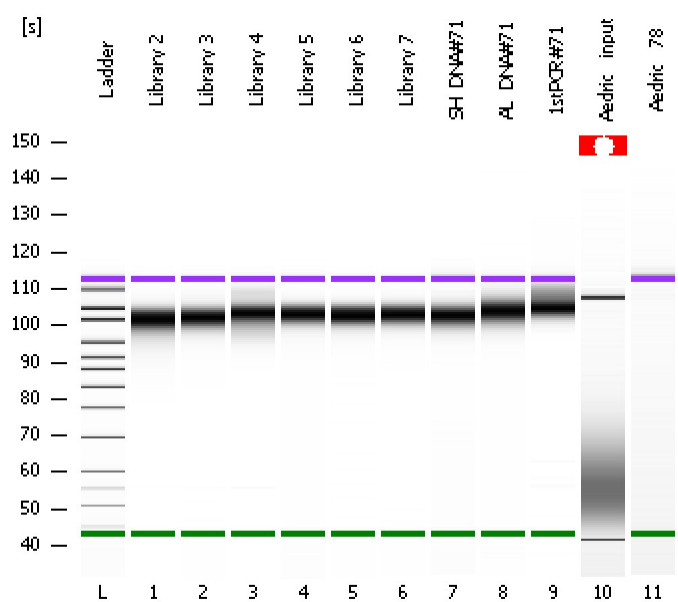


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
Modified: 10/15/2019 3:53:47 PM

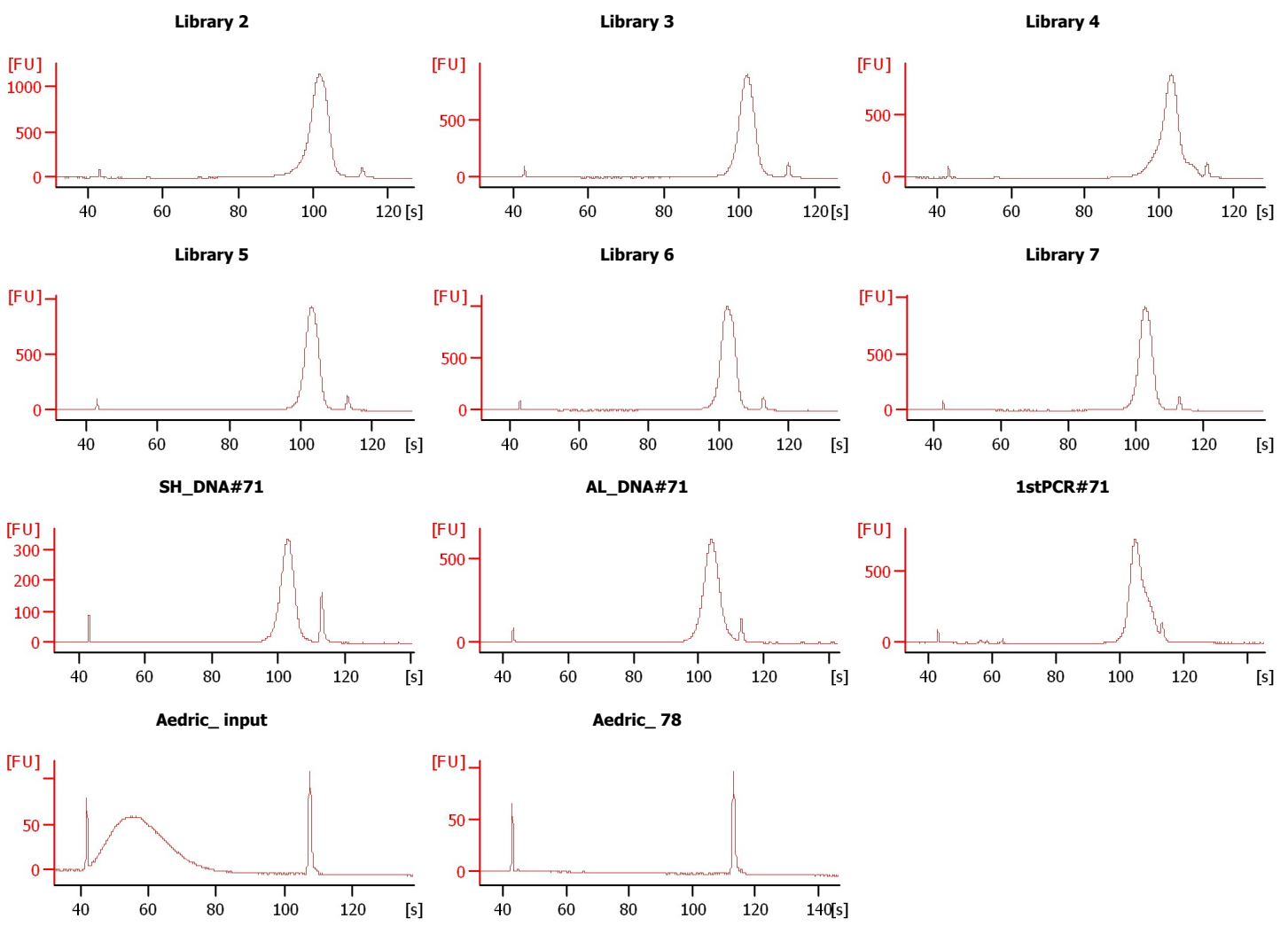
Electrophoresis File Run Summary



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:



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Library 2		<input type="checkbox"/>	✓			
Library 3		<input type="checkbox"/>	✓			
Library 4		<input type="checkbox"/>	✓			
Library 5		<input type="checkbox"/>	✓			
Library 6		<input type="checkbox"/>	✓			
Library 7		<input type="checkbox"/>	✓			
SH_DNA#71		<input type="checkbox"/>	✓			
AL_DNA#71		<input type="checkbox"/>	✓			
1stPCR#71		<input type="checkbox"/>	✓			
Aedric_input		<input type="checkbox"/>	✓			
Aedric_78		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
Modified: 10/15/2019 3:53:47 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

