Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electrophoresis File Run Summary

Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

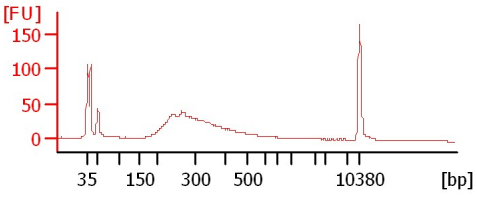
Assay Information:

Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy
Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

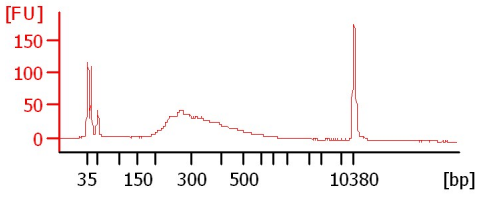
Chip Information:

Chip Lot #:
Reagent Kit Lot #:
Chip Comments:

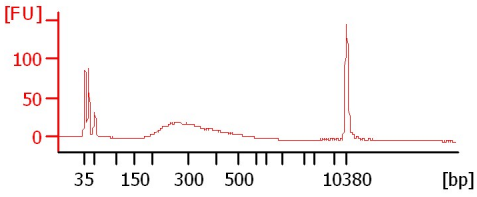
Mac3 library



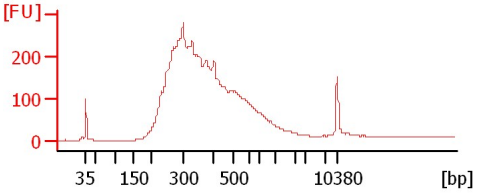
G7T2-3 library



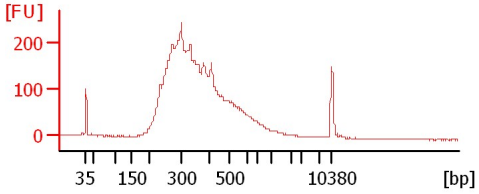
ACP-NK library



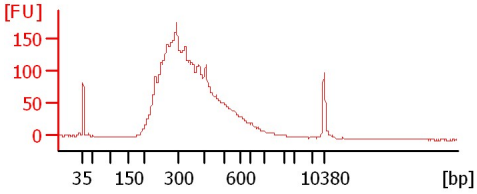
INU-5 library



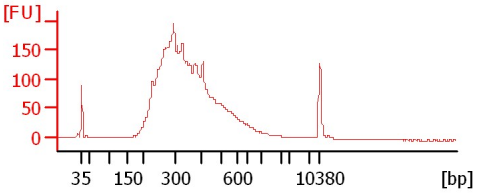
INU-6 library



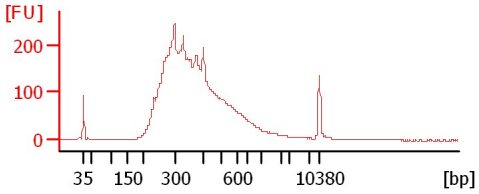
GLU-1 library



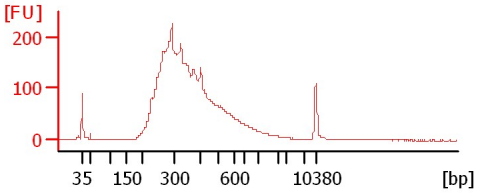
GLU-2 library



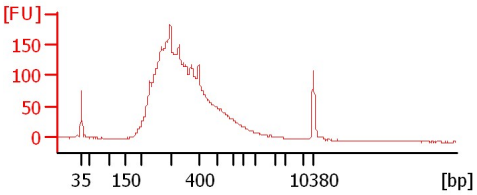
GLU-3 library



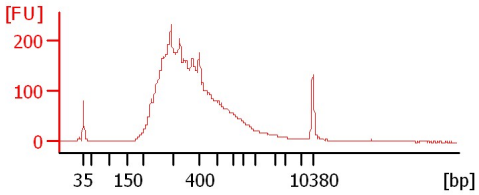
GLU-5 library



GLU-6 library



sample 11



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Mac3 library		☐	✔			
G7T2-3 library		☐	✔			
ACP-NK library		☐	✔			
INU-5 library		☐	✔			
INU-6 library		☐	✔			
GLU-1 library		☐	✔			
GLU-2 library		☐	✔			
GLU-3 library		☐	✔			
GLU-5 library		☐	✔			
GLU-6 library		☐	✔			
sample 11		☐	✔			
Ladder		☐	✔			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.) : 12
Minimum Visible Range [s] : 32
Maximum Visible Range [s] : 138
Start Analysis Time Range [s] : 33
End Analysis Time Range [s] : 137.5
Ladder Concentration [pg/µl] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/µl] : 125
Upper Marker Concentration [pg/µl] : 75
Used Upper Marker for Quantitation
Standard Curve Fit is Point to Point
Show Data Aligned to Lower and Upper Marker

Integrator Settings

Integration Start Time [s] : 33.05
Integration End Time [s] : 137
Slope Threshold : 0.8
Height Threshold [FU] : 5
Area Threshold : 0.1
Width Threshold [s] : 0.6
Baseline Plateau [s] : 0.5

