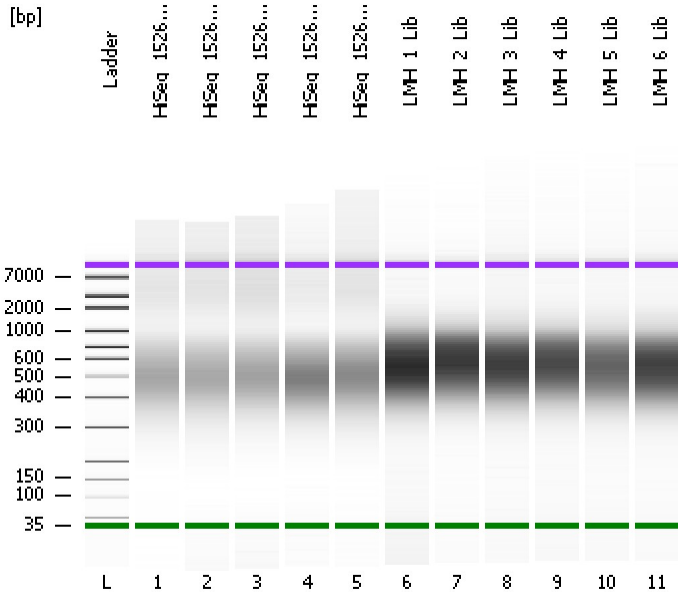


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
Modified: 10/24/2018 4:21:13 PM

Electrophoresis File Run Summary



Instrument Information:

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

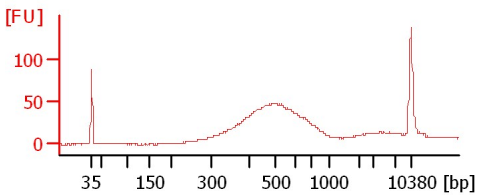
Assay Information:

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

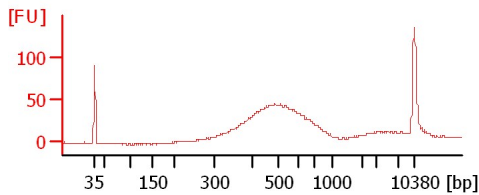
Chip Information:

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

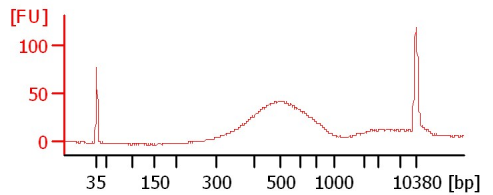
HiSeq_1526_12



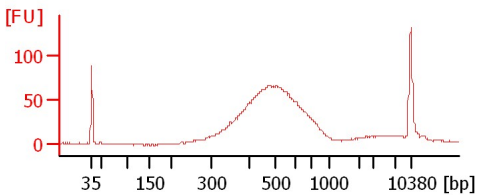
HiSeq_1526_13



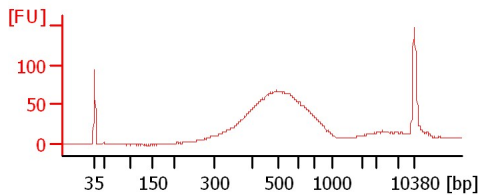
HiSeq_1526_14



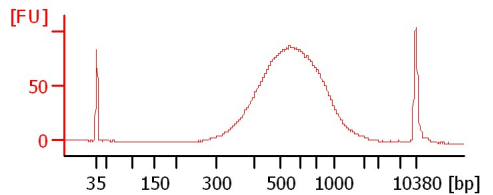
HiSeq_1526_15



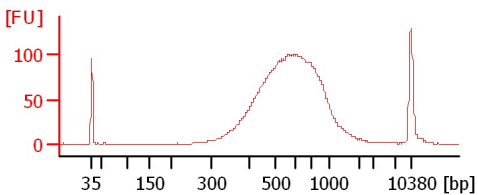
HiSeq_1526_16



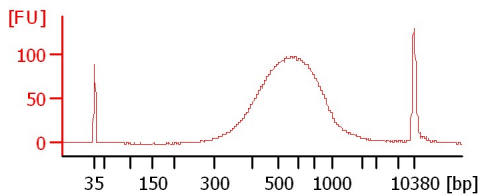
LMH_1_Lib



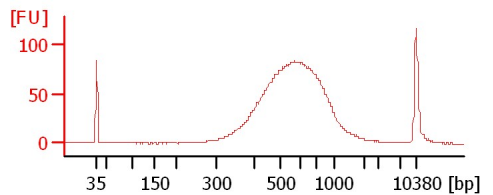
LMH_2_Lib



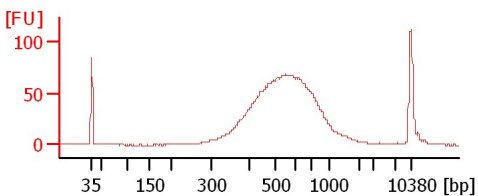
LMH_3_Lib



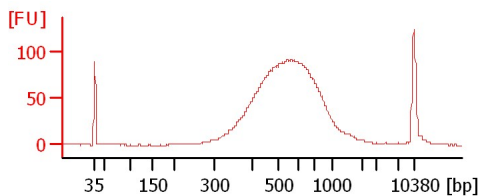
LMH_4_Lib



LMH_5_Lib



LMH_6_Lib



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
Modified: 10/24/2018 4:21:13 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
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HiSeq_1526_13		<input type="checkbox"/>	✓			
HiSeq_1526_14		<input type="checkbox"/>	✓			
HiSeq_1526_15		<input type="checkbox"/>	✓			
HiSeq_1526_16		<input type="checkbox"/>	✓			
LMH_1_Lib		<input type="checkbox"/>	✓			
LMH_2_Lib		<input type="checkbox"/>	✓			
LMH_3_Lib		<input type="checkbox"/>	✓			
LMH_4_Lib		<input type="checkbox"/>	✓			
LMH_5_Lib		<input type="checkbox"/>	✓			
LMH_6_Lib		<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\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
Modified: 10/24/2018 4:21:13 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