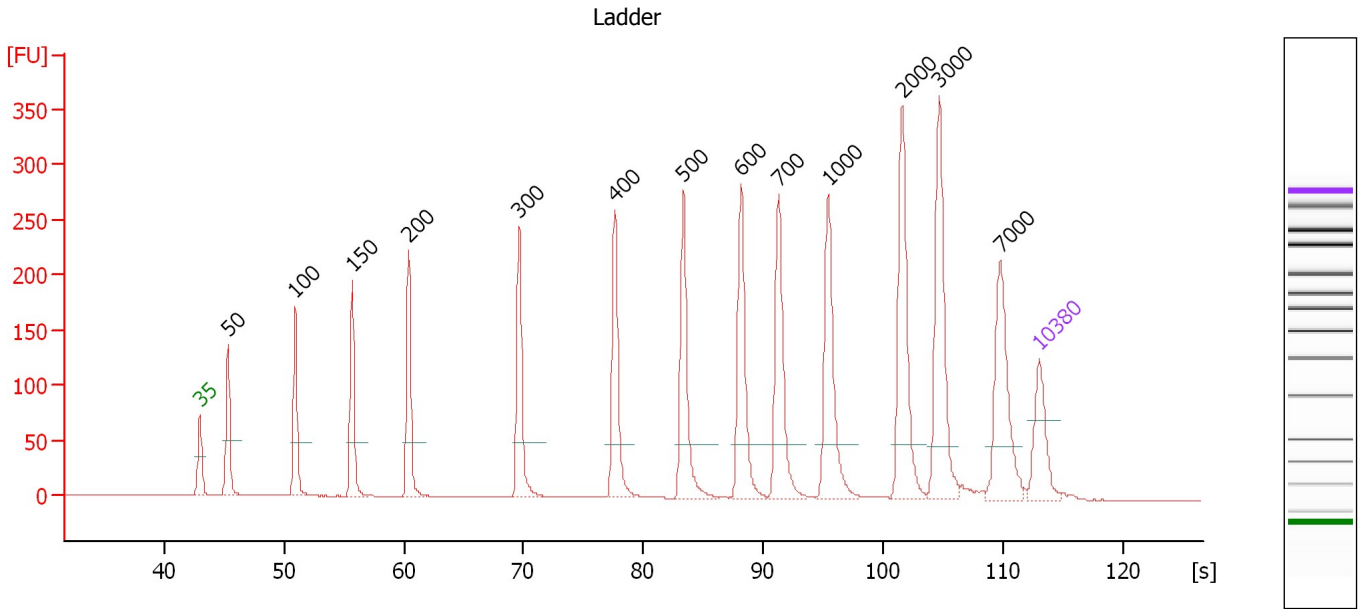
Ladder

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\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

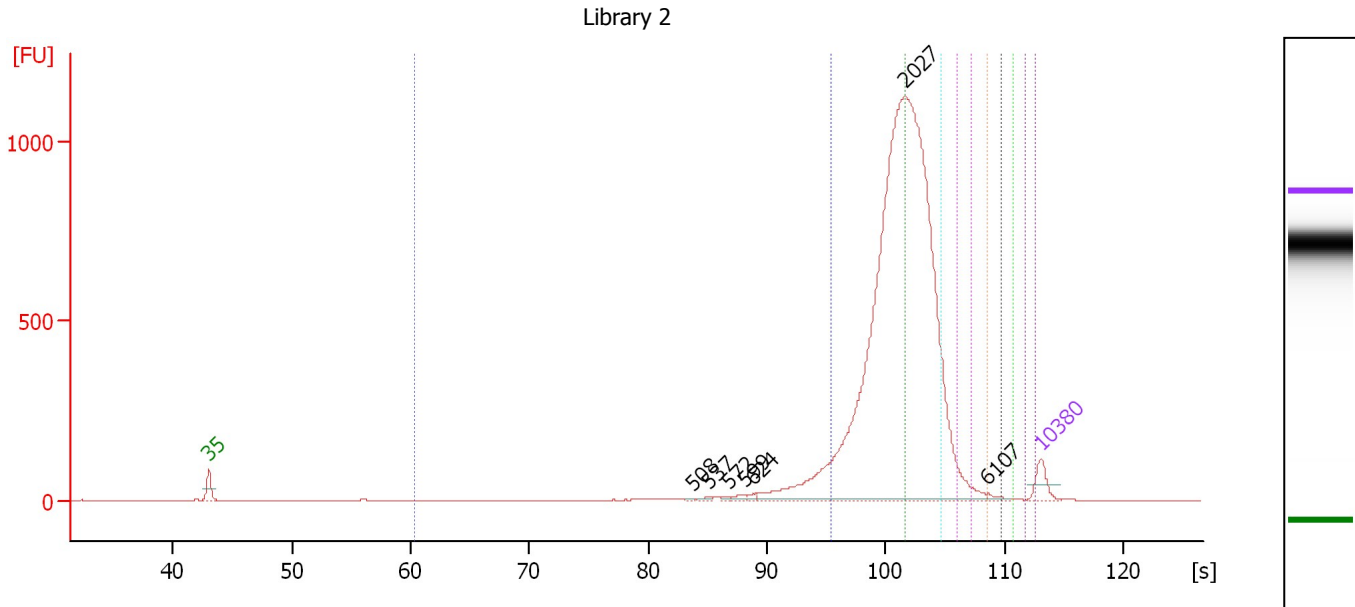
Peak table for Ladder

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	45.37
3	100	150.00	2,272.7	Ladder Peak	50.97
4	150	150.00	1,515.2	Ladder Peak	55.71
5	200	150.00	1,136.4	Ladder Peak	60.41
6	300	150.00	757.6	Ladder Peak	69.63
7	400	150.00	568.2	Ladder Peak	77.60
8	500	150.00	454.5	Ladder Peak	83.33
9	600	150.00	378.8	Ladder Peak	88.12
10	700	150.00	324.7	Ladder Peak	91.25
11	1,000	150.00	227.3	Ladder Peak	95.37
12	2,000	150.00	113.6	Ladder Peak	101.54
13	3,000	150.00	75.8	Ladder Peak	104.63
14	7,000	150.00	32.5	Ladder Peak	109.73
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Library 2

Number of peaks found:	7	Corr. Area 5:	84.8
Noise:	0.2	Corr. Area 6:	16.9
Corr. Area 1:	442.1	Corr. Area 7:	6.2
Corr. Area 2:	3,281.0	Corr. Area 8:	3.5
Corr. Area 3:	2,784.2	Corr. Area 9:	0.6
Corr. Area 4:	303.2		

Peak table for sample 1 : Library 2

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	508	4.56	13.6		83.73
3	537	10.11	28.5		85.08
4	572	8.36	22.1		86.79
5	599	15.05	38.1		88.05
6	624	12.88	31.3		88.86
7	2,027	4,562.54	3,410.4		101.63
8	6,107	13.63	3.4		108.59
9	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 1 : Library 2

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	767	366.90	442.1	795.2	6	22.6
1,000	2,000	1,683	2,299.26	3,281.0	2,069.8	47	14.4
2,000	3,000	2,417	1,883.27	2,784.2	1,180.5	40	11.4
3,000	4,000	3,353	206.01	303.2	93.1	4	8.3
4,000	5,000	4,373	58.07	84.8	20.1	1	6.8
6,000	7,000	6,388	11.76	16.9	2.8	0	4.5
7,000	8,000	7,377	4.30	6.2	0.9	0	4.1
8,000	9,000	8,398	2.38	3.5	0.4	0	3.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...