Filter Settings

Filter Width [s] : 0.5
Polynomial Order : 4

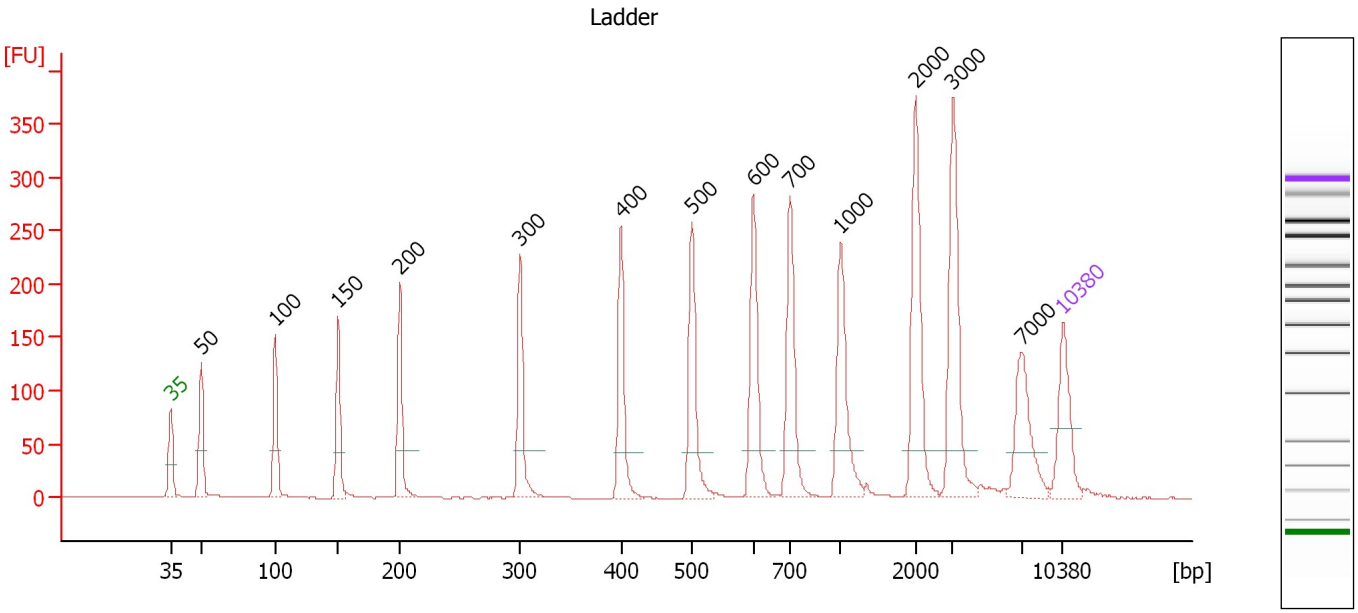
Ladder

Ladder Peak	Size	Area
1	35	160
2	50	210
3	100	208
4	150	221
5	200	242
6	300	270
7	400	305
8	500	306
9	600	336
10	700	321
11	1000	366
12	2000	413
13	3000	411
14	7000	400
15	10380	214

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

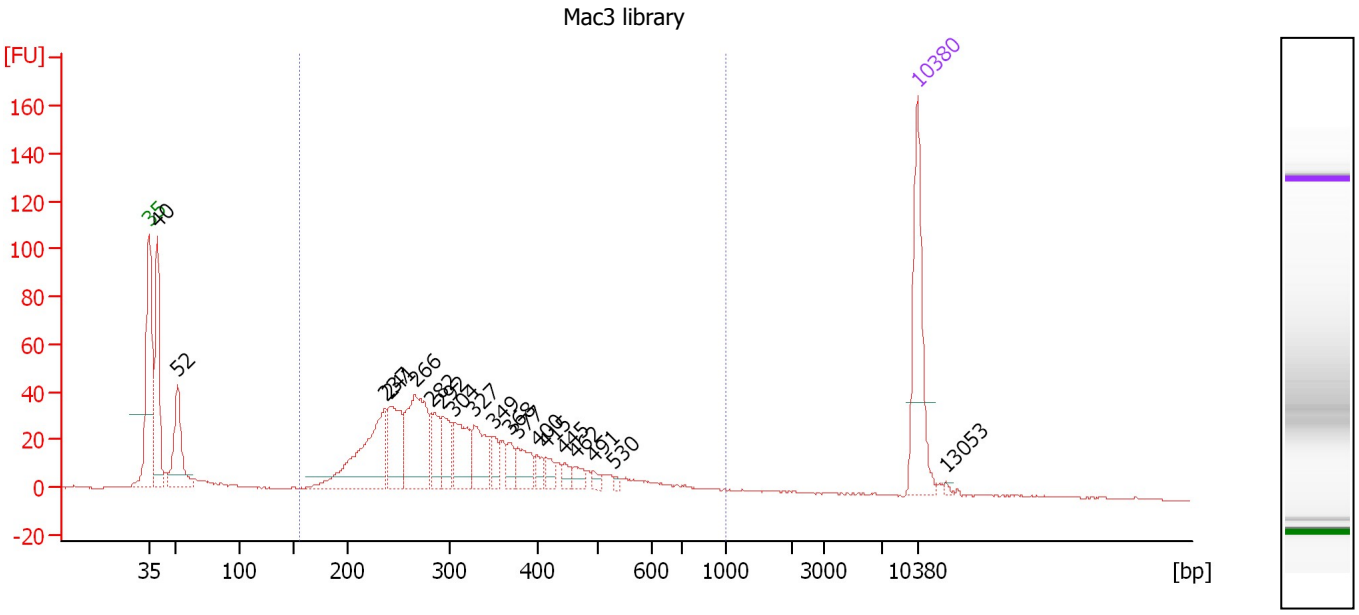
Peak table for Ladder

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1		35	125.00	5,411.3	Lower Marker	43.00
2		50	150.00	4,545.5	Ladder Peak	45.39
3		100	150.00	2,272.7	Ladder Peak	51.19
4		150	150.00	1,515.2	Ladder Peak	56.17
5		200	150.00	1,136.4	Ladder Peak	61.00
6		300	150.00	757.6	Ladder Peak	70.44
7		400	150.00	568.2	Ladder Peak	78.34
8		500	150.00	454.5	Ladder Peak	83.89
9		600	150.00	378.8	Ladder Peak	88.73
10		700	150.00	324.7	Ladder Peak	91.60
11		1,000	150.00	227.3	Ladder Peak	95.57
12		2,000	150.00	113.6	Ladder Peak	101.46
13		3,000	150.00	75.8	Ladder Peak	104.38
14		7,000	150.00	32.5	Ladder Peak	109.74
15		10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Mac3 library

Number of peaks found:	19	Corr. Area 1:	886.4
Noise:	0.3		

Peak table for sample 1 : Mac3 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	40	135.48	5,188.2		43.73
3	52	89.54	2,631.2		45.57
4	237	131.37	840.7		64.47
5	241	74.18	466.6		64.86
6	266	113.62	646.9		67.24
7	282	36.76	197.5		68.75
8	292	32.06	166.1		69.72
9	304	51.71	257.9		70.74
10	327	49.70	230.2		72.58
11	349	19.28	83.6		74.33
12	368	15.31	63.0		75.84
13	377	27.92	112.2		76.52
14	400	11.12	42.2		78.32
15	415	11.76	43.0		79.14
16	445	8.54	29.1		80.84
17	462	9.77	32.1		81.76
18	491	6.20	19.1		83.42
19	530	2.93	8.4		85.36
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,053	0.00	0.0		115.57