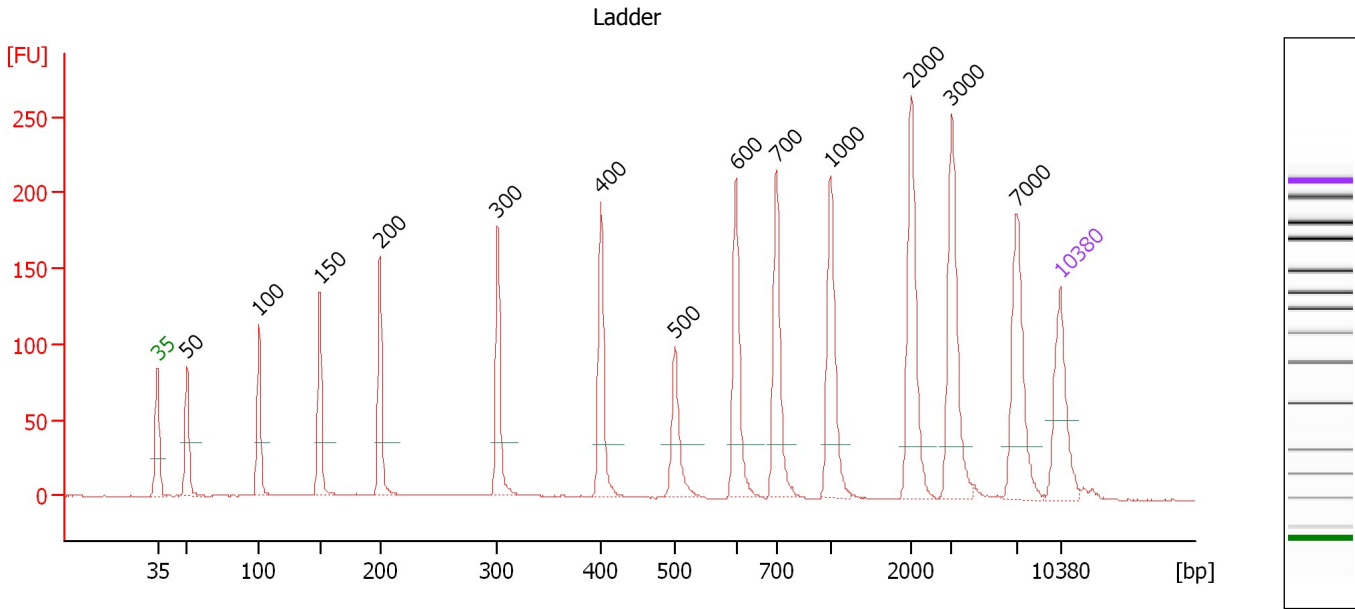
Ladder

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\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

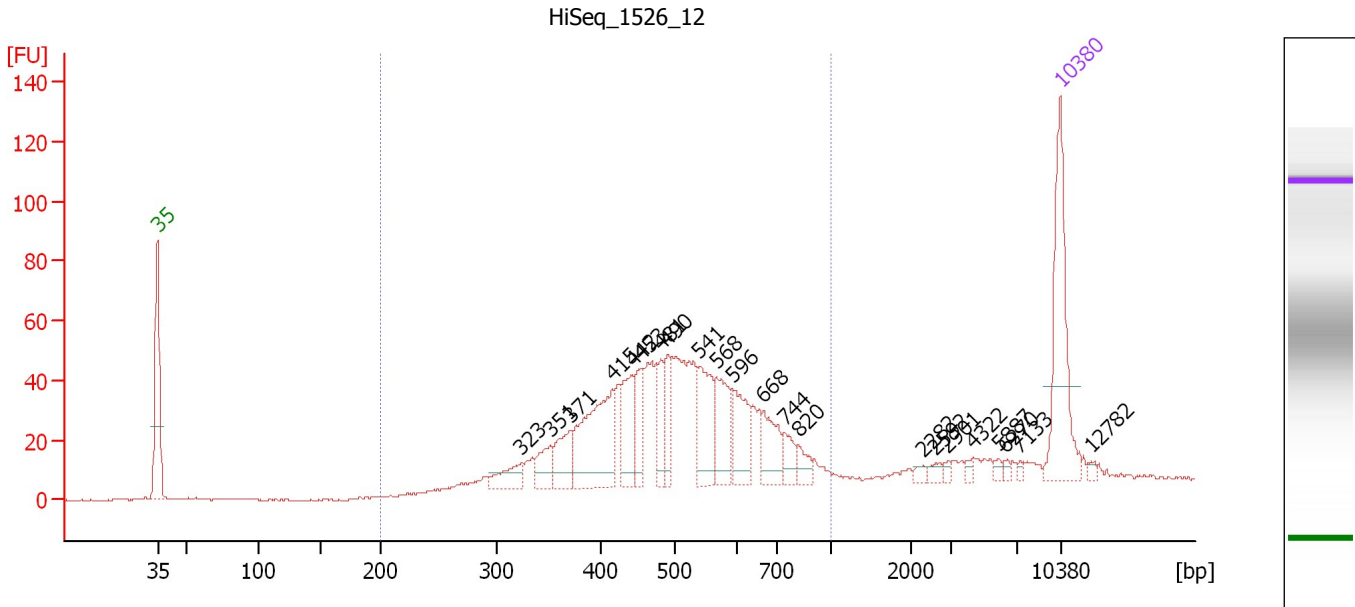
Peak table for Ladder

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	45.29
3	100	150.00	2,272.7	Ladder Peak	50.84
4	150	150.00	1,515.2	Ladder Peak	55.59
5	200	150.00	1,136.4	Ladder Peak	60.26
6	300	150.00	757.6	Ladder Peak	69.37
7	400	150.00	568.2	Ladder Peak	77.38
8	500	150.00	454.5	Ladder Peak	83.11
9	600	150.00	378.8	Ladder Peak	87.86
10	700	150.00	324.7	Ladder Peak	90.99
11	1,000	150.00	227.3	Ladder Peak	95.17
12	2,000	150.00	113.6	Ladder Peak	101.47
13	3,000	150.00	75.8	Ladder Peak	104.59
14	7,000	150.00	32.5	Ladder Peak	109.65
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : HiSeq_1526_12

Number of peaks found: 22 Corr. Area 1: 863.5
 Noise: 0.3

Peak table for sample 1 : HiSeq_1526_12


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	323	23.40	109.7		71.22
3	351	22.04	95.3		73.42
4	371	32.28	131.9		75.05
5	415	97.47	356.2		78.22
6	442	41.70	142.8		79.81
7	453	27.04	90.4		80.42
8	481	26.45	83.4		82.01
9	490	27.07	83.7		82.53
10	541	49.64	139.1		85.04
11	568	42.95	114.7		86.32
12	596	37.81	96.1		87.69
13	668	31.47	71.4		89.97
14	744	15.79	32.1		91.60
15	820	11.69	21.6		92.66
16	2,282	3.51	2.3		102.35
17	2,592	4.60	2.7		103.31
18	2,901	2.49	1.3		104.28
19	4,322	3.48	1.2		106.26
20	5,887	3.13	0.8		108.25
21	6,270	2.64	0.6		108.73
22	7,133	2.27	0.5		109.79
23	10,380	75.00	10.9	Upper Marker	113.00
24	12,782	0.00	0.0		115.38

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...

