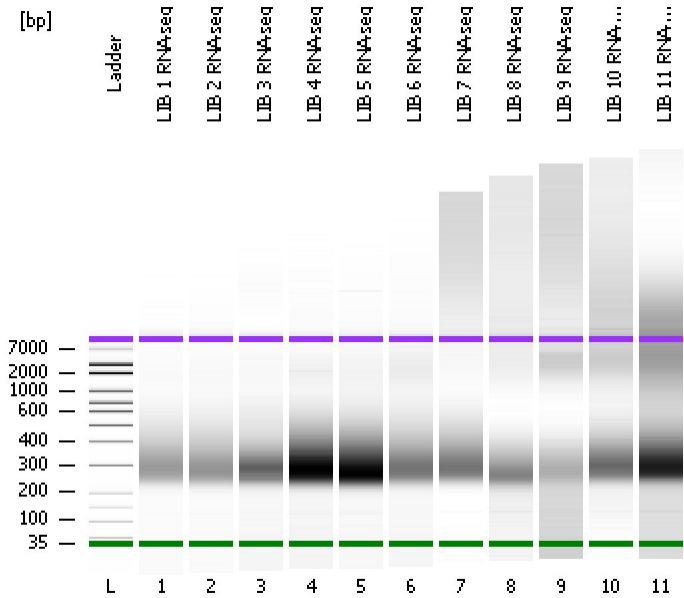


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
Modified: 9/26/2019 12:06:02 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

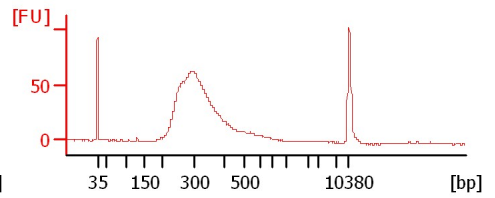
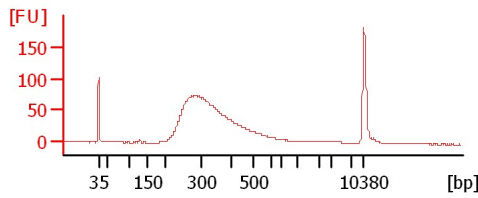
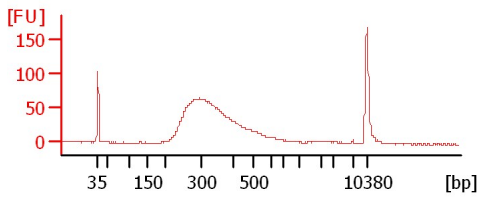
Chip Information:

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

LIB 1 RNA-seq

LIB 2 RNA-seq

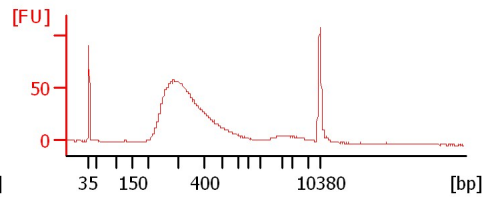
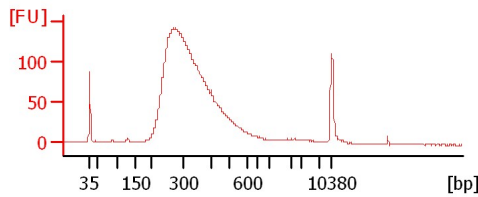
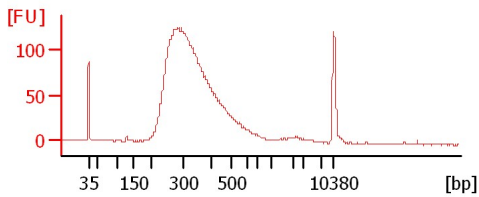
LIB 3 RNA-seq



LIB 4 RNA-seq

LIB 5 RNA-seq

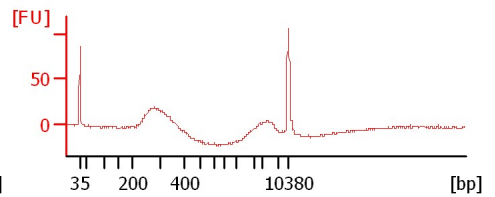
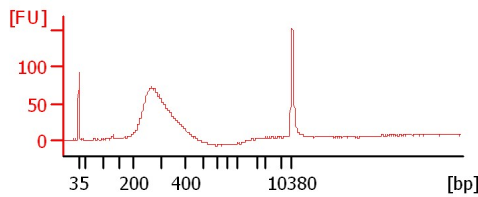
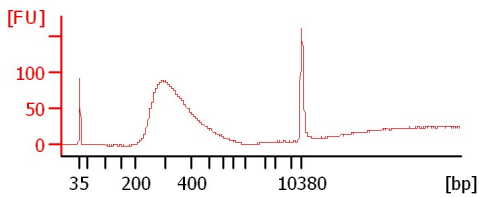
LIB 6 RNA-seq



LIB 7 RNA-seq

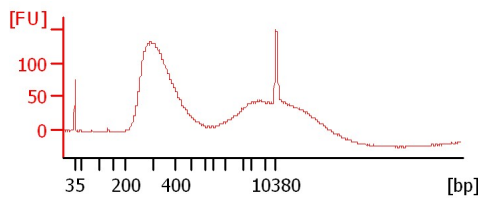
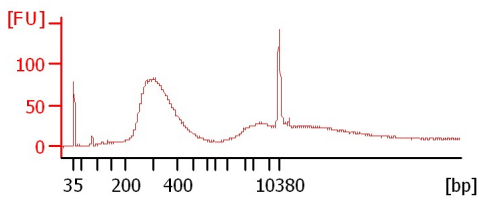
LIB 8 RNA-seq

LIB 9 RNA-seq



LIB 10 RNA-seq

LIB 11 RNA-seq



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
LIB 1 RNA-seq		<input type="checkbox"/>	✓			
LIB 2 RNA-seq		<input type="checkbox"/>	✓			
LIB 3 RNA-seq		<input type="checkbox"/>	✓			
LIB 4 RNA-seq		<input type="checkbox"/>	✓			
LIB 5 RNA-seq		<input type="checkbox"/>	✓			
LIB 6 RNA-seq		<input type="checkbox"/>	✓			
LIB 7 RNA-seq		<input type="checkbox"/>	✓			
LIB 8 RNA-seq		<input type="checkbox"/>	✓			
LIB 9 RNA-seq		<input type="checkbox"/>	✓			
LIB 10 RNA-seq		<input type="checkbox"/>	✓			
LIB 11 RNA-seq		<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\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
Modified: 9/26/2019 12:06:02 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

