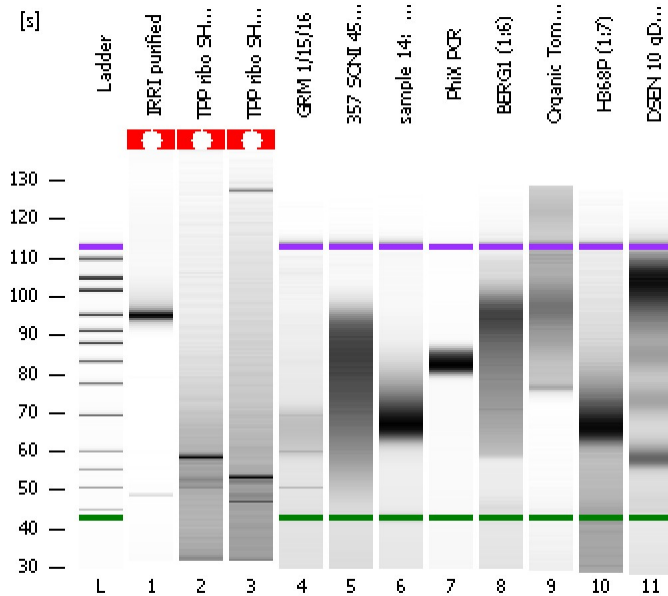


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
Modified: 1/27/2016 3:47:39 PM

Electrophoresis File Run Summary



Instrument Information:

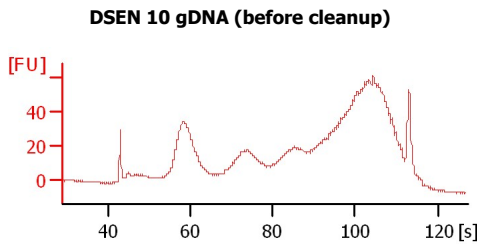
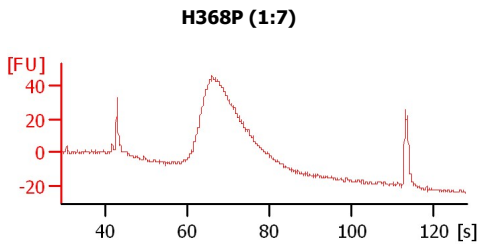
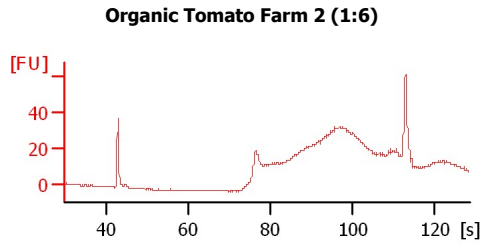
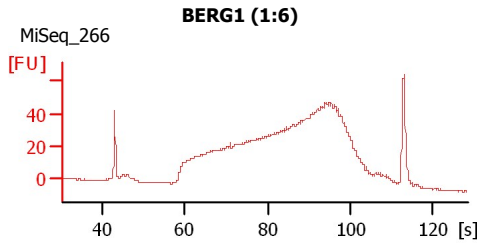
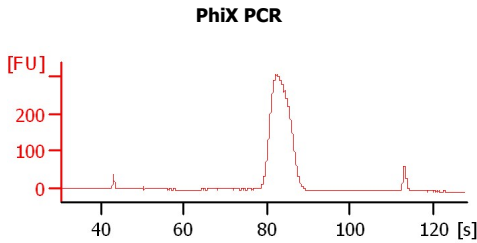
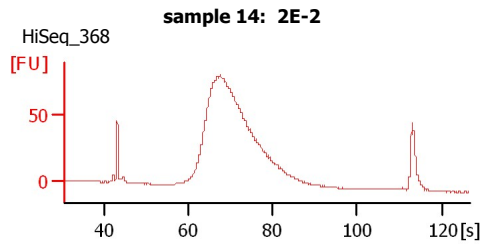
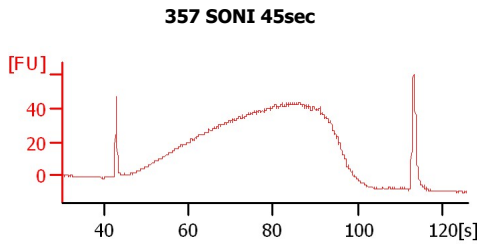
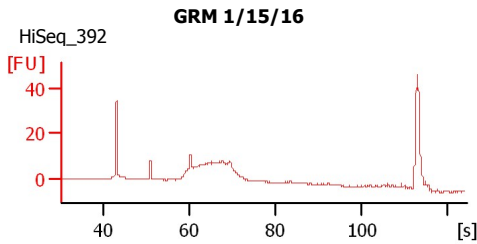
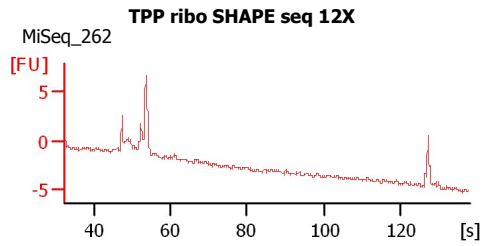
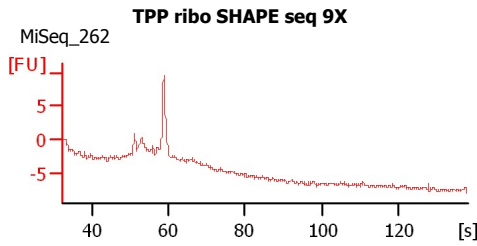
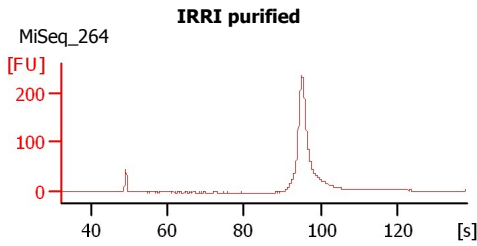
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
IRRI purified	MiSeq_264	<input type="checkbox"/>	✓			
TPP ribo SHAPE seq 9X	MiSeq_262	<input type="checkbox"/>	✓			
TPP ribo SHAPE seq 12X	MiSeq_262	<input type="checkbox"/>	✓			
GRM 1/15/16	HiSeq_392	<input type="checkbox"/>	✓			
357 SONI 45sec		<input type="checkbox"/>	✓			
sample 14: 2E-2	HiSeq_368	<input type="checkbox"/>	✓			
PhiX PCR		<input type="checkbox"/>	✓			
BERG1 (1:6)	MiSeq_266	<input type="checkbox"/>	✓			
Organic Tomato Farm 2 (1:6)		<input type="checkbox"/>	✓			
H368P (1:7)		<input type="checkbox"/>	✓			
DSEN 10 gDNA (before cleanup)		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
Modified: 1/27/2016 3:47:39 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