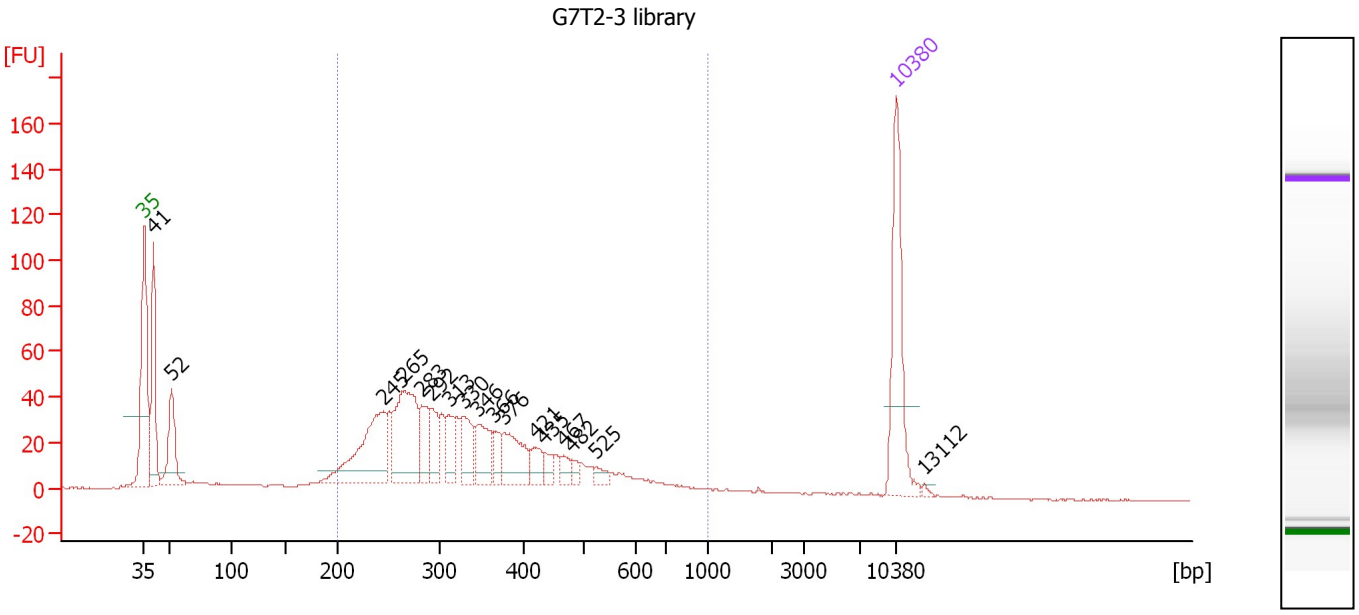
Region table for sample 1 : Mac3 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
156	1,000	335	720.08	886.4	3,726.4	74	36.3

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : G7T2-3 library

Number of peaks found:	17	Corr. Area 1:	996.7
Noise:	0.3		

Peak table for sample 2 : G7T2-3 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	41	111.53	4,162.4		43.89
3	52	68.52	2,011.3		45.58
4	245	111.85	690.4		65.29
5	265	107.79	615.2		67.18
6	283	37.23	199.1		68.87
7	292	28.89	149.8		69.71
8	313	27.62	133.8		71.45
9	330	34.51	158.6		72.79
10	346	36.74	160.8		74.08
11	366	14.53	60.2		75.62
12	376	44.29	178.3		76.46
13	421	16.06	57.8		79.49
14	435	10.02	34.9		80.28
15	467	10.60	34.4		82.07
16	482	5.45	17.1		82.91
17	525	7.53	21.7		85.10
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,112	0.00	0.0		115.63

Region table for sample 2 : G7T2-3 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	354	697.40	996.7	3,387.4	76	34.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
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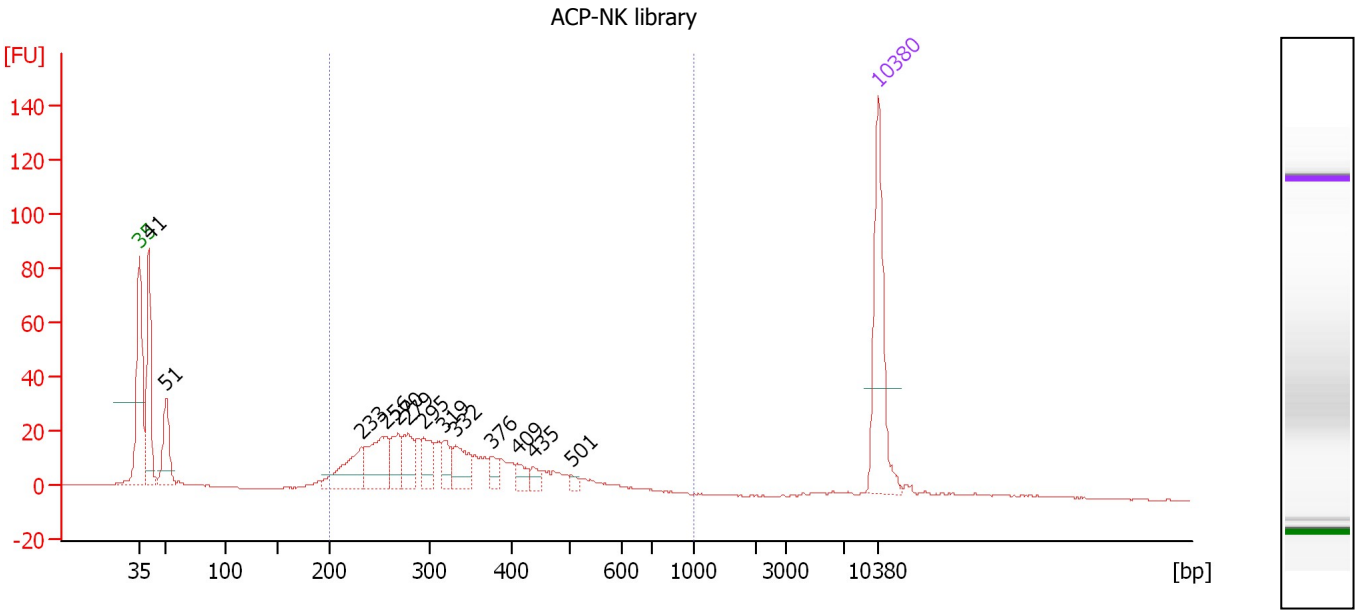
Electropherogram Summary Continued ...