... Region table for sample 1 :

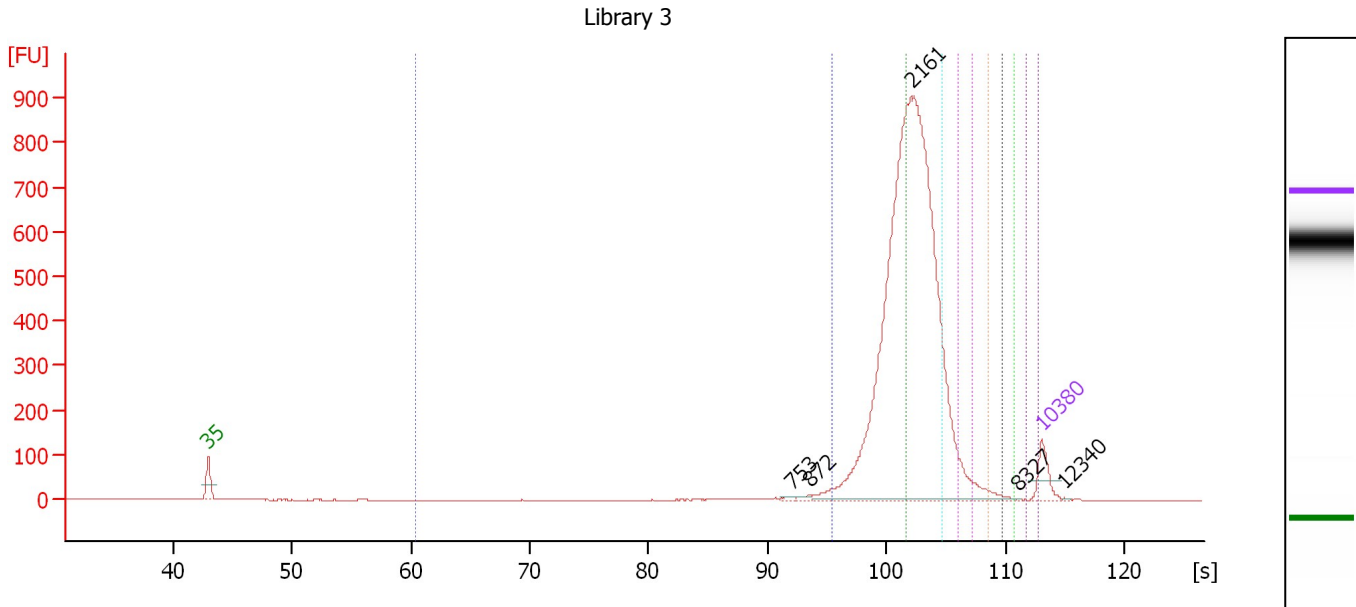
Library 2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
9,000	10,000	9,753	0.39	0.6	0.1	0	2.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
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Electropherogram Summary Continued ...



Overall Results for sample 2 : Library 3

Number of peaks found:	5	Corr. Area 5:	85.0
Noise:	0.4	Corr. Area 6:	21.4
Corr. Area 1:	83.6	Corr. Area 7:	7.7
Corr. Area 2:	1,777.3	Corr. Area 8:	3.9
Corr. Area 3:	2,282.0	Corr. Area 9:	0.5
Corr. Area 4:	292.5		

Peak table for sample 2 : Library 3

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	753	7.34	14.8		91.98
3	872	9.39	16.3		93.61
4	2,161	2,910.11	2,040.2		102.04
5	8,327	1.45	0.3		111.02
6	10,380	75.00	10.9	Upper Marker	113.00
7	12,340	0.00	0.0		114.89

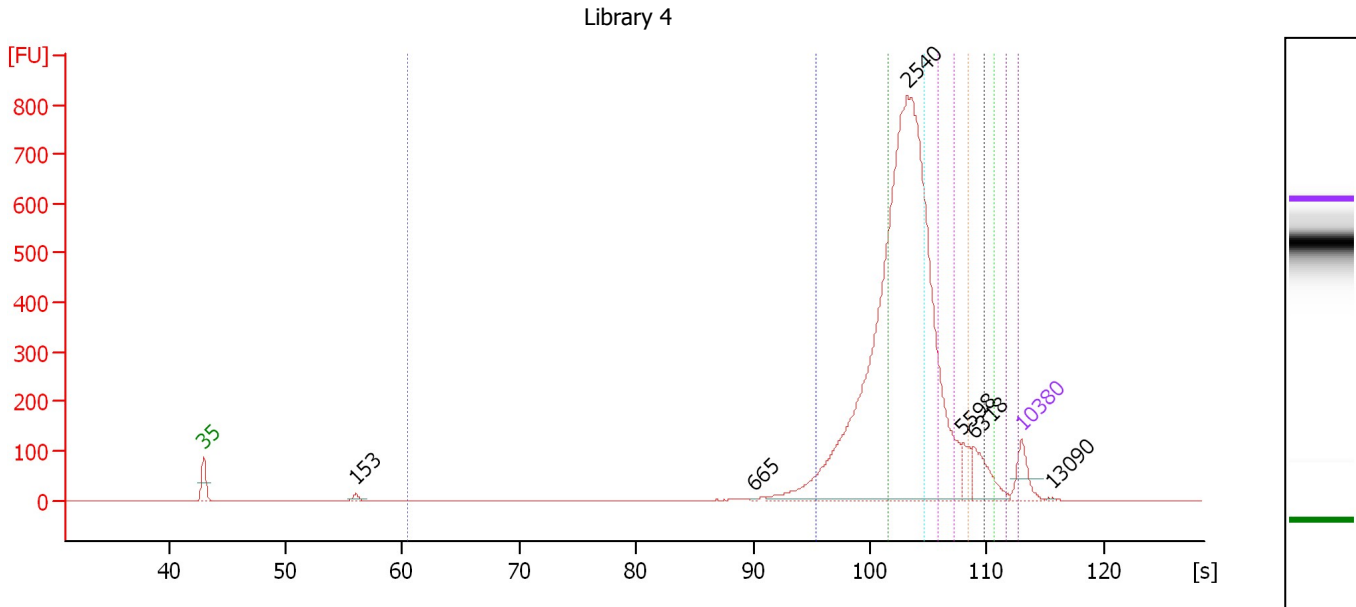
Region table for sample 2 : Library 3

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	782	65.79	83.6	137.1	2	21.0
1,000	2,000	1,748	1,178.41	1,777.3	1,021.2	39	11.9
2,000	3,000	2,425	1,471.75	2,282.0	919.7	50	11.3
3,000	4,000	3,356	189.50	292.5	85.6	6	8.6
4,000	5,000	4,379	55.47	85.0	19.2	2	6.6
6,000	7,000	6,406	14.19	21.4	3.4	0	4.8
7,000	8,000	7,406	5.08	7.7	1.0	0	3.9
8,000	9,000	8,351	2.50	3.9	0.5	0	3.2
9,000	10,000	9,752	0.34	0.5	0.1	0	2.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

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 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Library 4

Number of peaks found:	6	Corr. Area 5:	223.4
Noise:	0.3	Corr. Area 6:	121.7
Corr. Area 1:	113.8	Corr. Area 7:	56.1
Corr. Area 2:	1,288.3	Corr. Area 8:	24.6
Corr. Area 3:	2,214.1	Corr. Area 9:	3.6
Corr. Area 4:	560.4		

Peak table for sample 3 : Library 4

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	153	19.11	189.0		56.01
3	665	3.70	8.4		90.16
4	2,540	2,792.18	1,665.5		103.21
5	5,598	59.31	16.1		107.94
6	6,318	114.93	27.6		108.86
7	10,380	75.00	10.9	Upper Marker	113.00
8	13,090	0.00	0.0		115.62