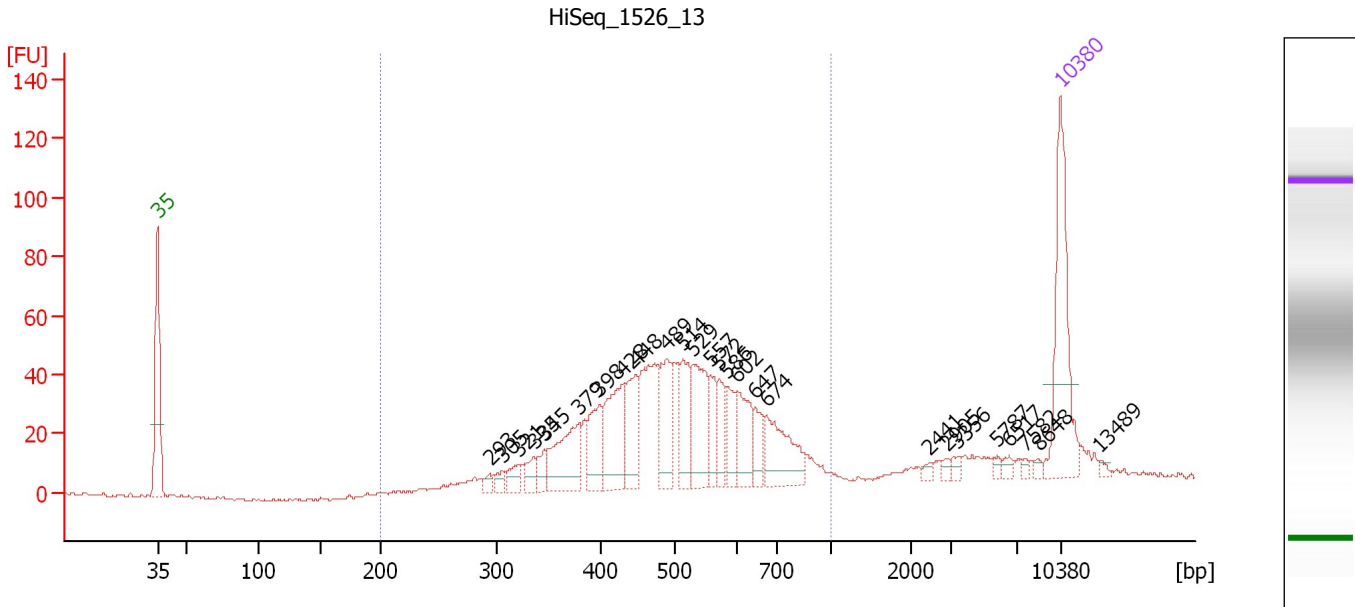
... Region table for sample 1 : HiSeq 1526 12

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	508	741.10	863.5	2,412.8	 89	25.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : HiSeq_1526_13

Number of peaks found: 26 Corr. Area 1: 788.7
 Noise: 0.6

Peak table for sample 2 : HiSeq_1526_13

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	293	6.82	35.2		68.77
3	305	8.18	40.6		69.78
4	321	14.81	69.9		71.05
5	335	14.03	63.4		72.20
6	345	12.75	56.0		72.99
7	379	60.36	241.2		75.71
8	398	37.59	143.0		77.25
9	428	63.86	225.8		79.01
10	448	40.66	137.6		80.11
11	489	53.04	164.3		82.48
12	514	39.14	115.4		83.76
13	529	54.66	156.4		84.51
14	557	24.08	65.5		85.83
15	572	22.50	59.6		86.53
16	586	24.66	63.8		87.19
17	602	34.97	87.9		87.94
18	647	18.59	43.5		89.34
19	674	50.47	113.4		90.18
20	2,441	2.91	1.8		102.84
21	2,905	3.23	1.7		104.29
22	3,356	4.11	1.9		105.04
23	5,787	2.82	0.7		108.12
24	6,517	4.03	0.9		109.04
25	7,582	2.26	0.5		110.23

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...

... Peak table for sample 2 : HiSeq_1526_13

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	8,648	2.65	0.5		111.29
27	10,380	75.00	10.9	Upper Marker	113.00
28	13,489	0.00	0.0		116.08