Overall Results for sample 3 : ACP-NK library

Number of peaks found:	13	Corr. Area 1:	469.8
Noise:	0.2		

Peak table for sample 3 : ACP-NK library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	41	108.53	4,040.1		43.91
3	51	58.31	1,727.3		45.53
4	233	51.62	336.3		64.08
5	256	60.67	358.9		66.30
6	270	29.48	165.7		67.56
7	279	32.81	178.0		68.47
8	295	29.37	151.0		69.94
9	319	18.89	89.8		71.91
10	332	33.64	153.5		72.97
11	376	12.16	49.0		76.46
12	409	11.53	42.7		78.83
13	435	7.79	27.1		80.30
14	501	4.52	13.7		83.94
15	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 3 : ACP-NK library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	336	395.67	469.8	1,966.3	72	29.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

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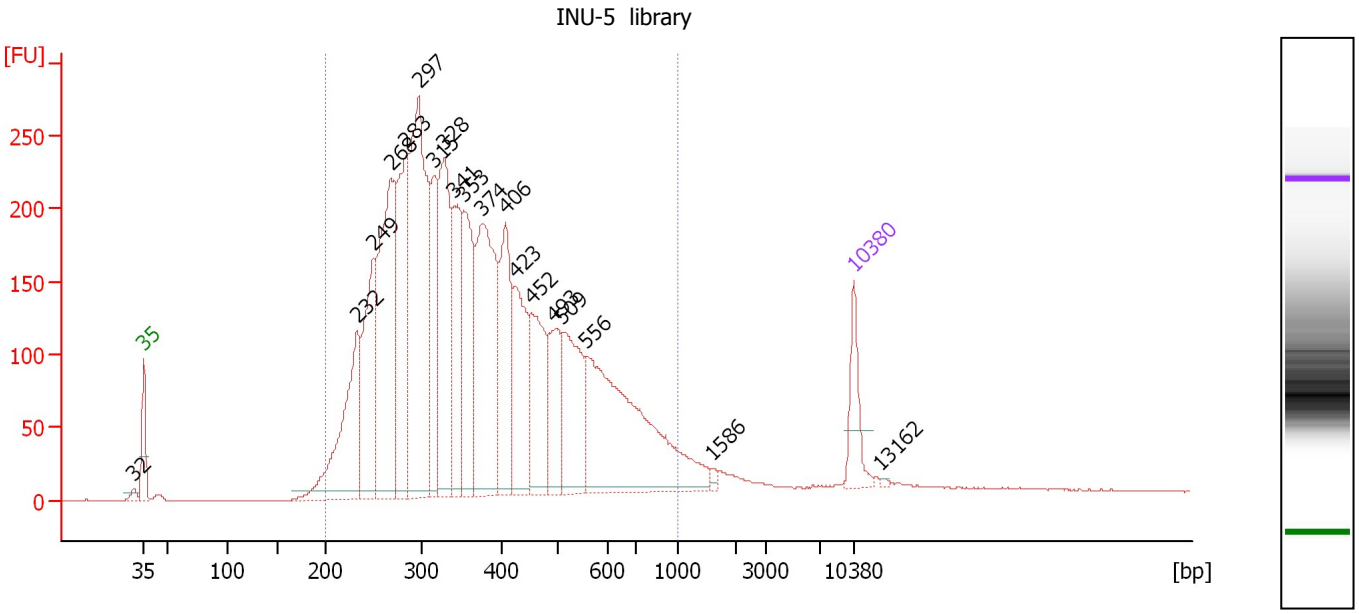
Electropherogram Summary Continued ...



Overall Results for sample 4 : INU-5 library

Number of peaks found:	19	Corr. Area 1:	6,187.6
Noise:	0.2		

Peak table for sample 4 : INU-5 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.05
2	35	125.00	5,411.3	Lower Marker	43.00
3	232	389.36	2,542.9		64.02
4	249	308.62	1,875.9		65.65
5	268	598.55	3,388.3		67.38
6	283	412.10	2,208.7		68.80
7	297	729.43	3,725.9		70.12
8	315	215.35	1,034.9		71.64
9	328	390.31	1,803.5		72.64
10	341	262.74	1,169.0		73.64
11	353	277.47	1,192.6		74.58
12	374	485.60	1,968.3		76.27
13	406	250.99	936.0		78.68
14	423	238.24	852.8		79.63
15	452	218.55	733.1		81.21
16	493	163.82	503.1		83.52
17	509	236.05	703.2		84.31
18	556	587.03	1,598.4		86.62
19	1,586	7.30	7.0		99.02
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,162	0.00	0.0		115.68