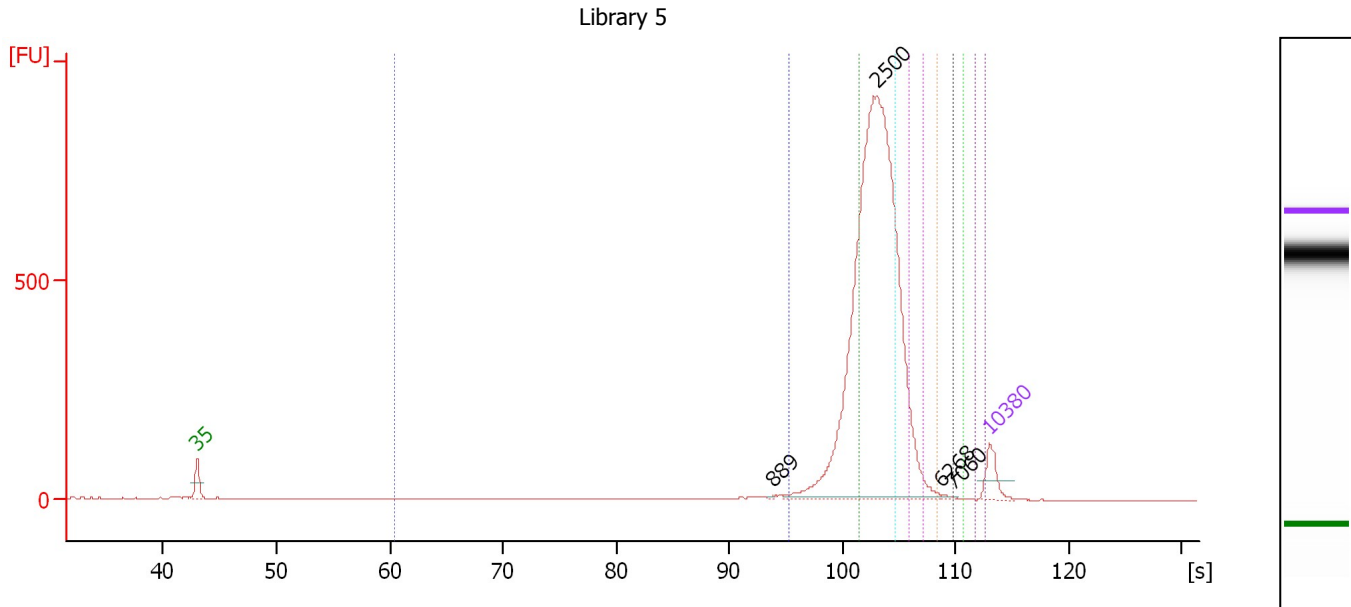
Region table for sample 3 : Library 4

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	854	85.32	113.8	154.6	2	13.8
1,000	2,000	1,667	841.12	1,288.3	764.6	27	15.7
2,000	3,000	2,505	1,392.97	2,214.1	842.5	46	11.1
3,000	4,000	3,399	354.14	560.4	157.9	12	8.6
4,000	5,000	4,396	142.19	223.4	49.0	5	6.6
6,000	7,000	6,440	78.65	121.7	18.5	3	4.6
7,000	8,000	7,397	36.12	56.1	7.4	1	4.0
8,000	9,000	8,369	15.55	24.6	2.8	1	3.5
9,000	10,000	9,619	2.25	3.6	0.4	0	3.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Library 5

Number of peaks found:	4	Corr. Area 5:	150.2
Noise:	0.6	Corr. Area 6:	13.0
Corr. Area 1:	66.7	Corr. Area 7:	4.8
Corr. Area 2:	893.3	Corr. Area 8:	2.4
Corr. Area 3:	2,546.8	Corr. Area 9:	0.3
Corr. Area 4:	546.9		

Peak table for sample 4 : Library 5

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	889	3.01	5.1		93.84
3	2,500	2,560.30	1,551.4		103.09
4	6,268	5.08	1.2		108.80
5	7,060	1.61	0.3		109.79
6	10,380	75.00	10.9	Upper Marker	113.00

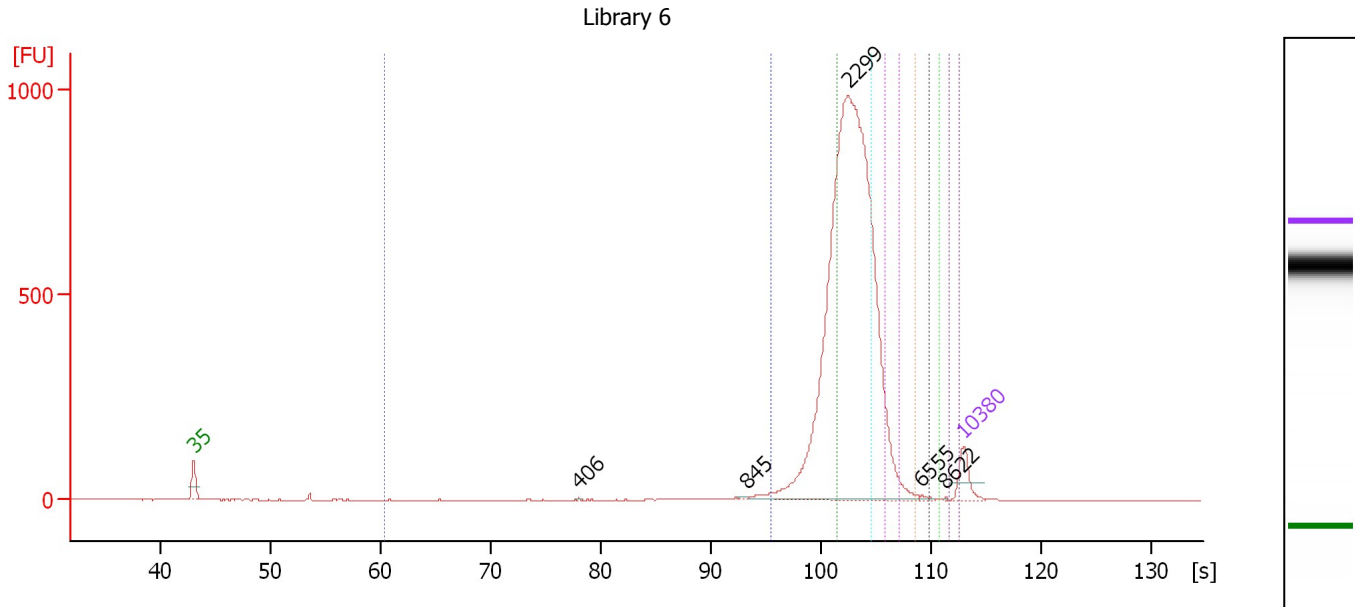
Region table for sample 4 : Library 5

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	708	51.83	66.7	129.5	2	28.1
1,000	2,000	1,785	565.27	893.3	479.8	21	11.2
2,000	3,000	2,488	1,574.62	2,546.8	959.1	59	11.3
3,000	4,000	3,386	339.65	546.9	152.0	13	8.4
4,000	5,000	4,339	93.95	150.2	32.8	4	6.4
6,000	7,000	6,400	8.23	13.0	1.9	0	4.5
7,000	8,000	7,389	3.01	4.8	0.6	0	4.0
8,000	9,000	8,364	1.48	2.4	0.3	0	3.7
9,000	10,000	9,768	0.16	0.3	0.0	0	2.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Library 6