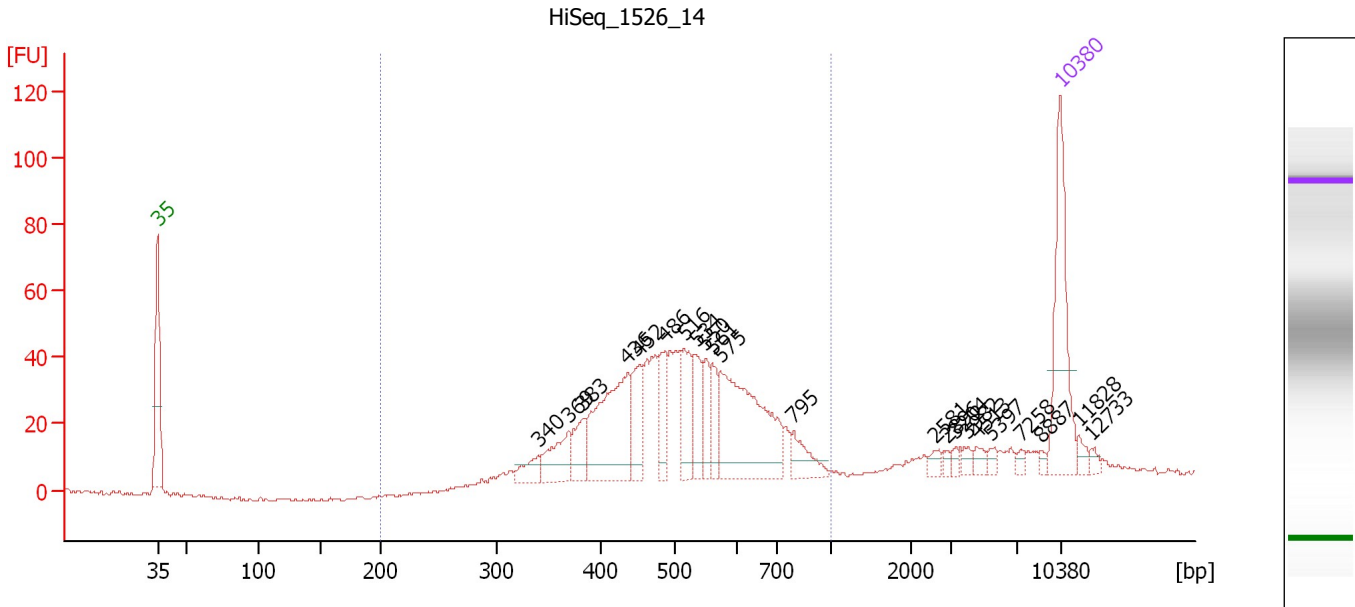
Region table for sample 2 : HiSeq_1526_13

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	512	671.44	788.7	2,151.0	88	24.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : HiSeq_1526_14

Number of peaks found: 22 Corr. Area 1: 697.0
 Noise: 0.5

Peak table for sample 3 : HiSeq_1526_14

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	340	18.18	81.1		72.56
3	369	38.16	156.8		74.89
4	383	27.16	107.5		76.01
5	436	118.80	412.6		79.46
6	452	39.54	132.6		80.35
7	486	34.75	108.3		82.32
8	516	40.04	117.7		83.84
9	534	33.41	94.7		84.74
10	550	25.80	71.0		85.50
11	561	24.18	65.3		85.99
12	575	135.34	356.7		86.67
13	795	22.16	42.3		92.31
14	2,581	5.89	3.5		103.28
15	2,896	3.45	1.8		104.27
16	3,204	3.73	1.8		104.85
17	3,982	5.92	2.3		105.83
18	4,513	5.99	2.0		106.51
19	5,397	4.78	1.3		107.63
20	7,258	3.68	0.8		109.91
21	8,887	2.93	0.5		111.52
22	10,380	75.00	10.9	Upper Marker	113.00
23	11,828	0.00	0.0		114.43
24	12,733	0.00	0.0		115.33

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...

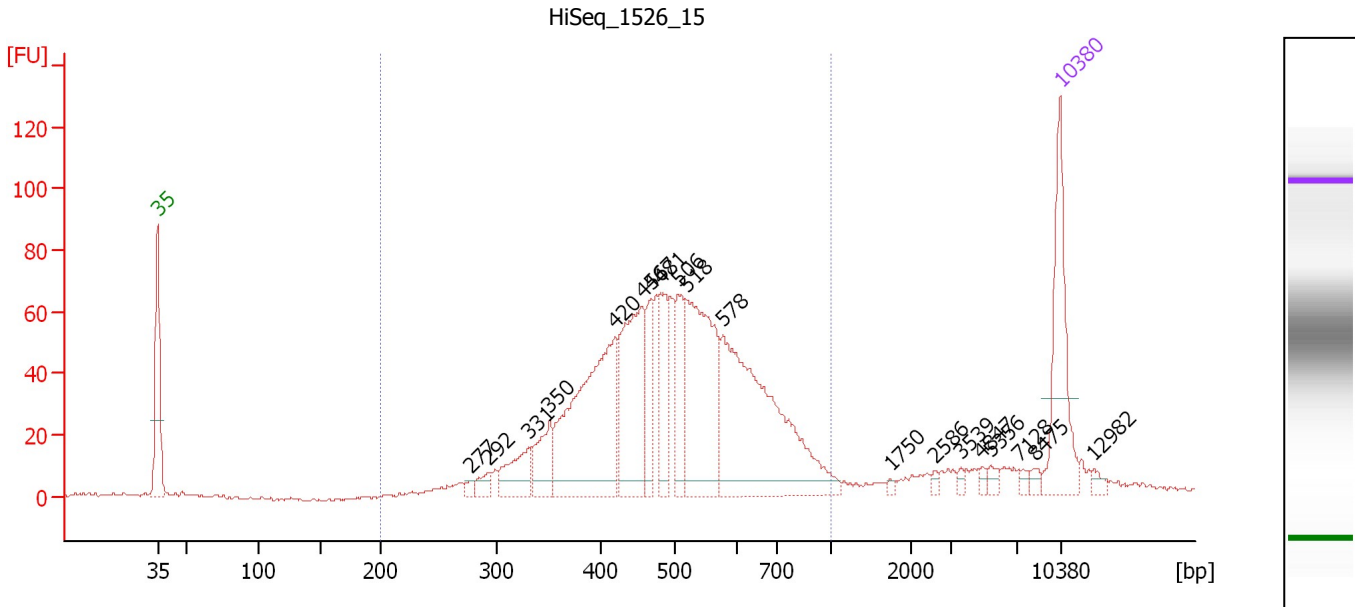
... Region table for sample 3 : HiSeq 1526 14

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	519	691.15	697.0	2,158.4	 86	23.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HiSeq 1526 15

Number of peaks found: 19 Corr. Area 1: 1,239.1
 Noise: 0.4

Peak table for sample 4 : HiSeq 1526 15

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	277	5.00	27.4		67.24
3	292	12.75	66.2		68.62
4	331	42.51	194.7		71.84
5	350	42.44	183.6		73.40
6	420	210.53	760.2		78.51
7	456	134.76	448.0		80.58
8	467	44.68	144.9		81.22
9	481	49.05	154.6		82.00
10	506	47.56	142.5		83.38
11	518	163.89	479.1		83.98
12	578	244.93	641.7		86.83
13	1,750	1.91	1.7		99.89
14	2,586	2.83	1.7		103.30
15	3,539	3.10	1.3		105.27
16	4,847	3.55	1.1		106.93
17	5,356	5.28	1.5		107.57
18	7,128	4.29	0.9		109.78
19	8,475	4.17	0.7		111.11
20	10,380	75.00	10.9	Upper Marker	113.00
21	12,982	0.00	0.0		115.58