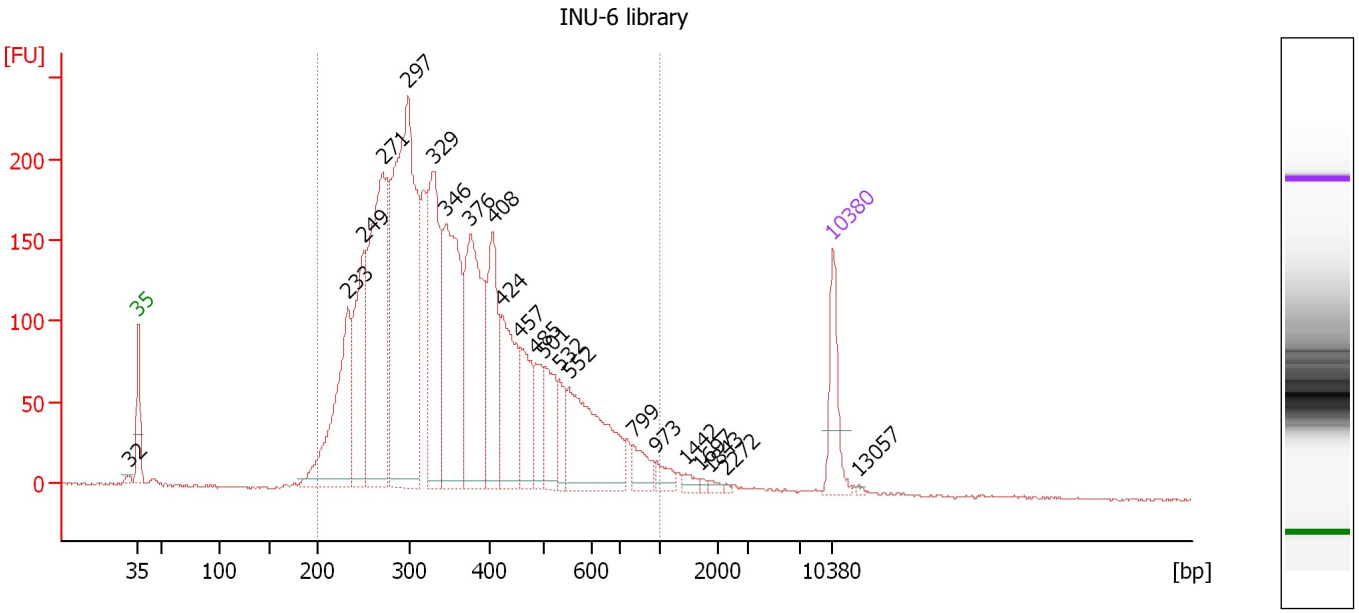
Region table for sample 4 : INU-5 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	390	5,656.81	6,187.6	25,398.5	95	36.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : INU-6 library

Number of peaks found:	22	Corr. Area 1:	4,908.8
Noise:	0.8		

Peak table for sample 5 : INU-6 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.03
2	35	125.00	5,411.3	Lower Marker	43.00
3	233	324.83	2,112.5		64.11
4	249	267.60	1,625.1		65.67
5	271	534.06	2,990.7		67.66
6	297	817.70	4,166.8		70.18
7	329	352.66	1,625.2		72.71
8	346	374.22	1,639.8		74.05
9	376	346.96	1,396.7		76.47
10	408	194.47	722.2		78.78
11	424	185.66	662.8		79.69
12	457	114.18	378.3		81.52
13	485	68.51	213.8		83.08
14	501	100.97	305.4		83.94
15	532	51.31	146.1		85.44
16	552	246.79	677.3		86.41
17	799	40.17	76.2		92.91
18	973	22.36	34.8		95.22
19	1,442	10.42	10.9		98.17
20	1,697	3.87	3.5		99.68
21	1,843	5.75	4.7		100.54
22	2,272	2.09	1.4		102.26
23	10,380	75.00	10.9	Upper Marker	113.00
24	13,057	0.00	0.0		115.58

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...

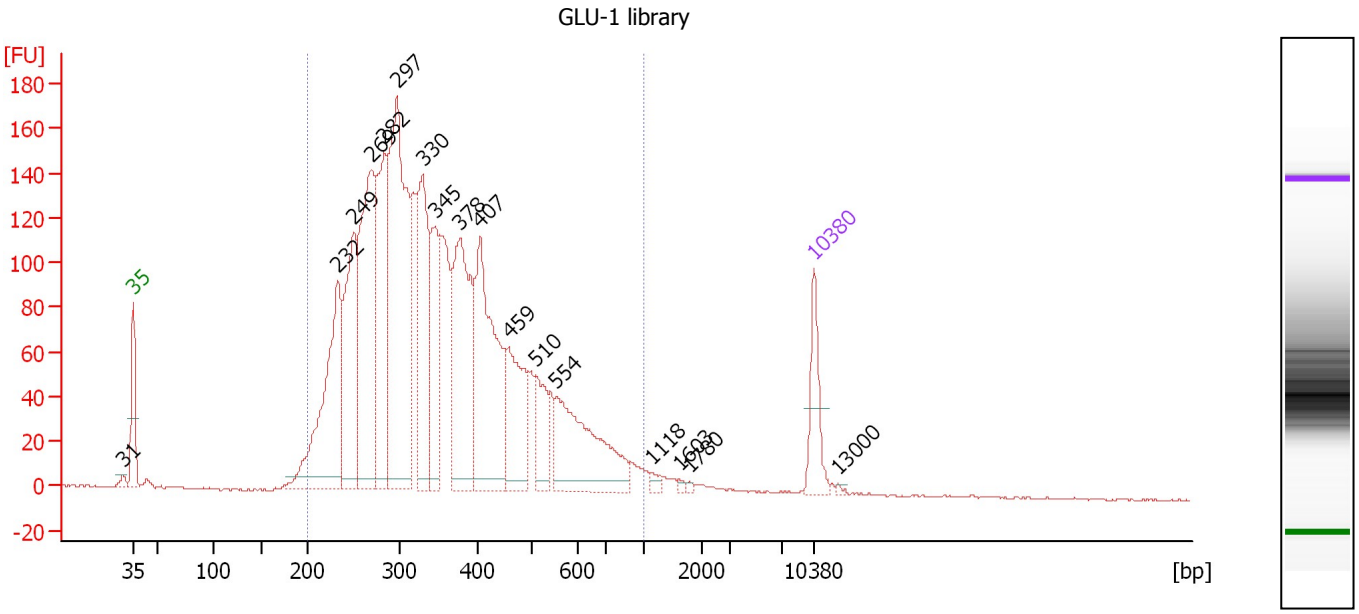
... Region table for sample 5 : INU-6 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	373	4,261.61	4,908.8	19,683.3	97	34.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : GLU-1 library

Number of peaks found:	17	Corr. Area 1:	3,594.9
Noise:	0.4		

Peak table for sample 6 : GLU-1 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.96
2	35	125.00	5,411.3	Lower Marker	43.00
3	232	436.30	2,850.7		64.01
4	249	316.18	1,921.3		65.66
5	269	518.85	2,921.2		67.52
6	282	325.91	1,748.0		68.78
7	297	681.47	3,476.2		70.16
8	330	300.20	1,379.2		72.79
9	345	182.30	800.4		74.00
10	378	352.08	1,412.4		76.57
11	407	398.82	1,485.4		78.71
12	459	176.82	583.5		81.62
13	510	93.19	277.0		84.36
14	554	266.09	727.7		86.50
15	1,118	9.65	13.1		96.27
16	1,603	3.99	3.8		99.12
17	1,780	2.90	2.5		100.16
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,000	0.00	0.0		115.52