Number of peaks found:	5	Corr. Area 5:	141.7
Noise:	0.4	Corr. Area 6:	14.4
Corr. Area 1:	79.5	Corr. Area 7:	4.8
Corr. Area 2:	1,263.8	Corr. Area 8:	4.6
Corr. Area 3:	2,748.2	Corr. Area 9:	0.0
Corr. Area 4:	541.1		

Peak table for sample 5 : Library 6

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	406	1.62	6.1		77.95
3	845	4.65	8.3		93.24
4	2,299	2,900.51	1,911.5		102.47
5	6,555	5.84	1.4		109.17
6	8,622	1.75	0.3		111.30
7	10,380	75.00	10.9	Upper Marker	113.00

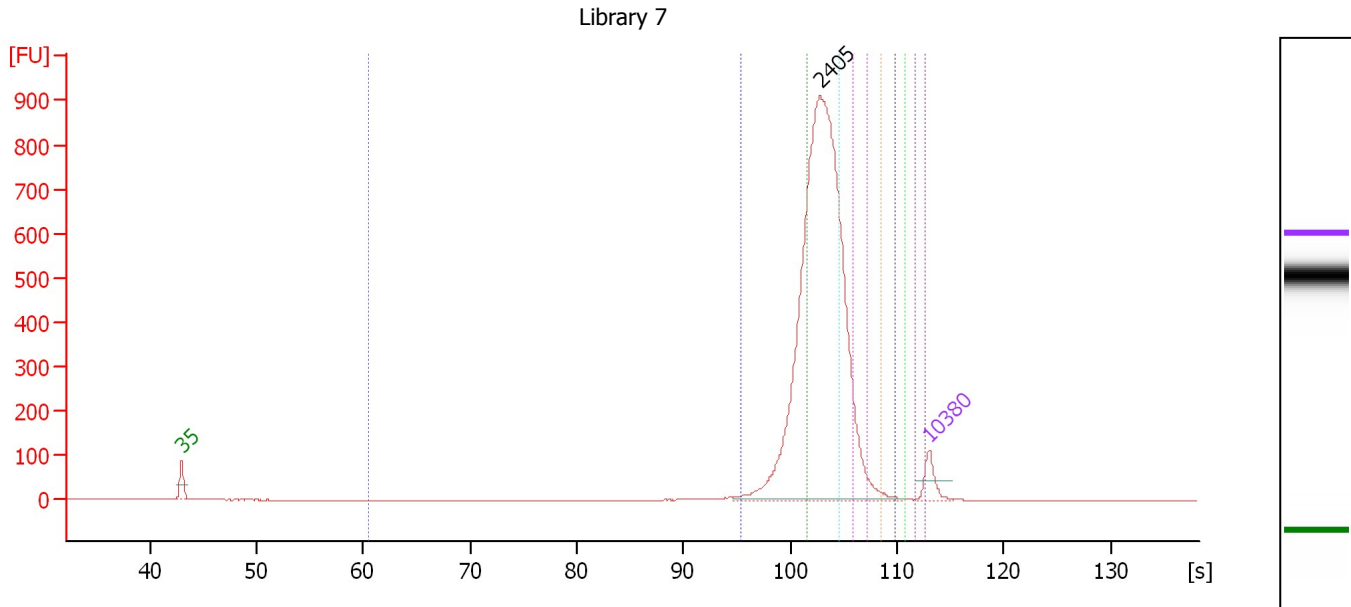
Region table for sample 5 : Library 6

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	733	60.48	79.5	143.2	2	26.1
1,000	2,000	1,784	792.15	1,263.8	672.9	26	11.1
2,000	3,000	2,471	1,682.66	2,748.2	1,031.8	56	11.6
3,000	4,000	3,382	332.77	541.1	149.1	11	8.4
4,000	5,000	4,328	87.77	141.7	30.7	3	6.4
6,000	7,000	6,410	9.03	14.4	2.1	0	4.7
7,000	8,000	7,395	3.02	4.8	0.6	0	3.9
8,000	9,000	8,488	2.83	4.6	0.5	0	3.0
9,000	10,000	9,744	0.00	0.0	0.0	0	1.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Library 7

Number of peaks found:	1	Corr. Area 5:	163.5
Noise:	0.2	Corr. Area 6:	15.1
Corr. Area 1:	29.7	Corr. Area 7:	5.2
Corr. Area 2:	918.8	Corr. Area 8:	2.5
Corr. Area 3:	2,486.4	Corr. Area 9:	0.0
Corr. Area 4:	532.9		

Peak table for sample 6 : Library 7

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	2,405	2,939.81	1,852.1		102.79
3	10,380	75.00	10.9	Upper Marker	113.00

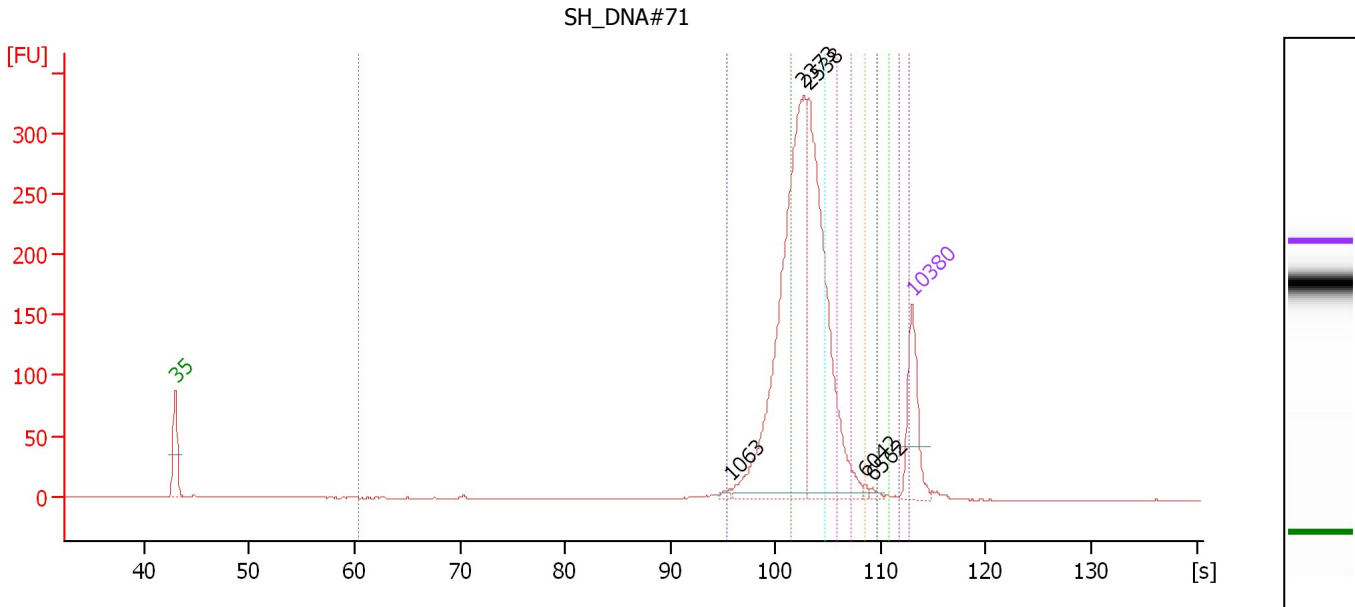
Region table for sample 6 : Library 7

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	827	25.17	29.7	47.6	1	16.0
1,000	2,000	1,789	663.83	918.8	562.3	22	10.8
2,000	3,000	2,484	1,755.82	2,486.4	1,071.0	59	11.3
3,000	4,000	3,383	377.94	532.9	169.3	13	8.4
4,000	5,000	4,333	116.77	163.5	40.8	4	6.7
6,000	7,000	6,375	10.97	15.1	2.6	0	4.7
7,000	8,000	7,389	3.74	5.2	0.8	0	4.0
8,000	9,000	8,372	1.76	2.5	0.3	0	3.5
9,000	10,000	9,767	0.00	0.0	0.0	0	2.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