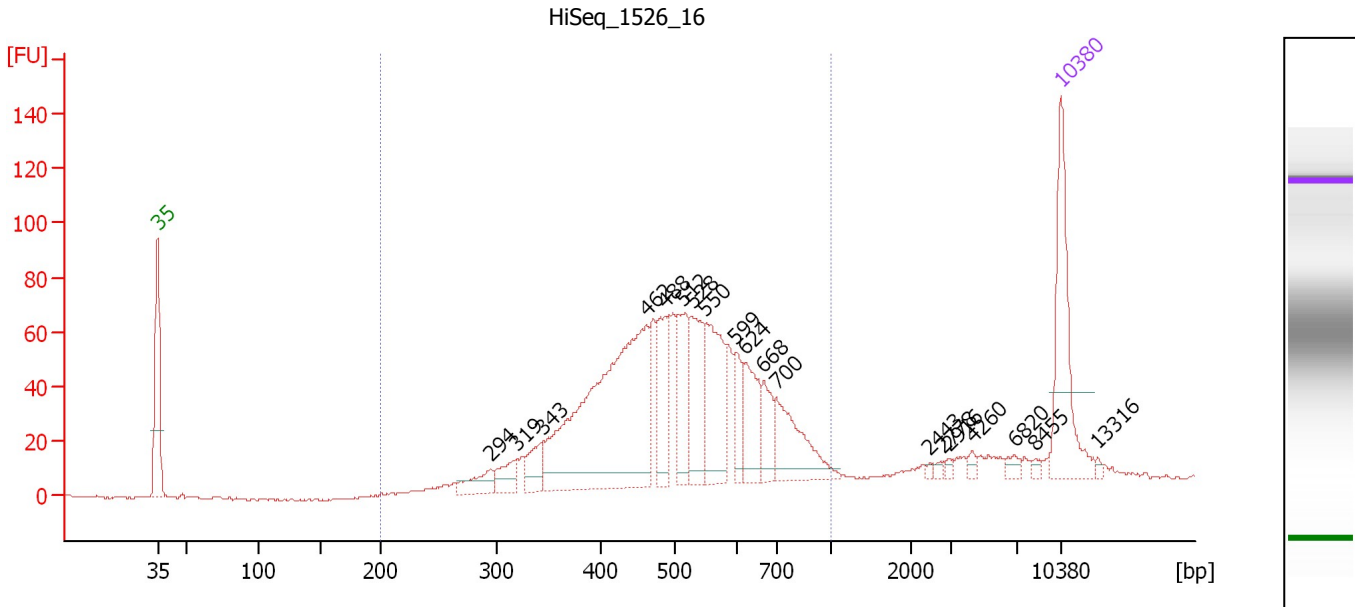
Region table for sample 4 : HiSeq 1526 15

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	506	1,062.89	1,239.1	3,464.7	91	25.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : HiSeq_1526_16

Number of peaks found: 19 Corr. Area 1: 1,237.4
 Noise: 0.2

Peak table for sample 5 : HiSeq_1526_16

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	294	23.00	118.7		68.78
3	319	22.96	109.1		70.88
4	343	25.22	111.4		72.83
5	462	308.40	1,011.5		80.93
6	488	53.41	165.9		82.41
7	512	57.97	171.4		83.69
8	528	62.67	179.7		84.46
9	550	79.05	217.6		85.51
10	599	27.02	68.4		87.79
11	624	46.95	114.1		88.60
12	668	27.45	62.3		89.98
13	700	61.43	133.0		90.99
14	2,443	1.81	1.1		102.85
15	2,778	2.42	1.3		103.90
16	2,916	2.51	1.3		104.33
17	4,260	3.50	1.2		106.19
18	6,820	4.97	1.1		109.43
19	8,455	3.00	0.5		111.09
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,316	0.00	0.0		115.91

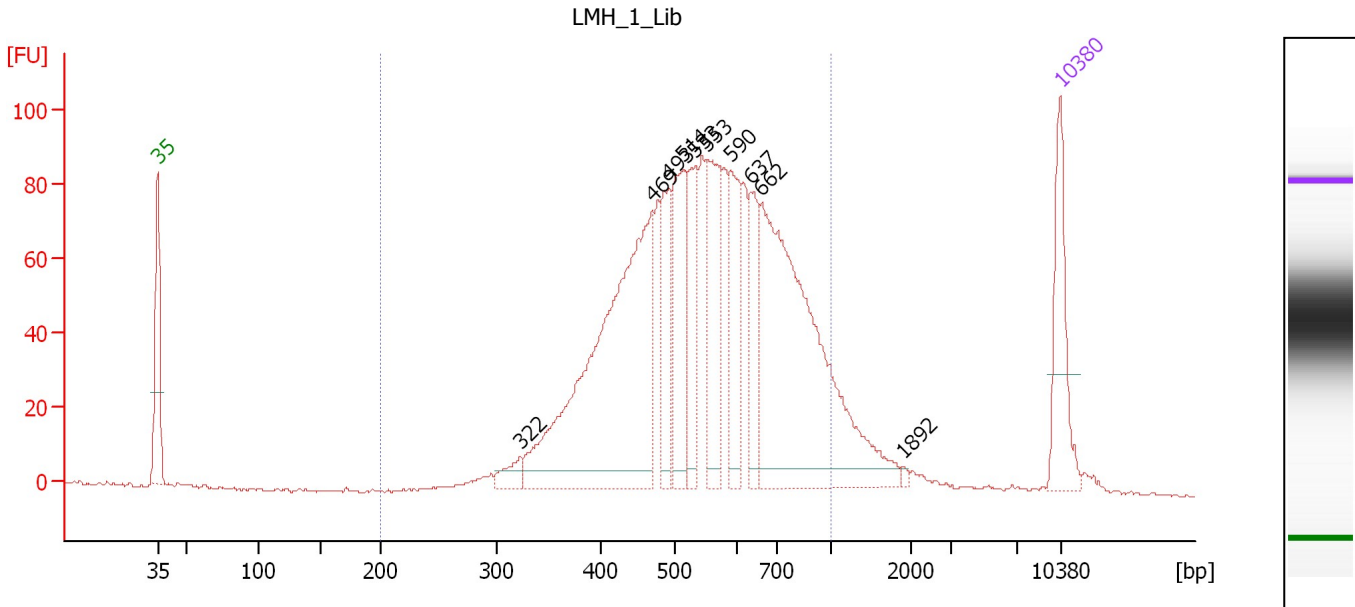
Region table for sample 5 : HiSeq_1526_16