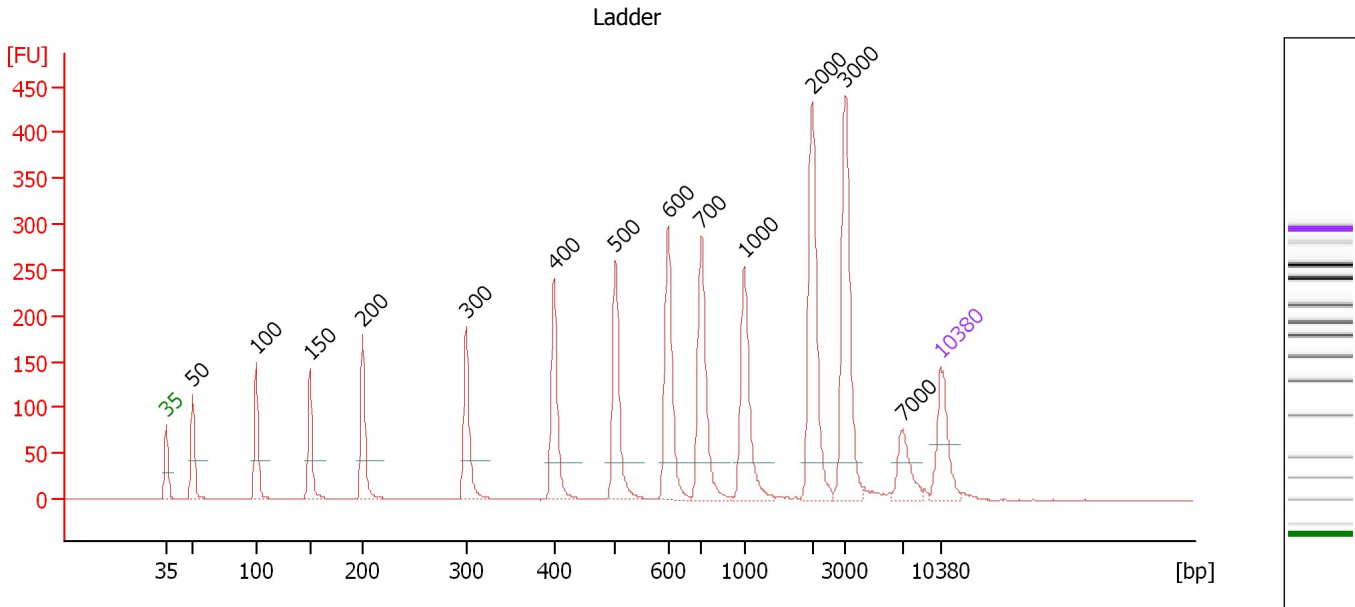
Ladder

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\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

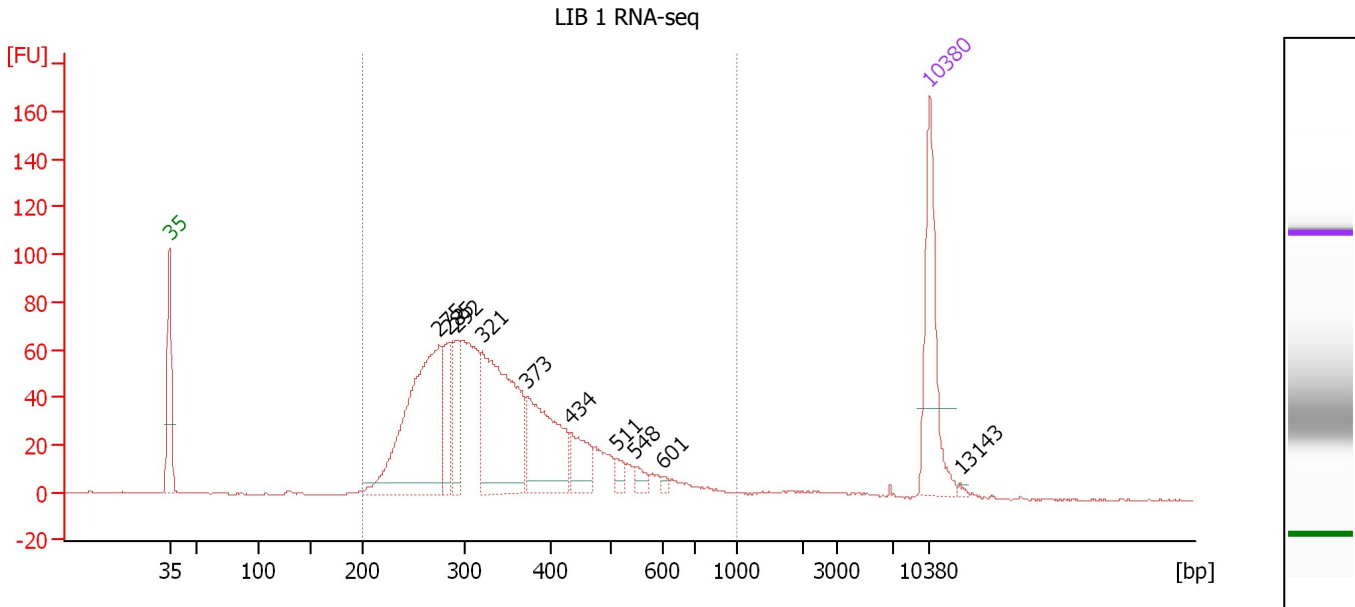
Peak table for Ladder

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	45.41
3	100	150.00	2,272.7	Ladder Peak	51.15
4	150	150.00	1,515.2	Ladder Peak	56.02
5	200	150.00	1,136.4	Ladder Peak	60.79
6	300	150.00	757.6	Ladder Peak	70.14
7	400	150.00	568.2	Ladder Peak	78.05
8	500	150.00	454.5	Ladder Peak	83.59
9	600	150.00	378.8	Ladder Peak	88.37
10	700	150.00	324.7	Ladder Peak	91.40
11	1,000	150.00	227.3	Ladder Peak	95.26
12	2,000	150.00	113.6	Ladder Peak	101.38
13	3,000	150.00	75.8	Ladder Peak	104.42
14	7,000	150.00	32.5	Ladder Peak	109.58
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : LIB 1 RNA-seq

Number of peaks found: 10 Corr. Area 1: 1,372.5
 Noise: 0.3

Peak table for sample 1 : LIB 1 RNA-seq

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	275	251.43	1,383.4		67.84
3	285	55.87	297.2		68.72
4	292	59.25	307.8		69.36
5	321	201.71	951.6		71.81
6	373	110.68	449.2		75.94
7	434	38.05	132.9		79.91
8	511	9.51	28.2		84.14
9	548	8.49	23.5		85.90
10	601	3.52	8.9		88.41
11	10,380	75.00	10.9	Upper Marker	113.00
12	13,143	0.00	0.0		115.80