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 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : SH_DNA#71

Number of peaks found:	5	Corr. Area 5:	58.0
Noise:	0.2	Corr. Area 6:	10.7
Corr. Area 1:	13.0	Corr. Area 7:	3.7
Corr. Area 2:	473.6	Corr. Area 8:	2.4
Corr. Area 3:	891.8	Corr. Area 9:	0.1
Corr. Area 4:	166.3		

Peak table for sample 7 : SH_DNA#71

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	1,063	4.35	6.2		95.76
3	2,373	454.61	290.3		102.69
4	2,538	334.49	199.7		103.20
5	6,042	3.28	0.8		108.51
6	6,562	4.15	1.0		109.17
7	10,380	75.00	10.9	Upper Marker	113.00

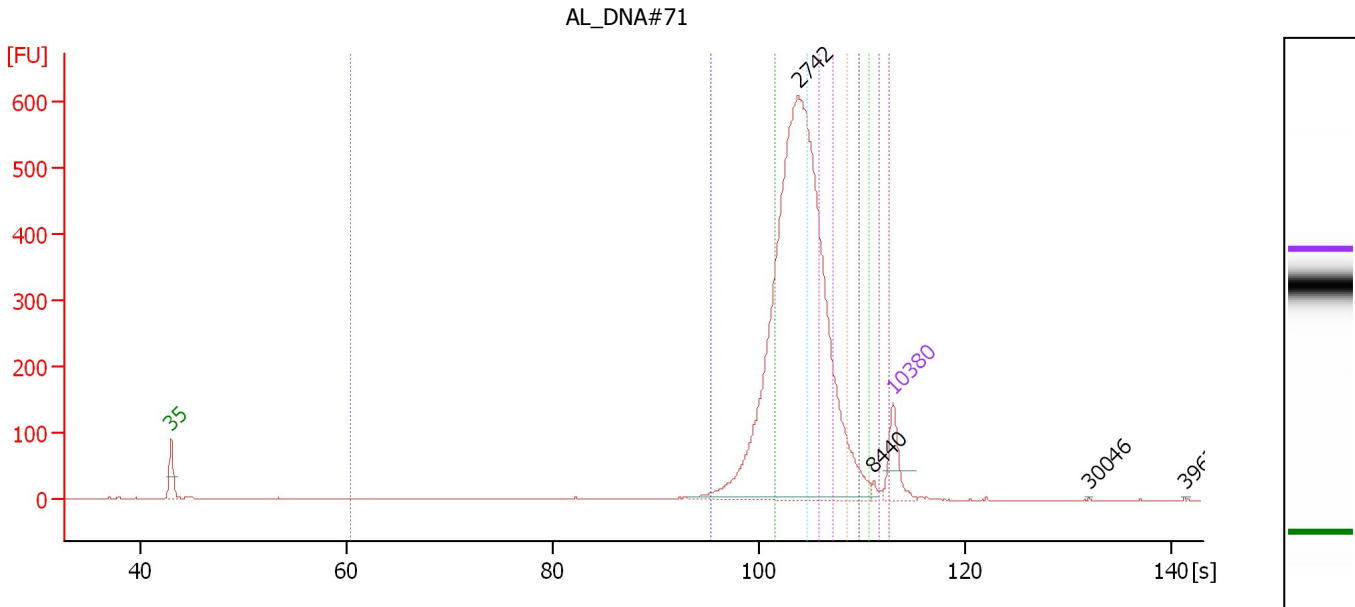
Region table for sample 7 : SH_DNA#71

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	841	7.80	13.0	16.3	1	20.5
1,000	2,000	1,750	240.98	473.6	208.7	29	12.0
2,000	3,000	2,460	441.47	891.8	271.9	54	11.4
3,000	4,000	3,378	82.69	166.3	37.1	10	8.5
4,000	5,000	4,363	29.06	58.0	10.1	4	6.7
6,000	7,000	6,406	5.46	10.7	1.3	1	4.6
7,000	8,000	7,388	1.86	3.7	0.4	0	4.2
8,000	9,000	8,409	1.18	2.4	0.2	0	3.7
9,000	10,000	9,764	0.04	0.1	0.0	0	2.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : AL_DNA#71

Number of peaks found:	4	Corr. Area 5:	359.0
Noise:	0.2	Corr. Area 6:	80.5
Corr. Area 1:	41.6	Corr. Area 7:	33.5
Corr. Area 2:	571.2	Corr. Area 8:	19.4
Corr. Area 3:	1,563.8	Corr. Area 9:	4.0
Corr. Area 4:	623.6		

Peak table for sample 8 : AL_DNA#71

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	2,742	1,863.76	1,029.9		103.83
3	8,440	7.63	1.4		111.13
4	10,380	75.00	10.9	Upper Marker	113.00
5	30,046	0.00	0.0		132.01
6	39,637	0.00	0.0		141.28