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	515	940.19	1,237.4	3,025.6	89	25.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : LMH_1_Lib

Number of peaks found: 10 Corr. Area 1: 1,693.4
 Noise: 0.4

Peak table for sample 6 : LMH_1_Lib

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	322	23.43	110.3		71.11
3	469	518.06	1,671.9		81.36
4	493	84.71	260.5		82.68
5	514	125.02	368.6		83.76
6	533	86.40	245.4		84.70
7	553	122.35	335.2		85.63
8	590	96.66	248.2		87.39
9	637	67.24	160.0		89.01
10	662	457.55	1,047.3		89.80
11	1,892	3.10	2.5		100.79
12	10,380	75.00	10.9	Upper Marker	113.00

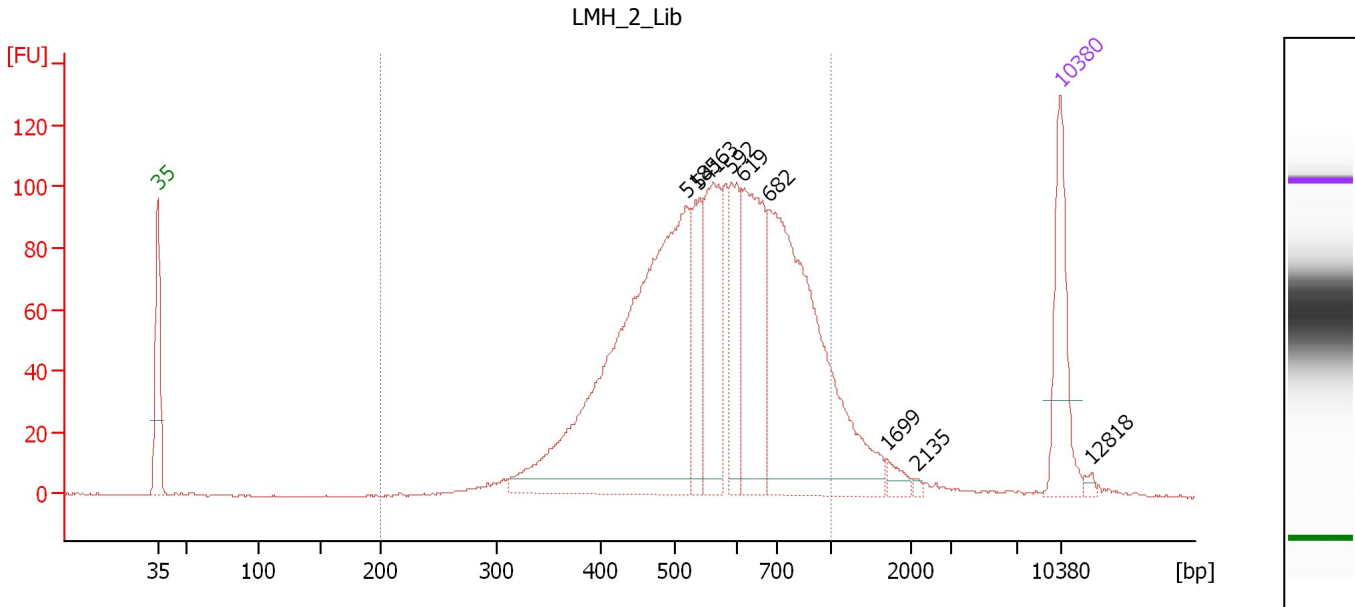
Region table for sample 6 : LMH_1_Lib

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	565	1,804.58	1,693.4	5,284.2	92	26.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

Created: 10/24/2018 3:39:56 PM
 Modified: 10/24/2018 4:21:13 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : LMH_2_Lib

Number of peaks found: 9 Corr. Area 1: 1,884.4
 Noise: 0.3

Peak table for sample 7 : LMH_2_Lib

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	518	604.32	1,766.1		83.98
3	541	82.68	231.7		85.04
4	563	139.97	376.8		86.10
5	592	85.67	219.1		87.50
6	619	167.88	410.8		88.46
7	682	407.79	906.1		90.42
8	1,699	10.95	9.8		99.57
9	2,135	2.42	1.7		101.89
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,818	0.00	0.0		115.41