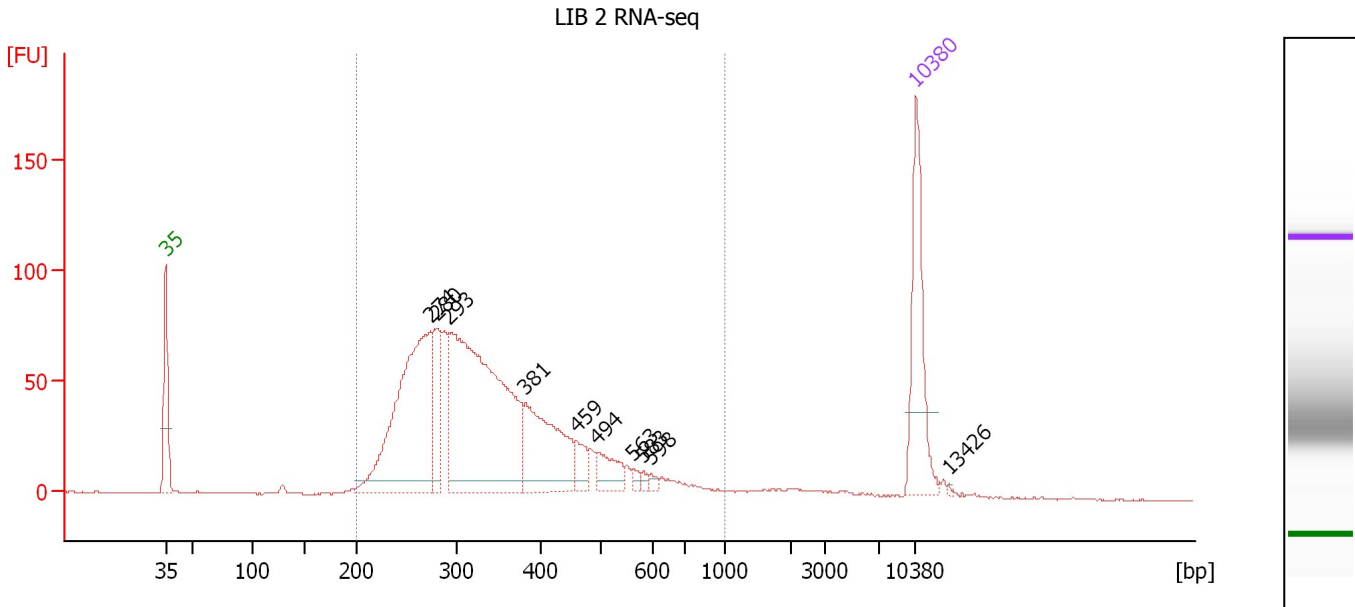
Region table for sample 1 : LIB 1 RNA-seq

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	353	924.96	1,372.5	4,366.4	97	30.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : LIB 2 RNA-seq

Number of peaks found: 10 Corr. Area 1: 1,558.7
 Noise: 0.3

Peak table for sample 2 : LIB 2 RNA-seq

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	274	272.91	1,511.1		67.68
3	280	65.83	356.2		68.27
4	293	384.16	1,988.0		69.47
5	381	115.91	460.4		76.58
6	459	17.99	59.4		81.31
7	494	27.80	85.3		83.25
8	563	4.62	12.5		86.58
9	583	3.28	8.5		87.58
10	598	3.81	9.7		88.27
11	10,380	75.00	10.9	Upper Marker	113.00
12	13,426	0.00	0.0		116.08

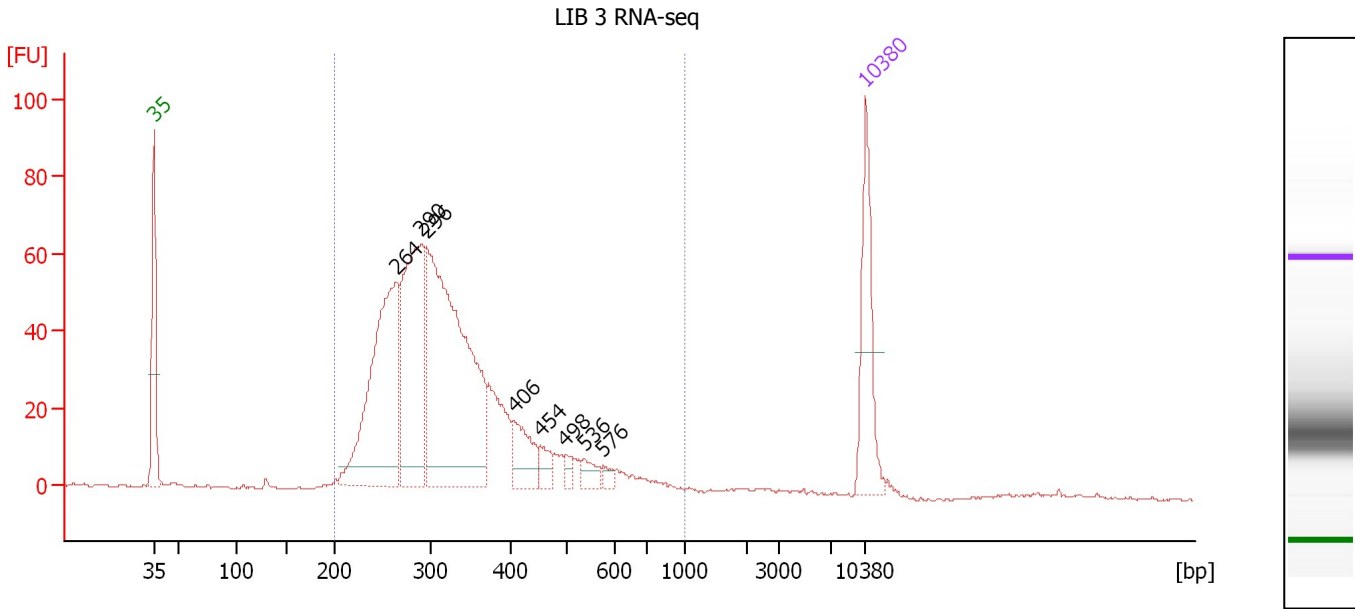
Region table for sample 2 : LIB 2 RNA-seq

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	350	986.16	1,558.7	4,718.2	96	31.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : LIB 3 RNA-seq

Number of peaks found: 8 Corr. Area 1: 1,087.4
 Noise: 0.3

Peak table for sample 3 : LIB 3 RNA-seq

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	264	305.92	1,753.6		66.81
3	290	247.78	1,294.1		69.22
4	296	457.74	2,340.8		69.79
5	406	48.39	180.5		78.39
6	454	18.69	62.4		81.01
7	498	10.07	30.6		83.48
8	536	14.56	41.2		85.31
9	576	6.65	17.5		87.20
10	10,380	75.00	10.9	Upper Marker	113.00