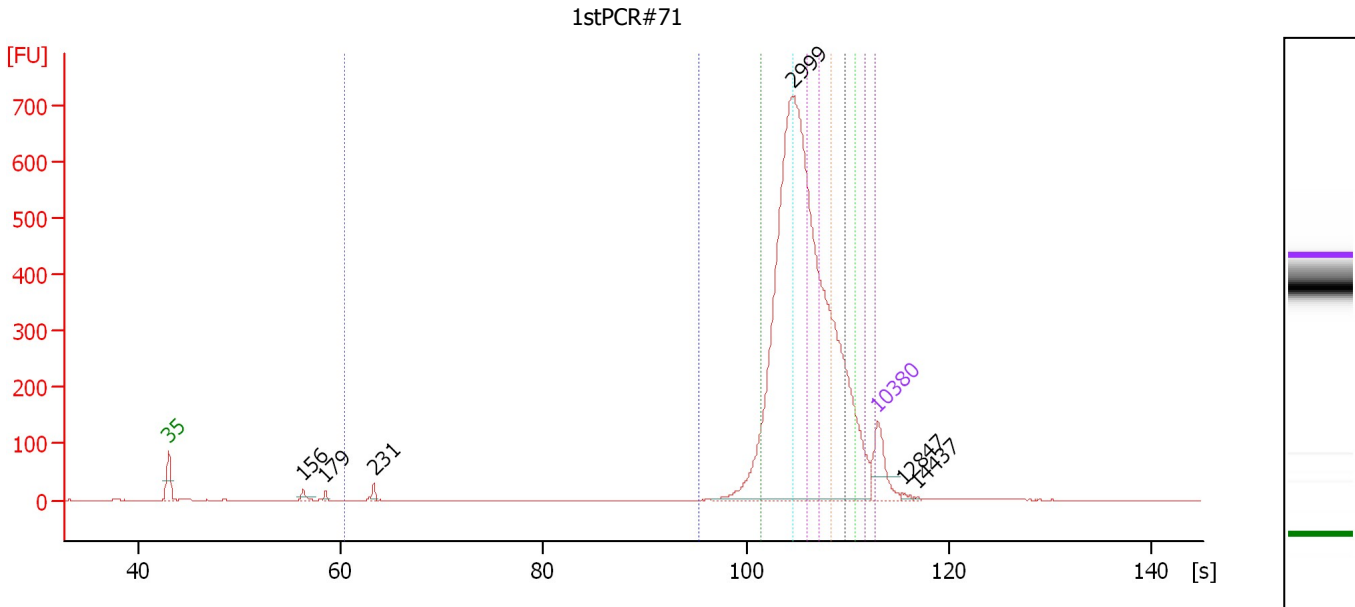
Region table for sample 8 : AL_DNA#71

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	724	28.54	41.6	72.3	1	29.0
1,000	2,000	1,745	320.98	571.2	278.6	16	12.7
2,000	3,000	2,524	854.67	1,563.8	513.1	44	11.1
3,000	4,000	3,429	342.40	623.6	151.3	18	8.8
4,000	5,000	4,399	198.51	359.0	68.4	10	6.6
6,000	7,000	6,398	45.17	80.5	10.7	2	4.6
7,000	8,000	7,384	18.72	33.5	3.8	1	4.1
8,000	9,000	8,390	10.62	19.4	1.9	1	3.3
9,000	10,000	9,617	2.16	4.0	0.3	0	3.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 1stPCR#71

Number of peaks found:	6	Corr. Area 5:	578.8
Noise:	0.2	Corr. Area 6:	334.2
Corr. Area 1:	18.6	Corr. Area 7:	187.8
Corr. Area 2:	147.3	Corr. Area 8:	108.4
Corr. Area 3:	1,289.3	Corr. Area 9:	39.7
Corr. Area 4:	806.9		

Peak table for sample 9 : 1stPCR#71

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	156	17.21	166.8		56.30
3	179	7.93	66.9		58.47
4	231	13.09	86.0		63.24
5	2,999	1,833.21	926.2		104.63
6	10,380	75.00	10.9	Upper Marker	113.00
7	12,847	0.00	0.0		115.38
8	14,437	0.00	0.0		116.92

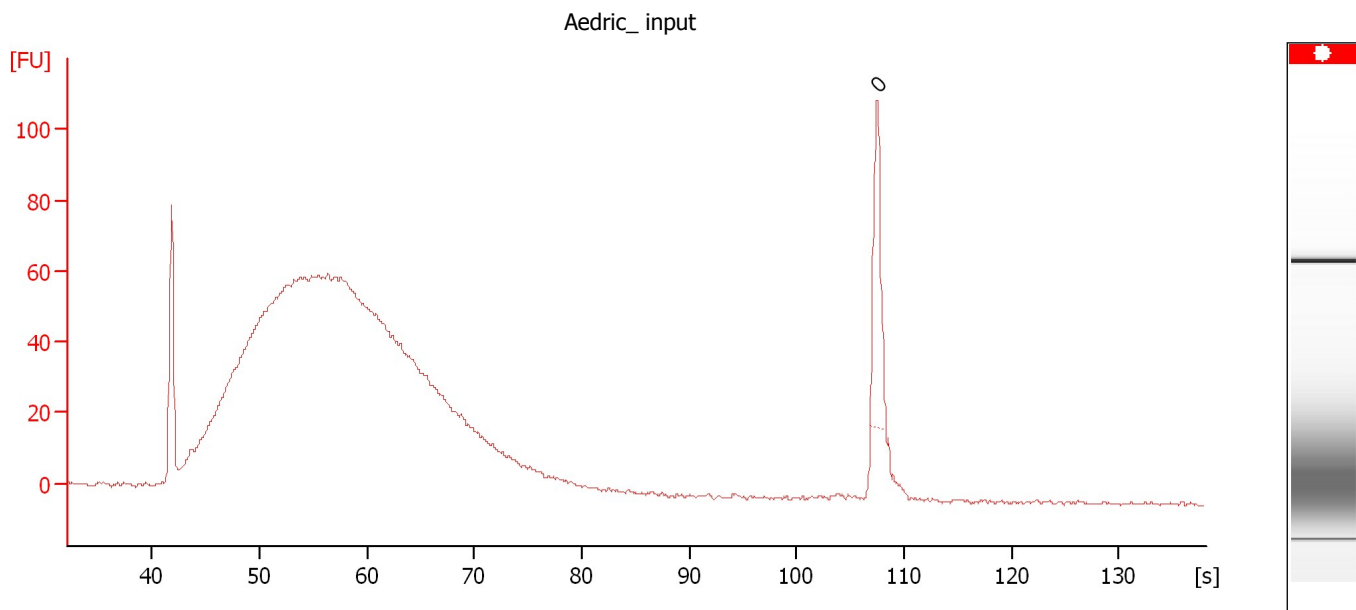
Region table for sample 9 : 1stPCR#71

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	235	14.07	18.6	92.5	0	26.4
1,000	2,000	1,819	69.09	147.3	57.6	4	9.3
2,000	3,000	2,615	593.40	1,289.3	343.9	32	10.0
3,000	4,000	3,439	373.03	806.9	164.3	20	8.6
4,000	5,000	4,418	269.47	578.8	92.4	14	6.8
6,000	7,000	6,421	157.91	334.2	37.3	8	4.6
7,000	8,000	7,411	88.36	187.8	18.1	5	4.2
8,000	9,000	8,411	50.00	108.4	9.0	3	3.5
9,000	10,000	9,433	17.95	39.7	2.9	1	3.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Aedric_input