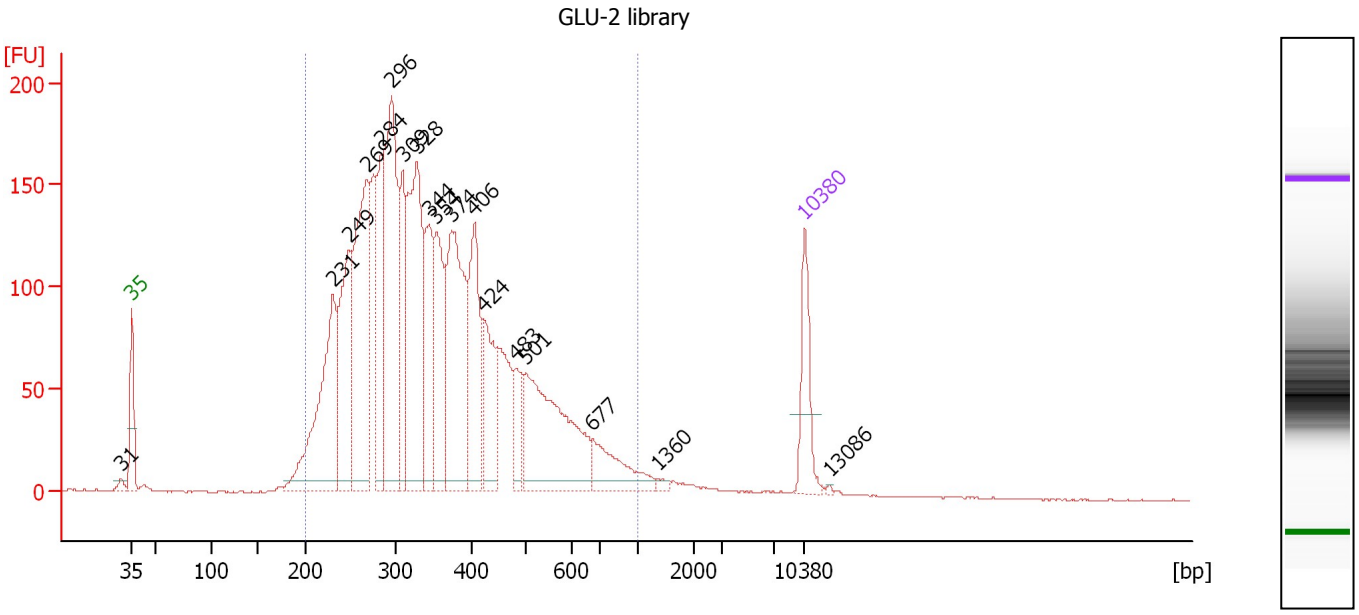
Region table for sample 6 : GLU-1 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	366	4,526.55	3,594.9	21,236.8	97	34.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : GLU-2 library

Number of peaks found:	18	Corr. Area 1:	3,945.5
Noise:	0.4		

Peak table for sample 7 : GLU-2 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.89
2	35	125.00	5,411.3	Lower Marker	43.00
3	231	360.57	2,366.7		63.91
4	249	275.75	1,677.5		65.63
5	269	416.50	2,345.4		67.52
6	284	179.25	955.1		68.96
7	296	427.33	2,186.6		70.07
8	309	156.35	765.7		71.18
9	328	381.64	1,761.0		72.68
10	344	161.78	713.0		73.90
11	354	192.26	823.7		74.67
12	374	328.89	1,332.4		76.28
13	406	199.93	746.2		78.67
14	424	122.26	437.0		79.66
15	483	58.17	182.5		82.94
16	501	312.18	944.2		83.94
17	677	89.77	201.1		90.92
18	1,360	6.11	6.8		97.69
19	10,380	75.00	10.9	Upper Marker	113.00
20	13,086	0.00	0.0		115.61

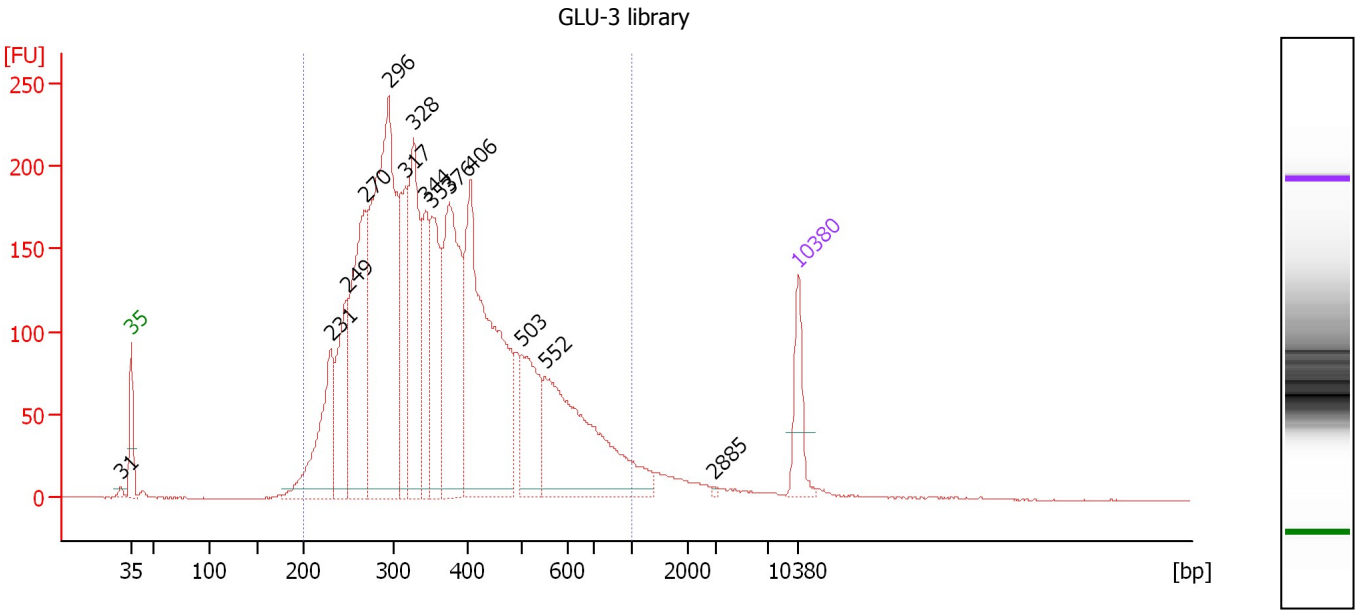
Region table for sample 7 : GLU-2 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	366	3,957.75	3,945.5	18,538.2	97	34.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : GLU-3 library