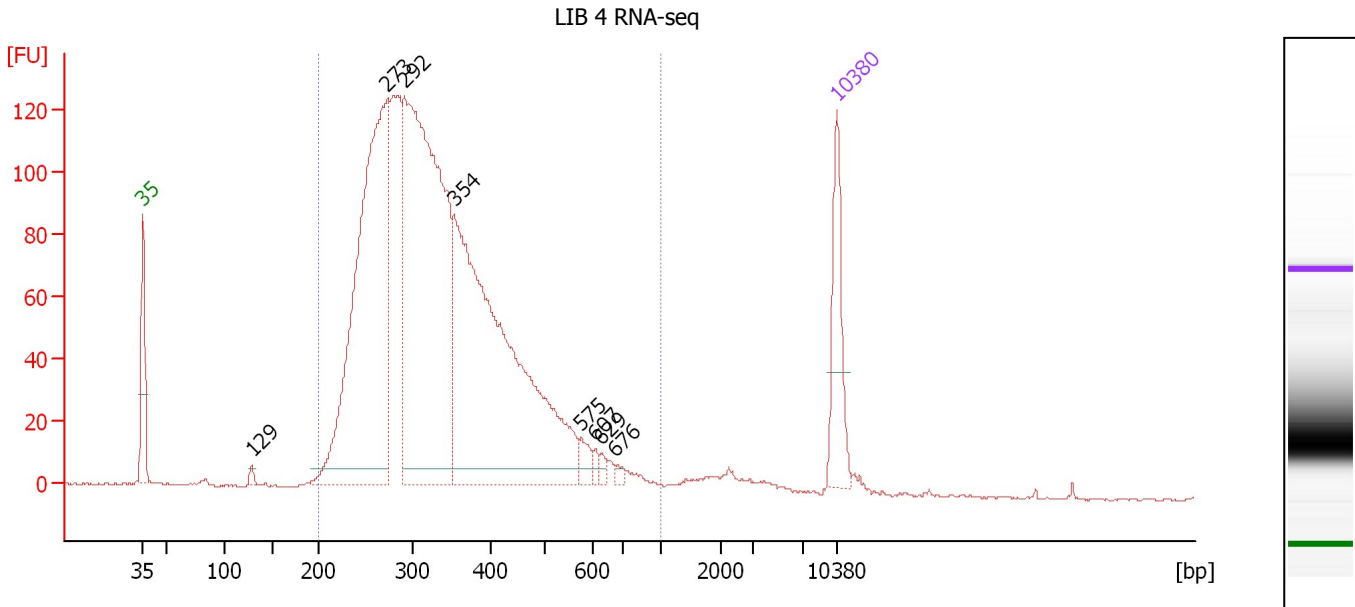
Region table for sample 3 : LIB 3 RNA-seq

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	335	1,261.14	1,087.4	6,198.7	98	29.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : LIB 4 RNA-seq

Number of peaks found: 8 Corr. Area 1: 2,558.2
 Noise: 0.5

Peak table for sample 4 : LIB 4 RNA-seq

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	129	6.16	72.3		53.97
3	273	707.18	3,923.8		67.62
4	292	831.80	4,315.5		69.40
5	354	737.48	3,158.0		74.40
6	575	16.49	43.4		87.19
7	607	7.09	17.7		88.59
8	629	6.89	16.6		89.24
9	676	5.47	12.2		90.69
10	10,380	75.00	10.9	Upper Marker	113.00

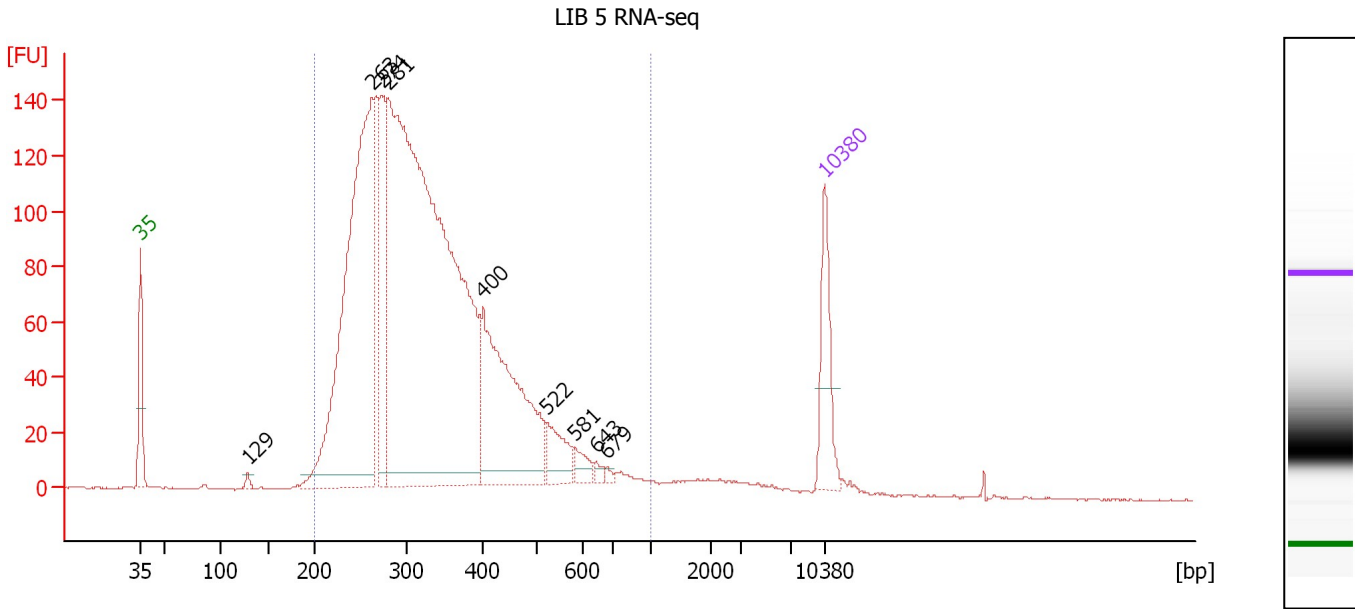
Region table for sample 4 : LIB 4 RNA-seq

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	345	2,575.02	2,558.2	12,375.9	97	29.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : LIB 5 RNA-seq

Number of peaks found: 9 Corr. Area 1: 2,849.6
 Noise: 0.2

Peak table for sample 5 : LIB 5 RNA-seq

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	129	8.36	98.2		53.98
3	263	806.95	4,645.8		66.70
4	274	171.51	947.5		67.73
5	281	1,660.34	8,962.6		68.34
6	400	339.76	1,286.6		78.05
7	522	55.27	160.3		84.66
8	581	21.06	54.9		87.45
9	643	7.59	17.9		89.68
10	679	4.51	10.1		90.78
11	10,380	75.00	10.9	Upper Marker	113.00

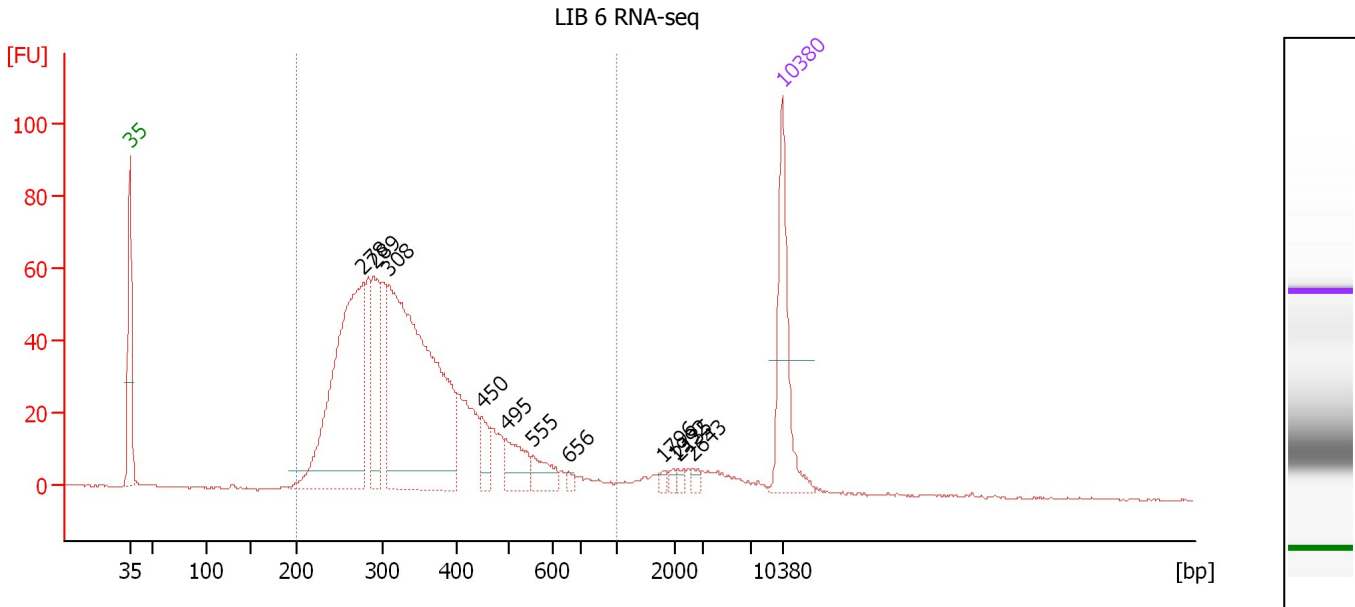
Region table for sample 5 : LIB 5 RNA-seq