Number of peaks found: 0 Noise: 0.6

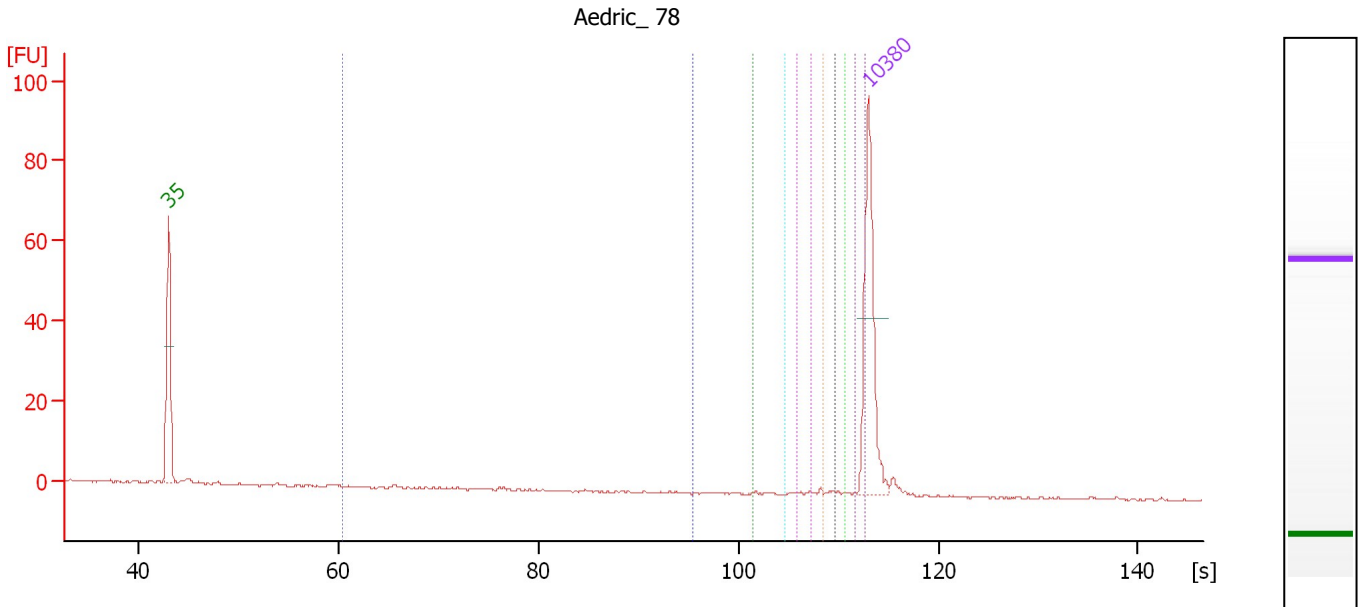
Peak table for sample 10 : Aedric_input

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		107.50

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Aedric_78

Number of peaks found:	0	Corr. Area 5:	0.2
Noise:	0.2	Corr. Area 6:	0.5
Corr. Area 1:	0.2	Corr. Area 7:	0.3
Corr. Area 2:	0.0	Corr. Area 8:	0.2
Corr. Area 3:	0.1	Corr. Area 9:	0.1
Corr. Area 4:	0.1		

Peak table for sample 11 : Aedric_78

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	10,380	75.00	10.9	Upper Marker	113.00

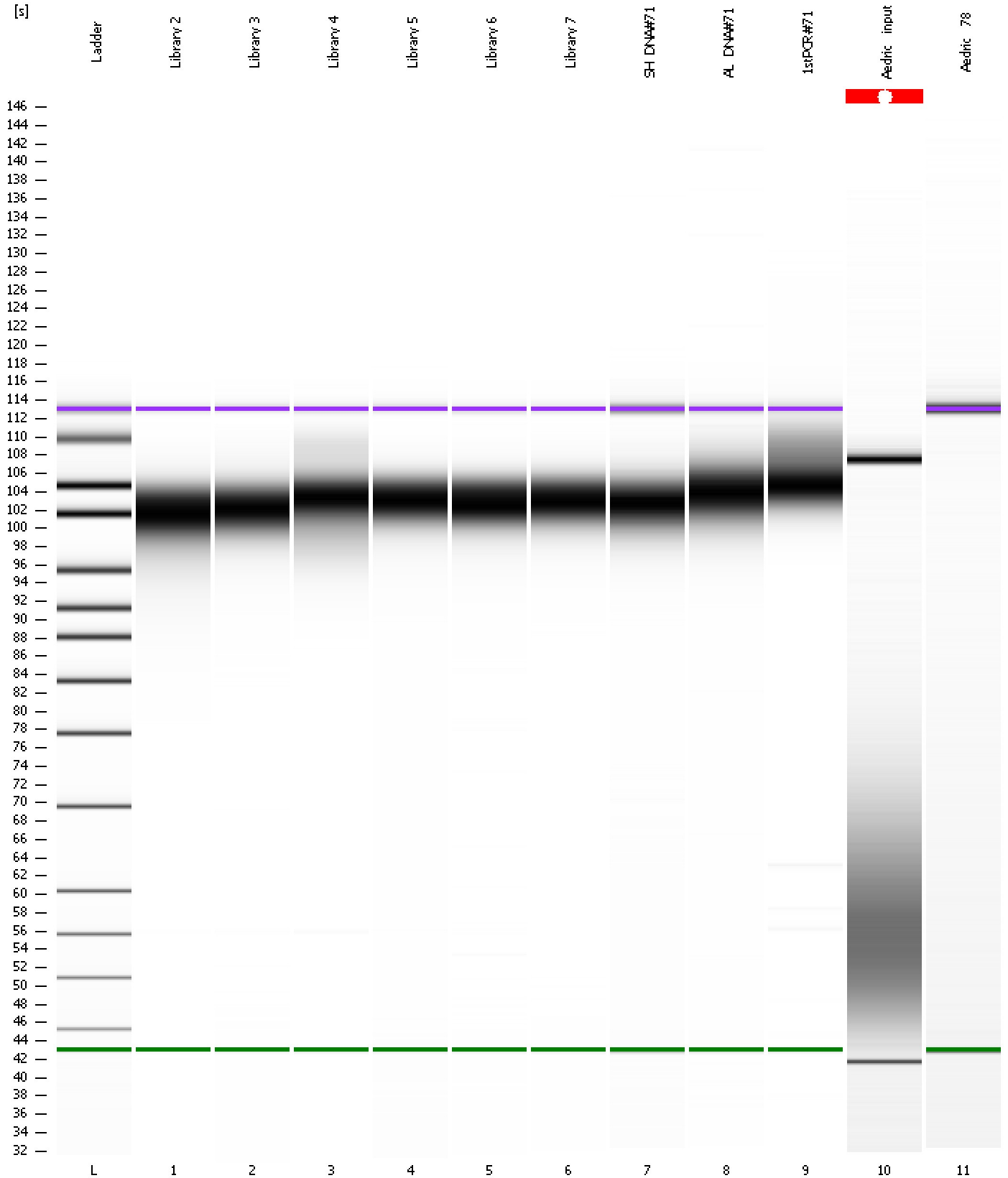
Region table for sample 11 : Aedric_78

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	323	0.24	0.2	1.2	2	28.2
1,000	2,000	1,742	0.01	0.0	0.0	0	1.5
2,000	3,000	2,063	0.06	0.1	0.0	1	1.9
3,000	4,000	3,686	0.09	0.1	0.0	1	4.7
4,000	5,000	4,526	0.18	0.2	0.1	2	7.6
6,000	7,000	6,552	0.46	0.5	0.1	4	4.5
7,000	8,000	7,336	0.27	0.3	0.1	3	4.0
8,000	9,000	8,419	0.21	0.2	0.0	2	3.6
9,000	10,000	9,787	0.06	0.1	0.0	1	1.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
Modified: 10/15/2019 3:53:47 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_002.xad

Created: 10/15/2019 2:32:44 PM
 Modified: 10/15/2019 3:53:47 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		10/15/2019 3:14:01 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\2019-10-15\2019-10-15_002.xad)		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/15/2019 2:32:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1