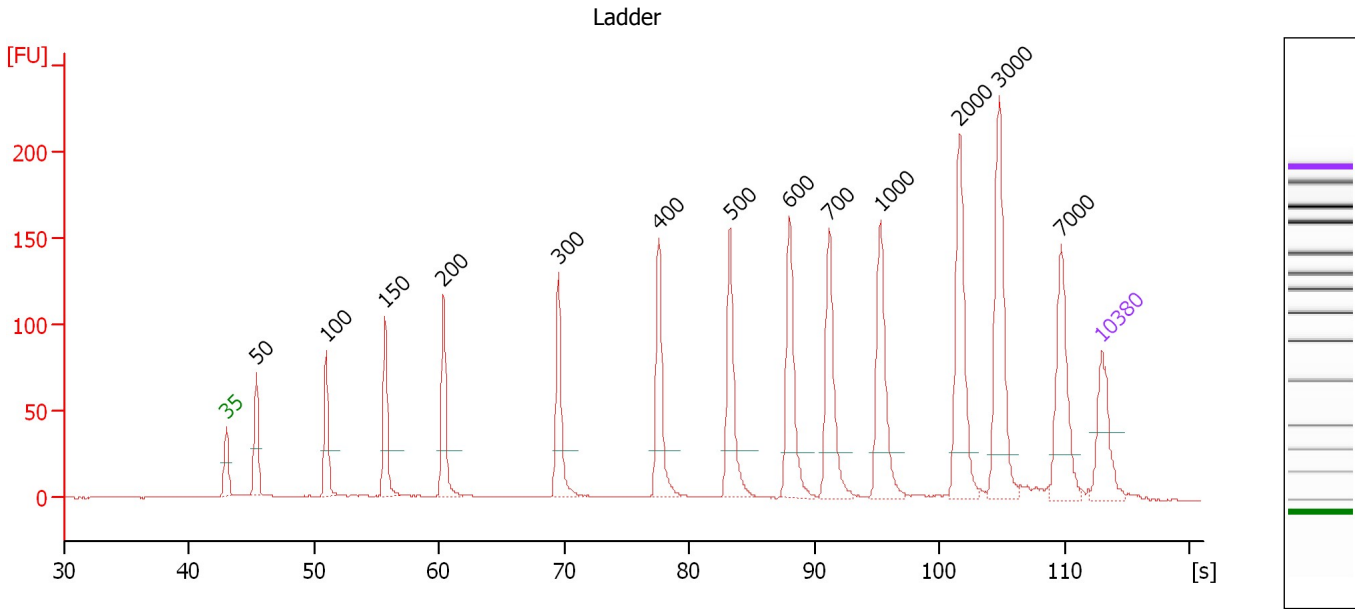
Ladder

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:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

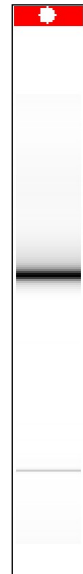
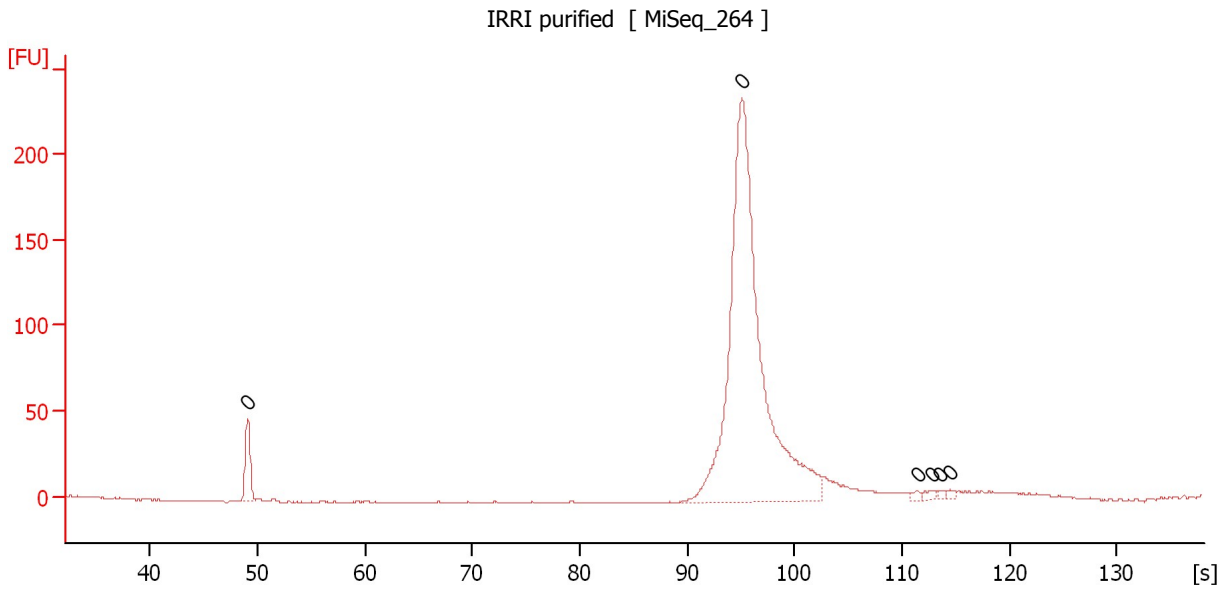
Peak table for Ladder