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	340	3,141.70	2,849.6	15,389.3	96	30.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : LIB 6 RNA-seq

Number of peaks found: 11 Corr. Area 1: 1,165.8
 Noise: 0.3

Peak table for sample 6 : LIB 6 RNA-seq

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	278	328.77	1,794.3		68.05
3	289	84.16	440.9		69.14
4	308	443.79	2,181.4		70.79
5	450	21.78	73.4		80.80
6	495	35.87	109.8		83.32
7	555	23.16	63.2		86.24
8	656	4.19	9.7		90.07
9	1,796	3.39	2.9		100.13
10	1,992	3.08	2.3		101.33
11	2,135	3.29	2.3		101.79
12	2,643	3.97	2.3		103.33
13	10,380	75.00	10.9	Upper Marker	113.00

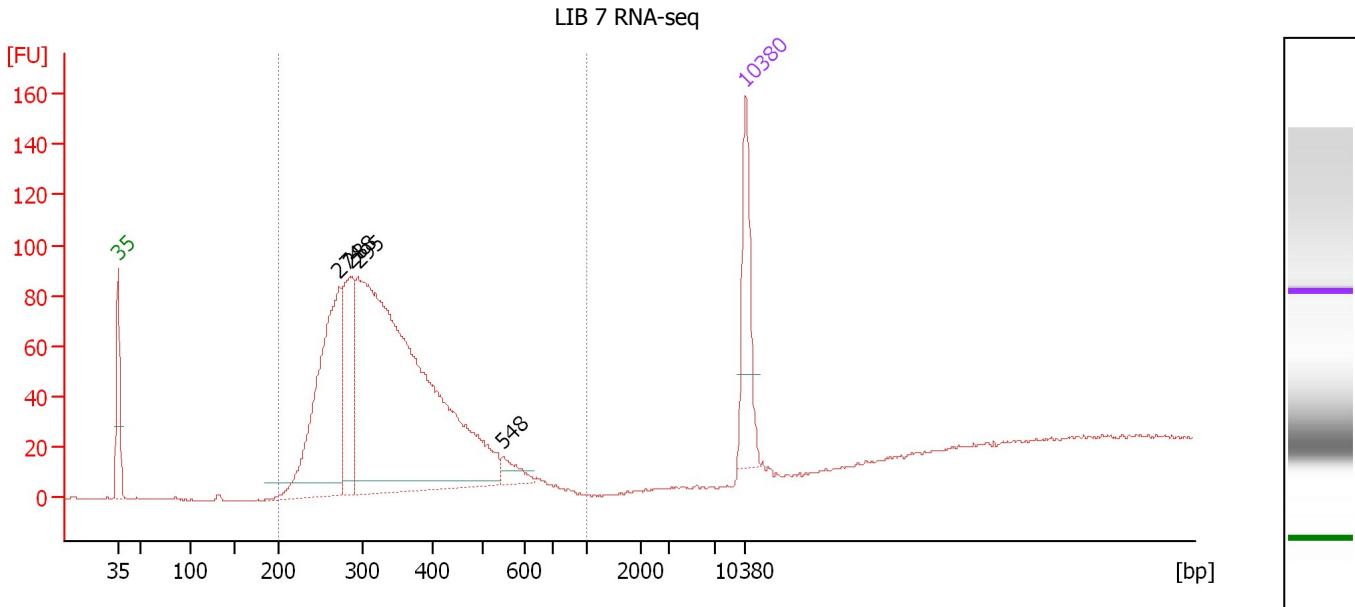
Region table for sample 6 : LIB 6 RNA-seq

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	352	1,162.64	1,165.8	5,517.2	92	31.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : LIB 7 RNA-seq

Number of peaks found: 4 Corr. Area 1: 1,402.3
 Noise: 0.2

Peak table for sample 7 : LIB 7 RNA-seq

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	274	331.47	1,835.0		67.68
3	288	155.03	816.5		68.99
4	295	905.14	4,644.0		69.70
5	548	18.71	51.7		85.88
6	10,380	75.00	10.9	Upper Marker	113.00

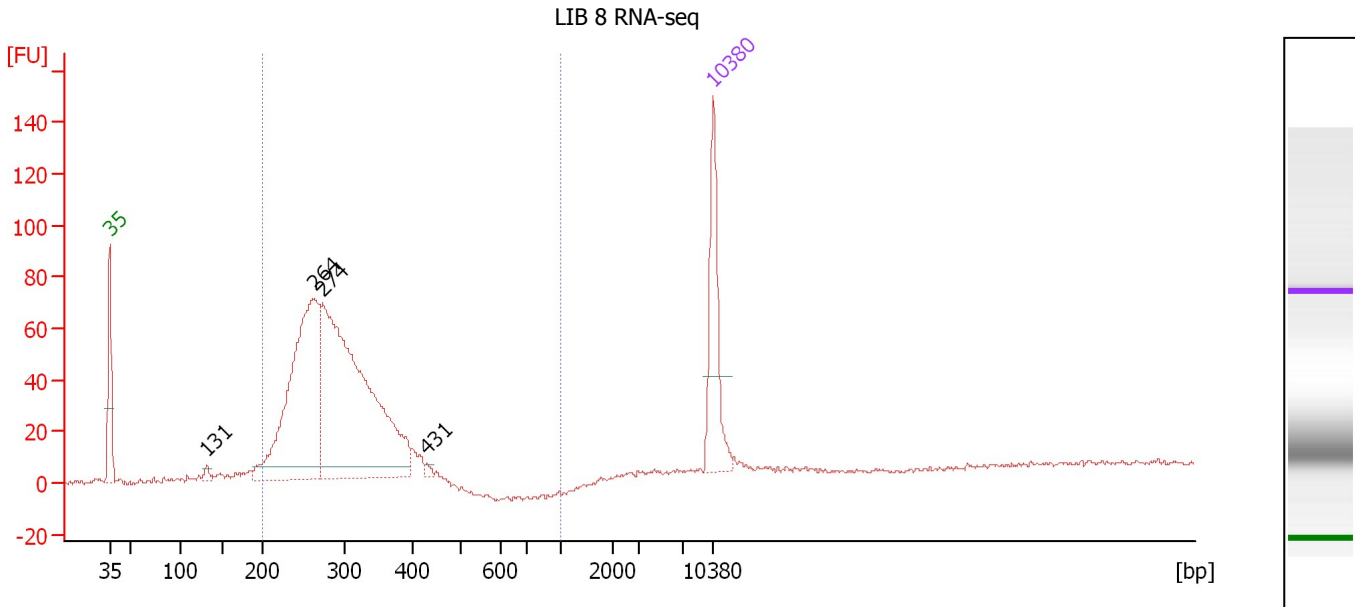
Region table for sample 7 : LIB 7 RNA-seq

