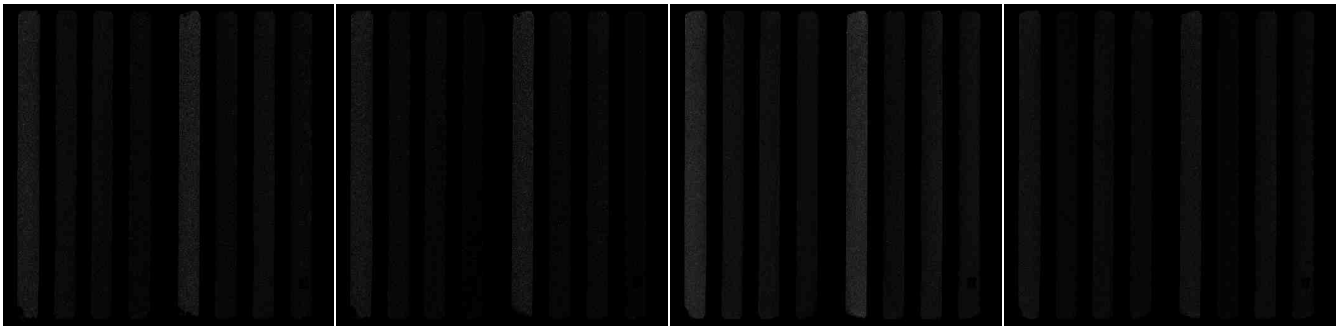
00032: G (4) 00033: T (5) 00034: A (6) 00035: C (7)



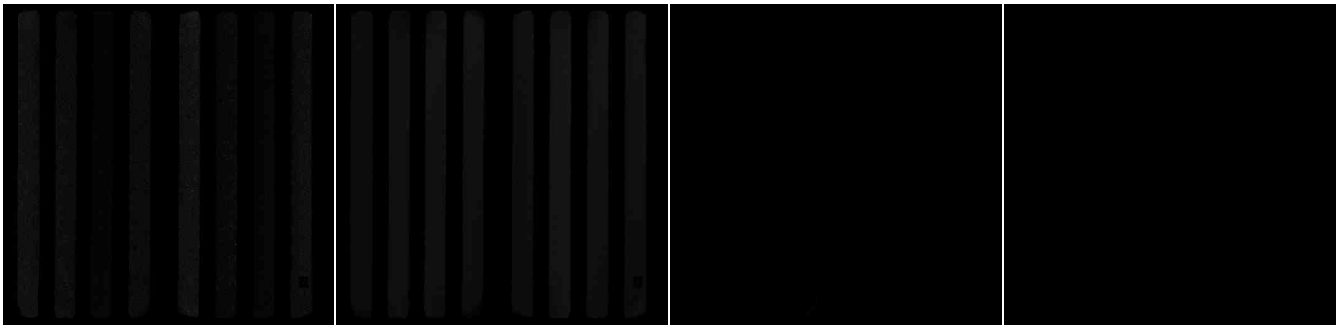
00036: G (8) 00037: T (9) 00038: A (10) 00039: C (11)



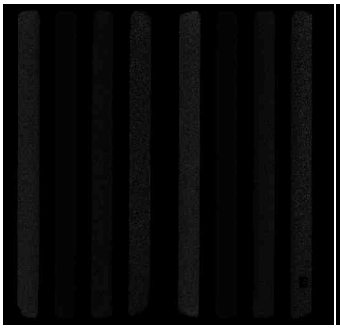
00040: G (12) 00041: T (13) 00042: A (14) 00043: C (15)



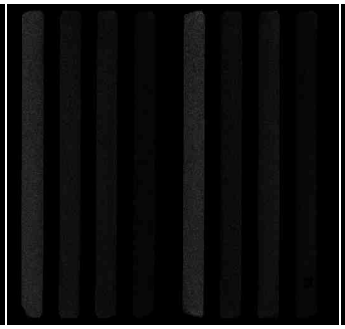
00044: G (16) 00045: T (17) 00426: A (398) 00427: C (399)



00428: G (400)



00429: Normalization (401)



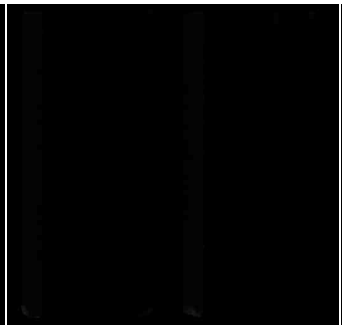
00430: Substrate (--)

00431: Substrate (--)

00432: T (402)

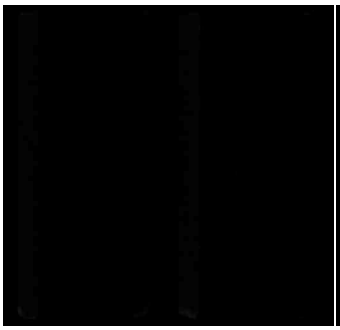
00433: A (403)

00824: T (794)



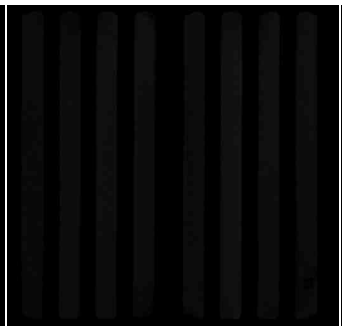
00825: A (795)

00826: C (796)



00827: G (797)

00828: T (798)



00829: A (799)

00830: C (800)

00831: G (801)

00832: Normalization (802)

00833: Substrate (--)

Details

Report generator v2.6