Number of peaks found:	14	Corr. Area 1:	4,995.4
Noise:	0.4		

Peak table for sample 8 : GLU-3 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.88
2	35	125.00	5,411.3	Lower Marker	43.00
3	231	278.87	1,829.1		63.93
4	249	224.59	1,367.7		65.61
5	270	452.27	2,542.1		67.56
6	296	939.12	4,812.5		70.03
7	317	195.09	933.1		71.76
8	328	351.85	1,624.7		72.66
9	344	191.79	845.5		73.89
10	353	257.39	1,105.1		74.61
11	376	413.84	1,669.6		76.41
12	406	698.61	2,610.0		78.64
13	503	189.41	571.0		84.02
14	552	454.97	1,248.0		86.42
15	2,885	2.72	1.4		104.05
16	10,380	75.00	10.9	Upper Marker	113.00

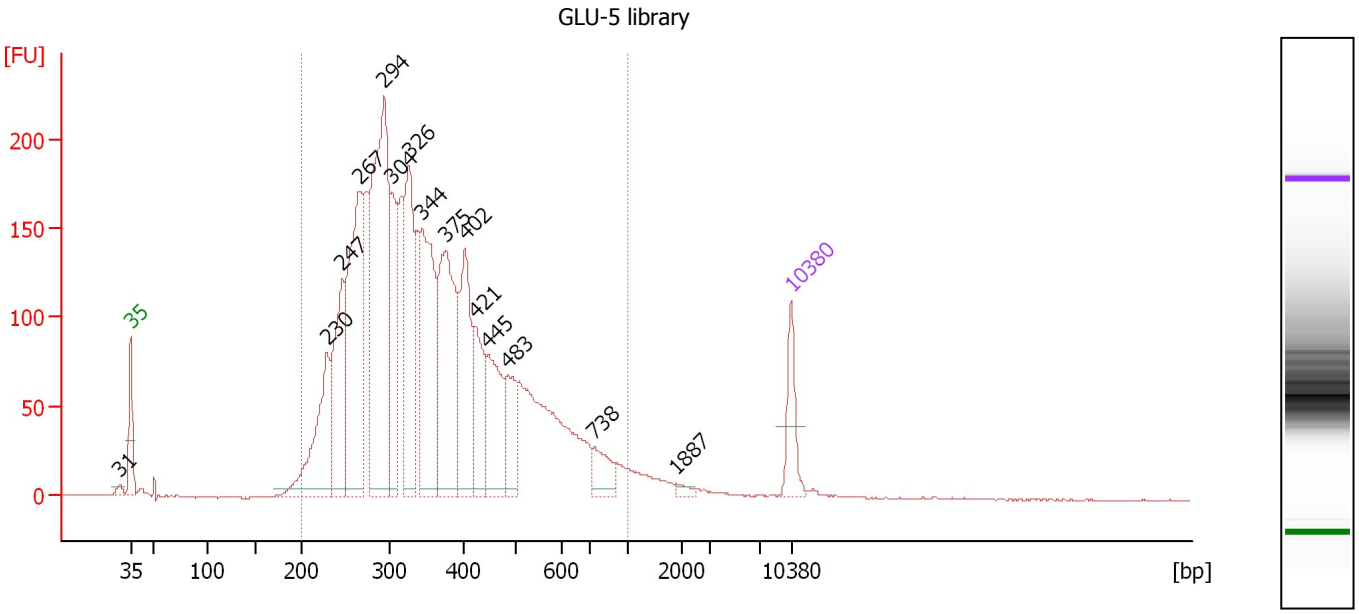
Region table for sample 8 : GLU-3 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	385	4,676.41	4,995.4	20,995.0	96	35.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : GLU-5 library

Number of peaks found:	15	Corr. Area 1:	4,262.3
Noise:	0.4		

Peak table for sample 9 : GLU-5 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.81
2	35	125.00	5,411.3	Lower Marker	43.00
3	230	325.11	2,145.8		63.79
4	247	321.03	1,969.8		65.43
5	267	566.84	3,220.3		67.29
6	294	802.99	4,143.2		69.84
7	304	237.95	1,186.5		70.74
8	326	360.02	1,673.1		72.49
9	344	425.68	1,875.4		73.90
10	375	438.41	1,769.6		76.39
11	402	262.62	990.8		78.42
12	421	176.87	636.7		79.50
13	445	202.61	689.3		80.85
14	483	117.89	369.8		82.94
15	738	65.43	134.4		92.10
16	1,887	10.49	8.4		100.80
17	10,380	75.00	10.9	Upper Marker	113.00