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	334	1,164.11	1,402.3	5,574.9	98	20.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : LIB 8 RNA-seq

Number of peaks found: 4 Corr. Area 1: 979.3
 Noise: 0.9

Peak table for sample 8 : LIB 8 RNA-seq

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	131	8.94	103.2		54.19
3	264	350.61	2,014.6		66.75
4	274	496.97	2,752.1		67.67
5	431	3.87	13.6		79.79
6	10,380	75.00	10.9	Upper Marker	113.00

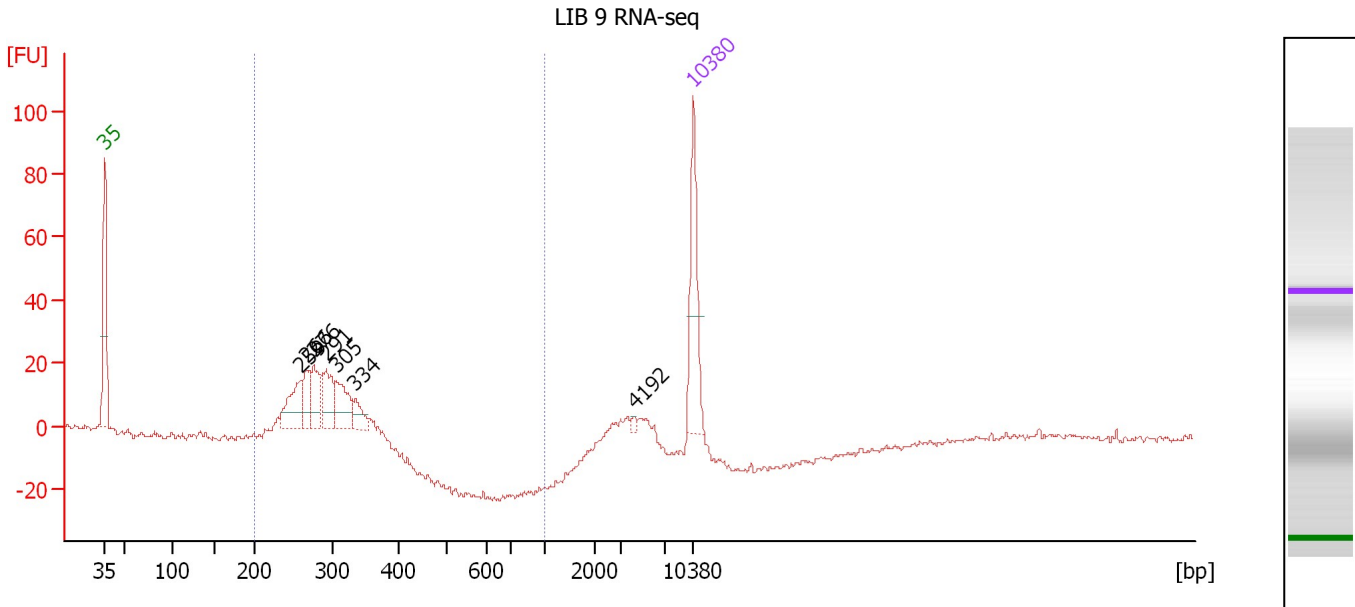
Region table for sample 8 : LIB 8 RNA-seq

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	292	831.28	979.3	4,447.1	93	16.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : LIB 9 RNA-seq

Number of peaks found: 7 Corr. Area 1: 190.6
 Noise: 0.9

Peak table for sample 9 : LIB 9 RNA-seq

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	259	60.26	352.8		66.29
3	266	34.08	193.9		66.99
4	276	45.98	252.6		67.88
5	291	44.97	233.9		69.34
6	305	50.67	251.6		70.54
7	334	21.67	98.3		72.83
8	4,192	3.16	1.1		105.96
9	10,380	75.00	10.9	Upper Marker	113.00

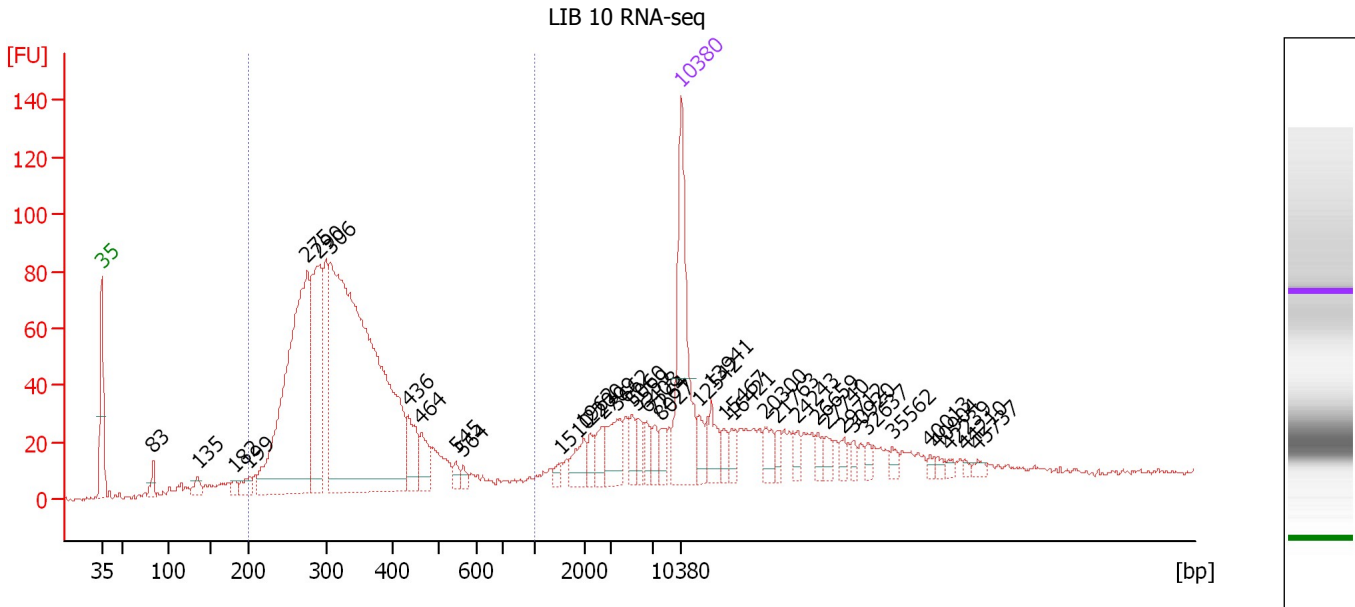
Region table for sample 9 : LIB 9 RNA-seq