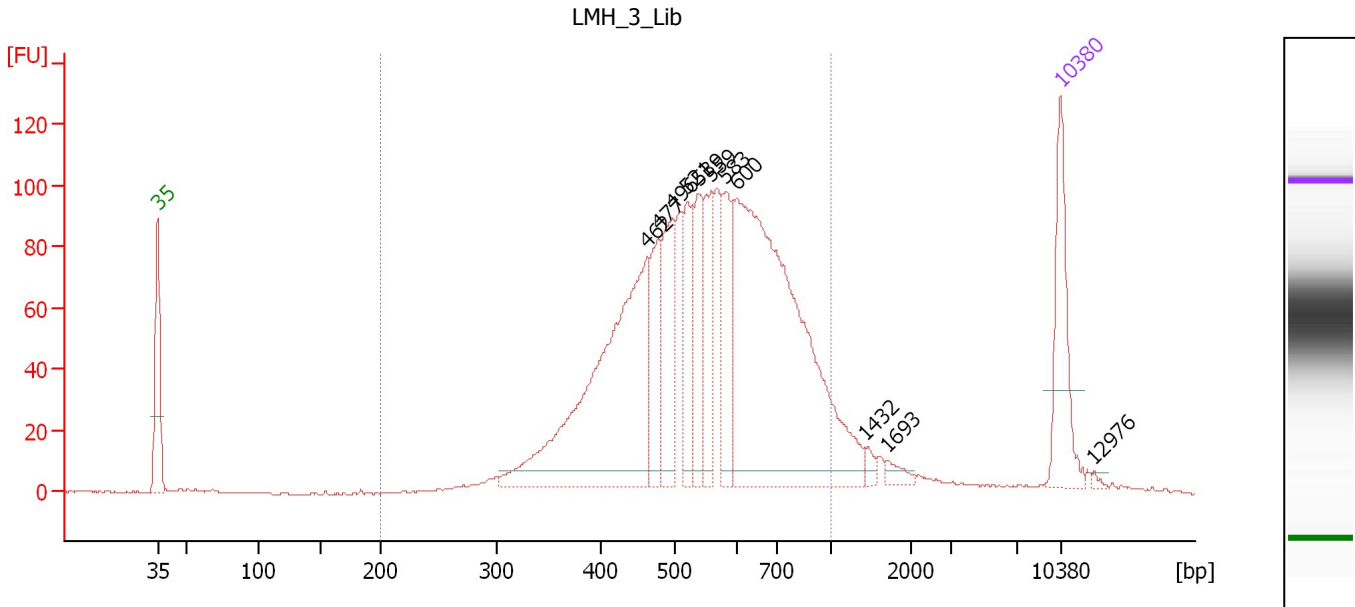
Region table for sample 7 : LMH_2_Lib

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	581	1,529.85	1,884.4	4,373.1	91	26.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : LMH_3 Lib

Number of peaks found: 11 Corr. Area 1: 1,835.9
 Noise: 0.3

Peak table for sample 8 : LMH_3 Lib

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	462	399.51	1,310.3		80.93
3	477	71.87	228.2		81.80
4	496	105.80	323.1		82.88
5	521	75.10	218.3		84.12
6	539	75.10	211.3		84.94
7	559	78.58	213.0		85.91
8	583	75.76	196.9		87.05
9	600	539.02	1,360.7		87.87
10	1,432	7.14	7.5		97.89
11	1,693	9.32	8.3		99.53
12	10,380	75.00	10.9	Upper Marker	113.00
13	12,976	0.00	0.0		115.57

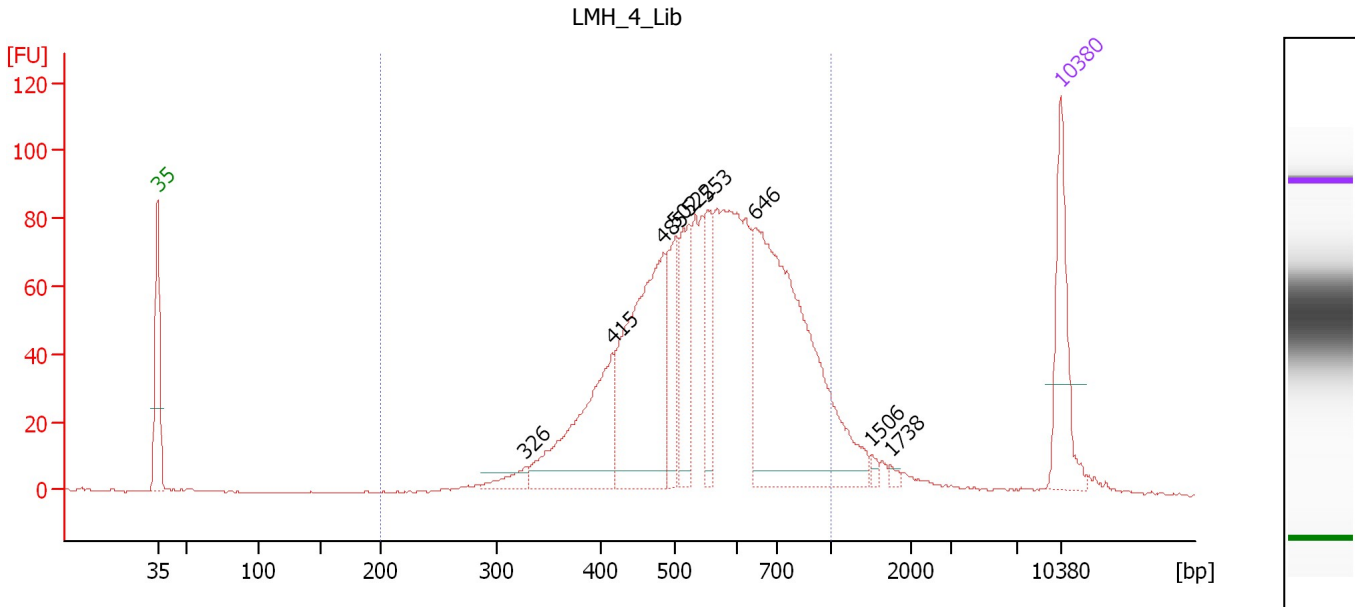
Region table for sample 8 : LMH_3 Lib

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	564	1,617.43	1,835.9	4,753.2	91	25.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : LMH_4_Lib

Number of peaks found: 9 Corr. Area 1: 1,527.4
 Noise: 0.2

Peak table for sample 9 : LMH_4_Lib

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	326	19.97	92.8		71.46
3	415	176.33	644.0		78.23
4	485	279.63	873.6		82.25
5	502	68.41	206.6		83.19
6	522	93.79	272.0		84.18
7	553	62.13	170.2		85.64
8	646	413.74	971.1		89.28
9	1,506	4.48	4.5		98.35
10	1,738	3.76	3.3		99.81
11	10,380	75.00	10.9	Upper Marker	113.00