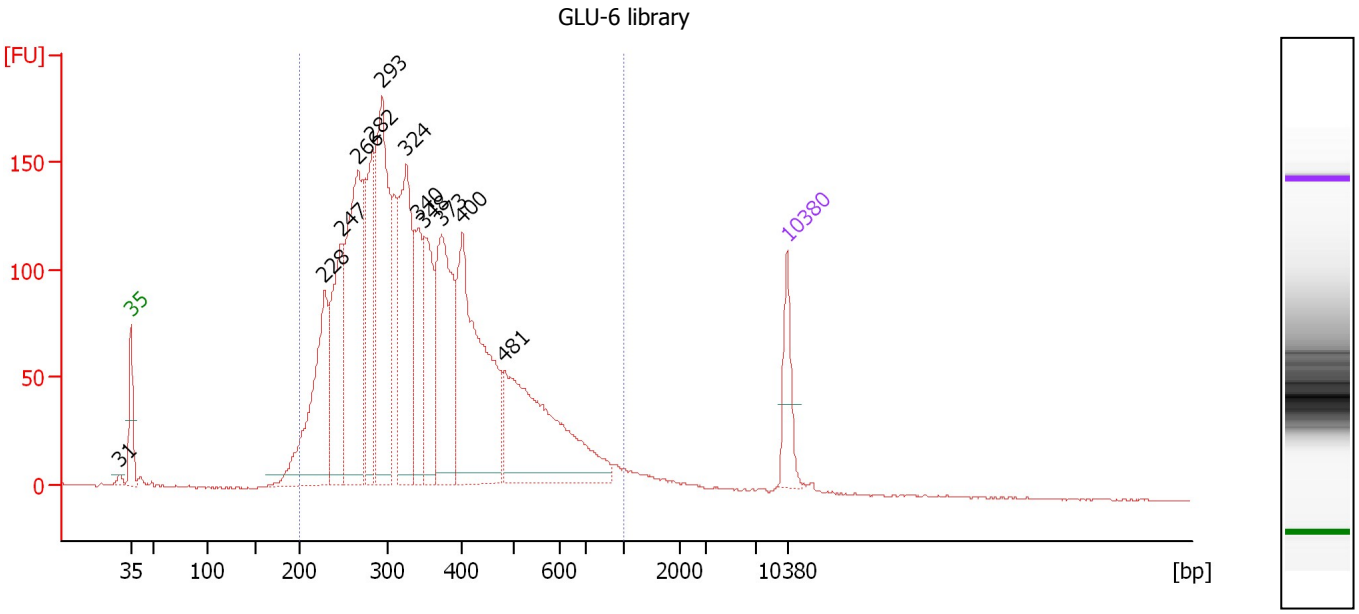
Region table for sample 9 : GLU-5 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	372	5,222.98	4,262.3	24,129.5	96	34.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : GLU-6 library

Number of peaks found:	12	Corr. Area 1:	3,637.0
Noise:	0.4		

Peak table for sample 10 : GLU-6 library

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.75
2	35	125.00	5,411.3	Lower Marker	43.00
3	228	418.02	2,783.3		63.60
4	247	342.50	2,097.0		65.48
5	266	507.82	2,897.3		67.19
6	282	273.38	1,469.6		68.72
7	293	542.30	2,807.1		69.75
8	324	395.92	1,848.9		72.37
9	340	177.10	788.5		73.62
10	348	224.13	975.2		74.24
11	373	368.71	1,498.9		76.18
12	400	529.59	2,005.5		78.34
13	481	466.00	1,467.7		82.84
14	10,380	75.00	10.9	Upper Marker	113.00

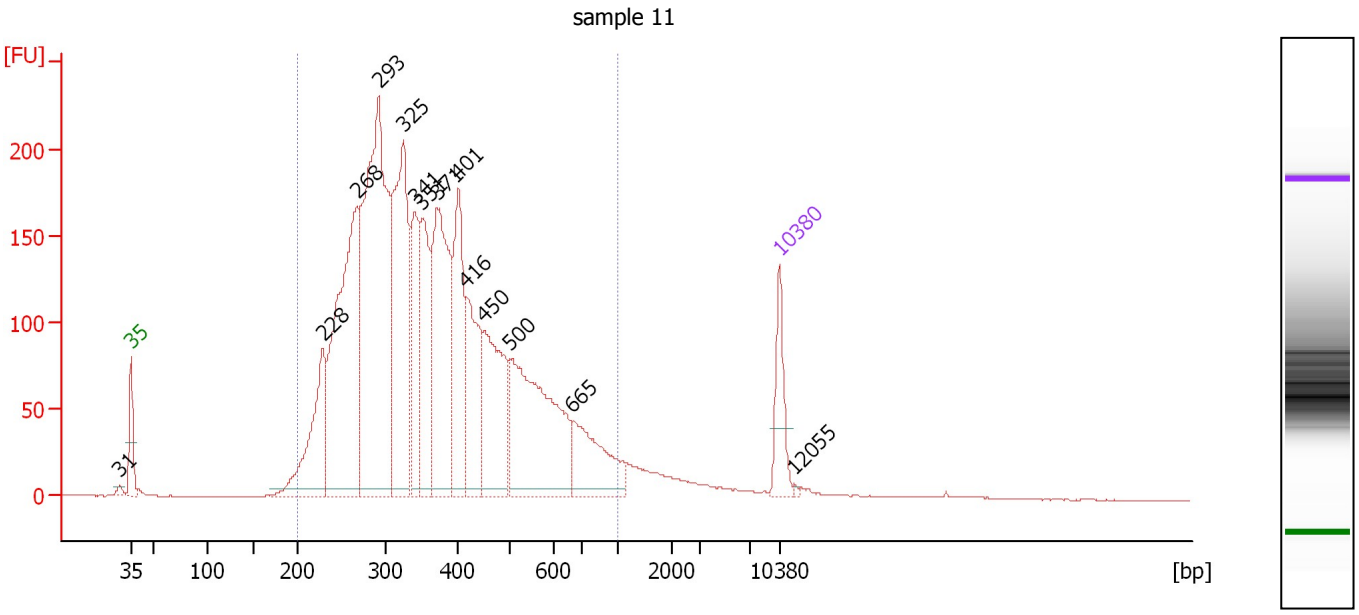
Region table for sample 10 : GLU-6 library

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	362	4,554.26	3,637.0	21,561.6	96	34.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found:	14	Corr. Area 1:	4,678.6
Noise:	0.3		

Peak table for sample 11 : sample 11

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.73
2	35	125.00	5,411.3	Lower Marker	43.00
3	228	279.43	1,853.5		63.68
4	268	686.10	3,886.0		67.37
5	293	941.39	4,875.5		69.73
6	325	495.11	2,310.7		72.38
7	341	205.85	915.4		73.65
8	351	236.71	1,022.1		74.46
9	371	427.89	1,745.9		76.07
10	401	255.01	964.2		78.37
11	416	210.30	765.5		79.24
12	450	265.24	892.1		81.14
13	500	438.88	1,329.1		83.91
14	665	166.30	379.0		90.59
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,055	0.00	0.0		114.61

Region table for sample 11 : sample 11

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	382	4,599.29	4,678.6	20,830.2	95	35.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Gel Image

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad

Created: 7/30/2019 11:26:11 AM
Modified: 7/30/2019 12:07:30 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 12)		Instrument	Run		7/30/2019 12:07:29 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2019-07-30\2019-07-30_001.xad)		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/30/2019 11:26:16 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1