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	287	263.65	190.6	1,406.9	90	10.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : LIB 10 RNA-seq

Number of peaks found: 39 Corr. Area 1: 1,458.3
 Noise: 0.9

Peak table for sample 10 : LIB 10 RNA-seq

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	83	10.74	196.4		49.18
3	135	9.66	108.1		54.59
4	182	5.28	44.1		59.03
5	199	11.09	84.4		60.71
6	275	277.82	1,531.3		67.79
7	290	125.79	657.1		69.21
8	306	508.60	2,517.3		70.63
9	436	25.84	89.9		80.03
10	464	18.59	60.7		81.57
11	545	5.71	15.9		85.76
12	564	4.42	11.9		86.66
13	1,510	3.63	3.6		98.38
14	1,962	13.70	10.6		101.15
15	2,200	6.77	4.7		101.99
16	2,709	11.56	6.5		103.53
17	3,662	19.96	8.3		105.27
18	5,060	8.94	2.7		107.08
19	5,559	8.59	2.3		107.72
20	6,408	7.95	1.9		108.81
21	7,264	7.45	1.6		109.84
22	8,027	7.14	1.3		110.62
23	10,380	75.00	10.9	Upper Marker	113.00
24	12,542	0.00	0.0		115.19
25	13,941	0.00	0.0		116.61

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...

... Peak table for sample 10 : LIB 10 RNA-seq

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	15,467	0.00	0.0		118.15
27	16,421	0.00	0.0		119.12
28	20,300	0.00	0.0		123.05
29	21,763	0.00	0.0		124.53
30	24,243	0.00	0.0		127.04
31	26,659	0.00	0.0		129.49
32	27,740	0.00	0.0		130.58
33	29,712	0.00	0.0		132.58
34	30,920	0.00	0.0		133.80
35	32,637	0.00	0.0		135.54
36	35,562	0.00	0.0		138.50
37	40,013	0.00	0.0		143.01
38	40,904	0.00	0.0		143.91
39	42,239	0.00	0.0		145.26
40	44,210	0.00	0.0		147.26
41	45,737	0.00	0.0		148.80

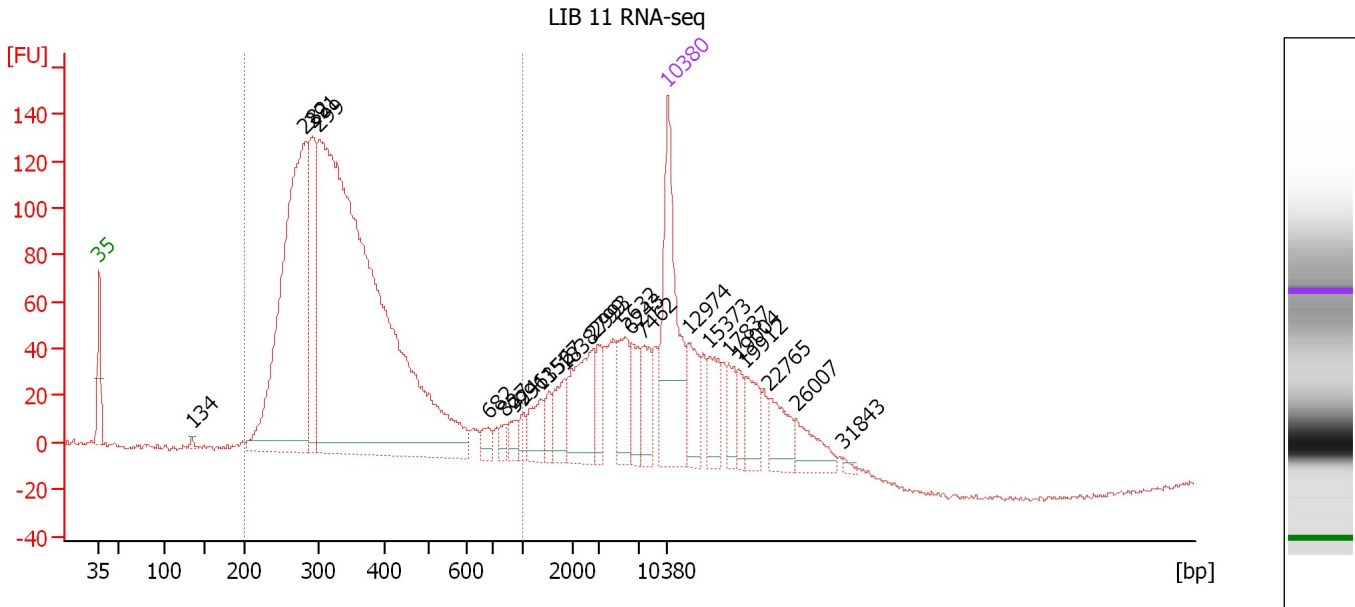
Region table for sample 10 : LIB 10 RNA-seq

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	341	1,051.88	1,458.3	5,039.7	67	28.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : LIB 11 RNA-seq

Number of peaks found: 24 Corr. Area 1: 2,417.7
 Noise: 1.1

Peak table for sample 11 : LIB 11 RNA-seq

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	134	2.80	31.6		54.49
3	282	316.32	1,702.5		68.41
4	291	96.43	501.5		69.33
5	299	745.92	3,779.0		70.05
6	682	8.22	18.2		90.87
7	827	6.04	11.1		93.04
8	924	7.60	12.5		94.29
9	996	6.63	10.1		95.20
10	1,356	16.42	18.4		97.44
11	1,527	8.95	8.9		98.49
12	1,838	19.44	16.0		100.39
13	2,799	41.33	22.4		103.81
14	2,993	12.08	6.1		104.40
15	5,632	25.44	6.8		107.81
16	6,243	17.83	4.3		108.60
17	7,462	19.15	3.9		110.05
18	10,380	75.00	10.9	Upper Marker	113.00
19	12,974	0.00	0.0		115.63
20	15,373	0.00	0.0		118.06
21	17,837	0.00	0.0		120.55
22	19,004	0.00	0.0		121.73
23	19,912	0.00	0.0		122.65
24	22,765	0.00	0.0		125.54
25	26,007	0.00	0.0		128.83

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
Modified: 9/26/2019 12:06:02 PM

Electropherogram Summary Continued ...**... Peak table for sample 11 : LIB 11 RNA-seq**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	31,843	0.00	0.0		134.74