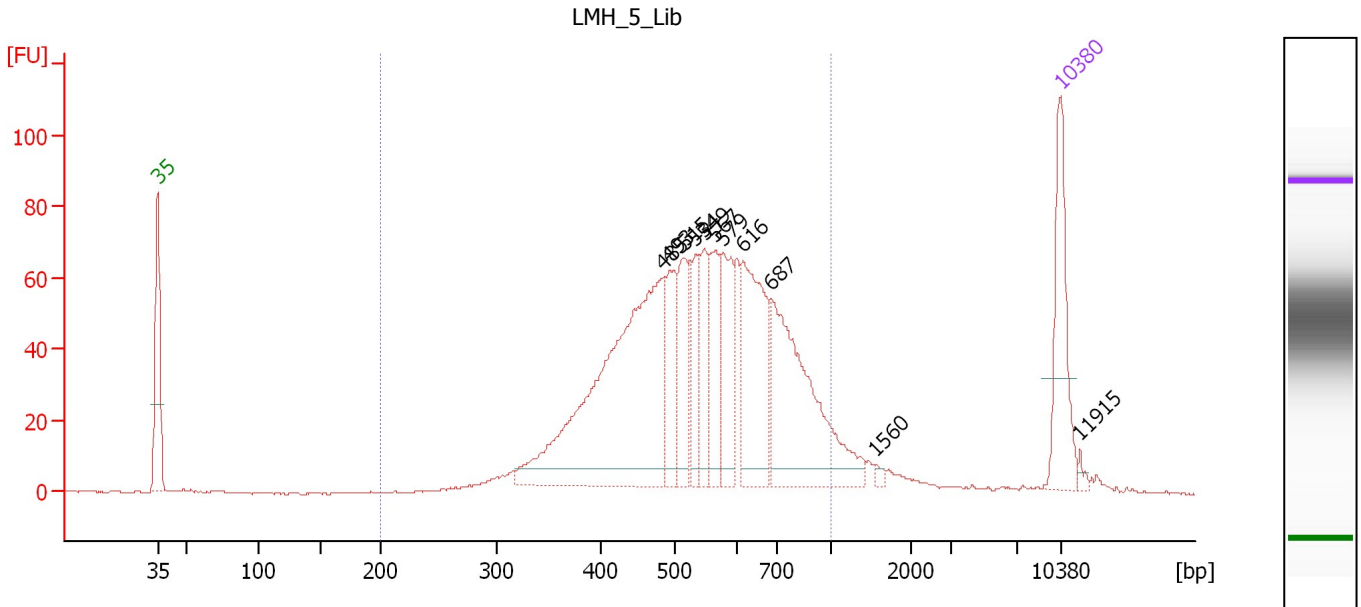
Region table for sample 9 : LMH_4_Lib

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	573	1,497.92	1,527.4	4,327.5	92	25.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : LMH 5 Lib

Number of peaks found: 11 Corr. Area 1: 1,293.5
 Noise: 0.2

Peak table for sample 10 : LMH 5 Lib

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	485	420.60	1,314.2		82.24
3	493	63.08	193.8		82.72
4	515	72.52	213.4		83.82
5	534	51.93	147.4		84.71
6	549	51.93	143.3		85.44
7	567	66.07	176.6		86.28
8	579	71.01	185.9		86.85
9	616	140.90	346.4		88.37
10	687	209.95	463.2		90.57
11	1,560	3.81	3.7		98.70
12	10,380	75.00	10.9	Upper Marker	113.00
13	11,915	0.00	0.0		114.52

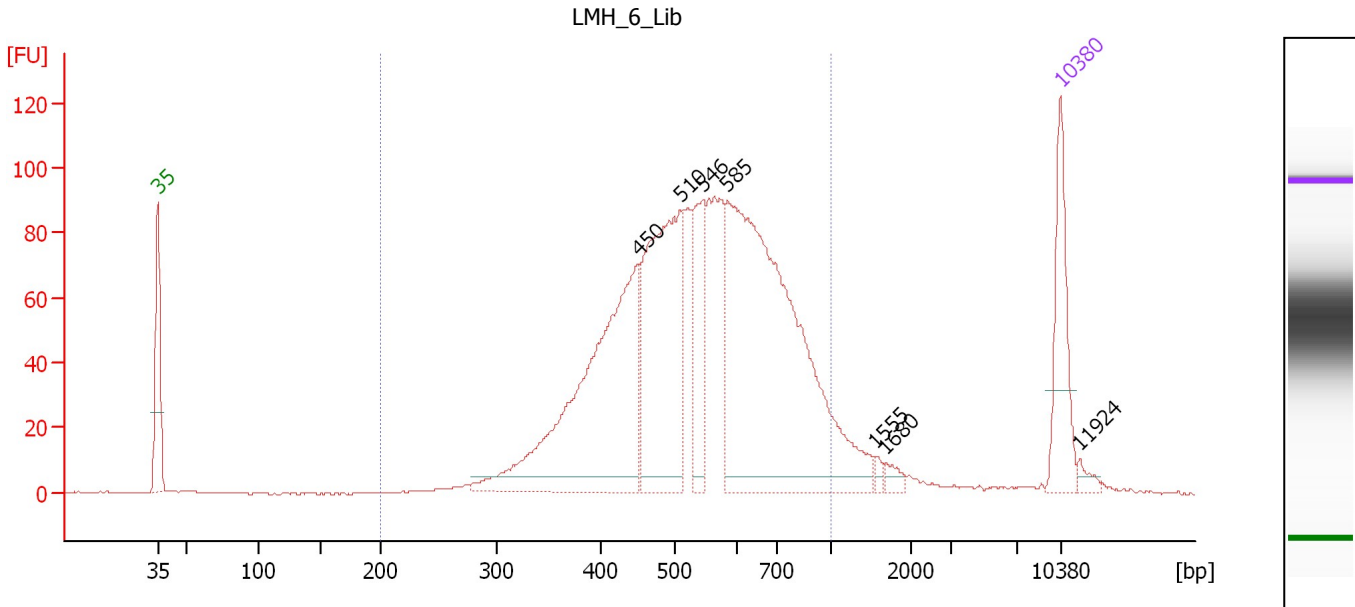
Region table for sample 10 : LMH 5 Lib

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	555	1,305.56	1,293.5	3,908.2	91	26.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : LMH_6 Lib

Number of peaks found: 7 Corr. Area 1: 1,751.0
 Noise: 0.2

Peak table for sample 11 : LMH_6 Lib

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	450	413.56	1,391.3		80.27
3	510	297.22	882.8		83.59
4	546	98.95	274.8		85.27
5	585	570.31	1,475.9		87.17
6	1,555	5.27	5.1		98.66
7	1,680	9.26	8.4		99.45
8	10,380	75.00	10.9	Upper Marker	113.00
9	11,924	0.00	0.0		114.53

Region table for sample 11 : LMH_6 Lib

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	554	1,608.08	1,751.0	4,827.7	91	26.3

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-24\2018-10-24_001.xad

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Gel Image