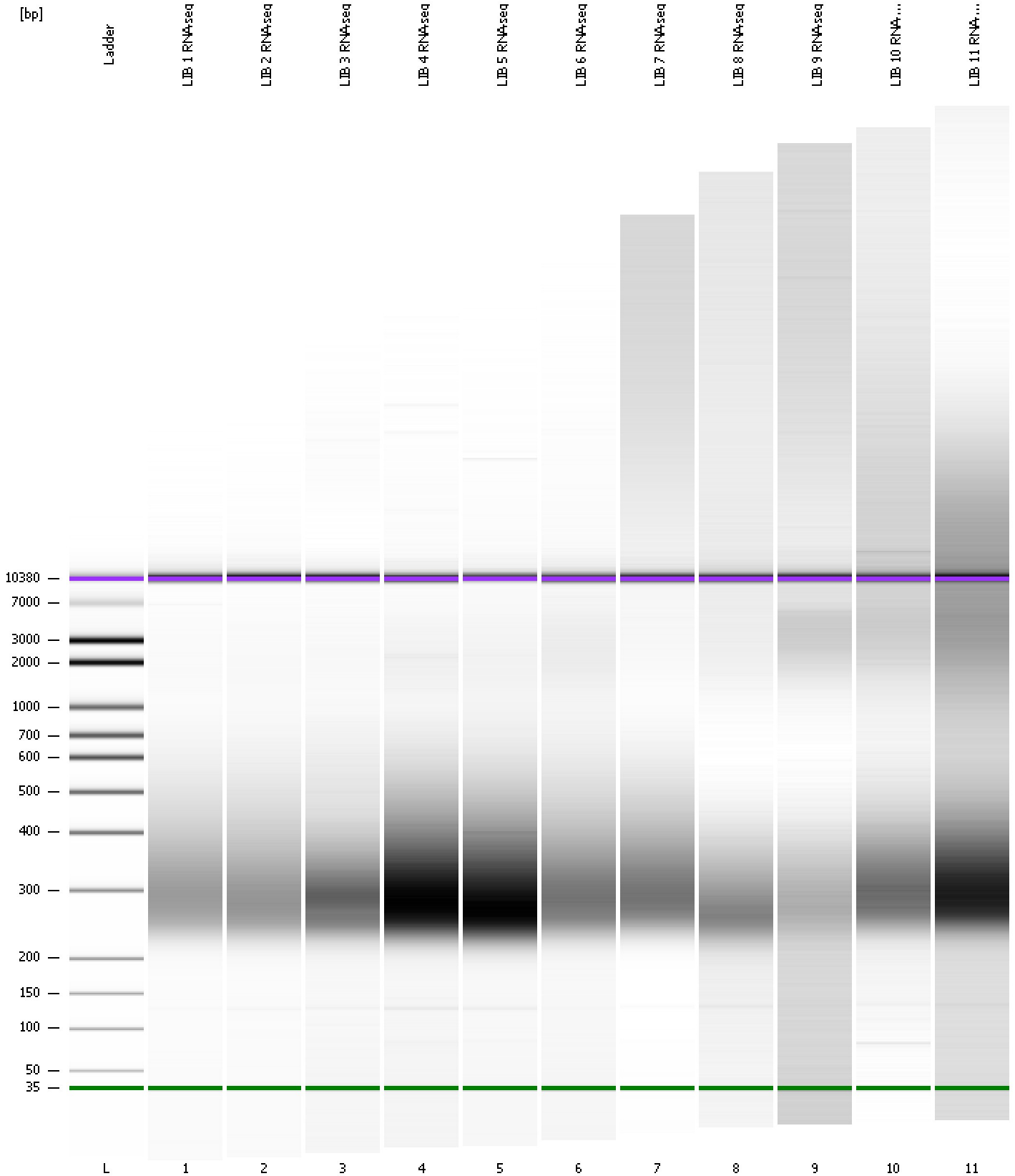
Region table for sample 11 : LIB 11 RNA-seq

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	359	1,083.71	2,417.7	5,058.5	 69	34.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
Modified: 9/26/2019 12:06:02 PM

Gel Image

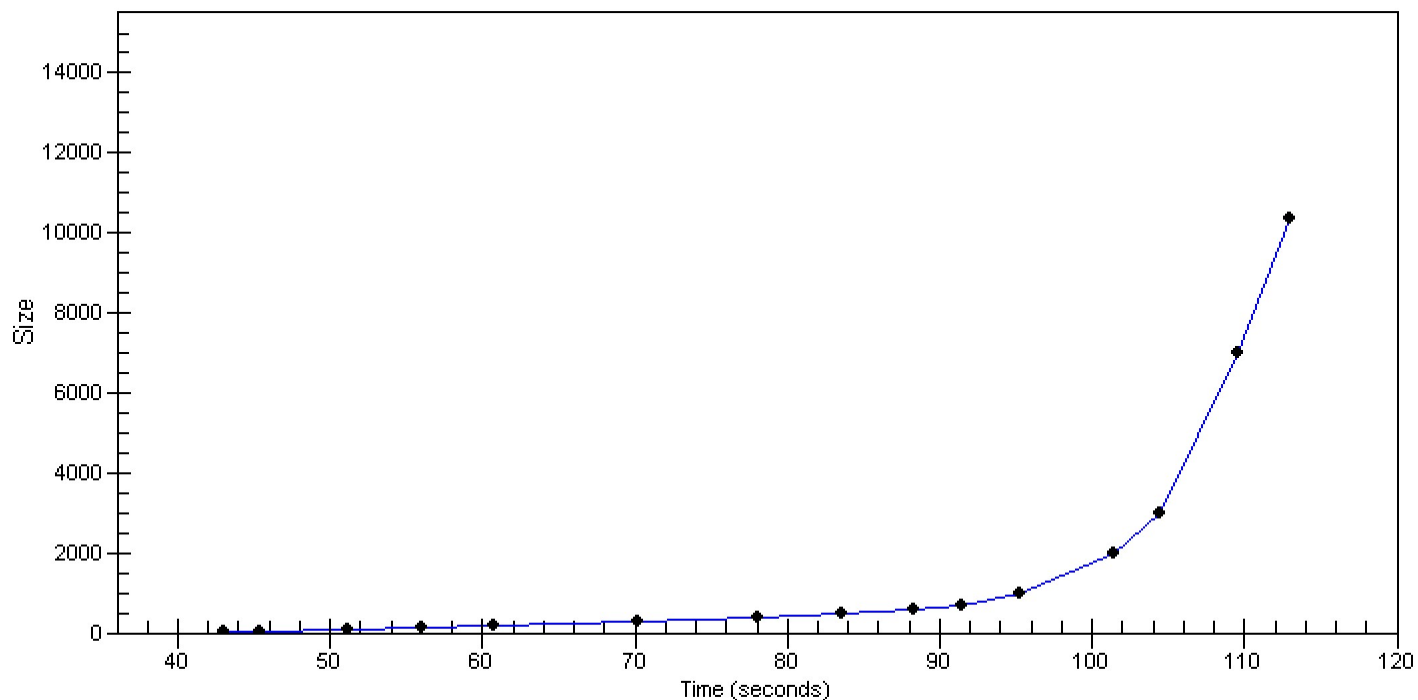


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
Modified: 9/26/2019 12:06:02 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-26\2019-09-26_001.xad

Created: 9/26/2019 11:18:50 AM
 Modified: 9/26/2019 12:06:02 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		9/26/2019 12:00:07 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\2019-09-26\2019-09-26_001.xad)		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		9/26/2019 11:18:56 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1