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	45.36
3	100	150.00	2,272.7	Ladder Peak	50.99
4	150	150.00	1,515.2	Ladder Peak	55.67
5	200	150.00	1,136.4	Ladder Peak	60.35
6	300	150.00	757.6	Ladder Peak	69.54
7	400	150.00	568.2	Ladder Peak	77.57
8	500	150.00	454.5	Ladder Peak	83.28
9	600	150.00	378.8	Ladder Peak	88.01
10	700	150.00	324.7	Ladder Peak	91.18
11	1,000	150.00	227.3	Ladder Peak	95.31
12	2,000	150.00	113.6	Ladder Peak	101.62
13	3,000	150.00	75.8	Ladder Peak	104.80
14	7,000	150.00	32.5	Ladder Peak	109.78
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 1 : IRRI purified

Number of peaks found: 0 Noise: 0.2

Peak table for sample 1 : IRRI purified

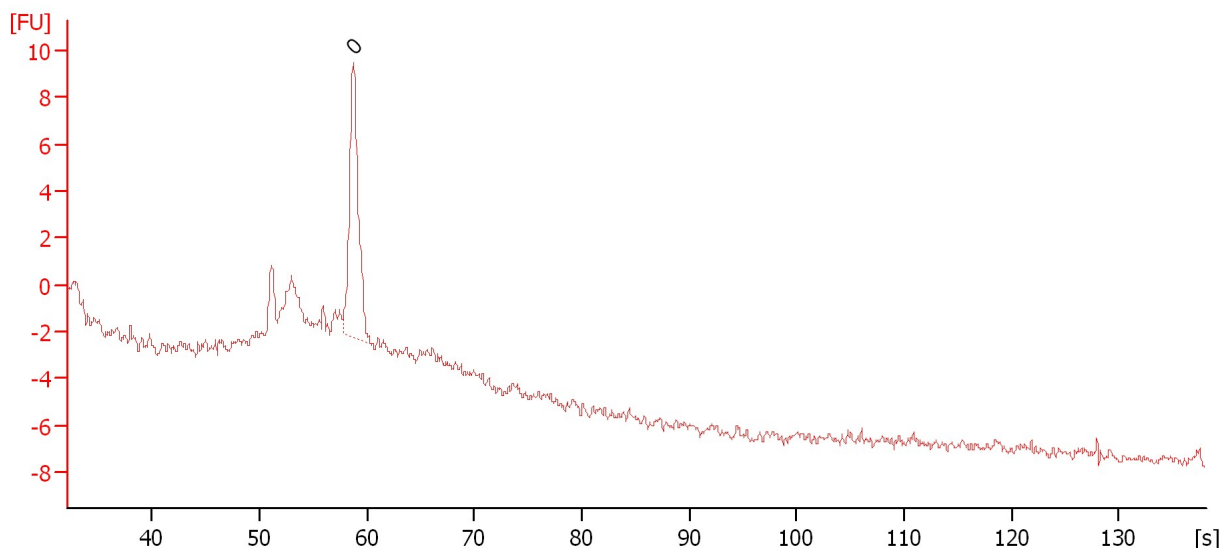
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		49.05
2	0	0.00	0.0		95.10
3	0	0.00	0.0		111.40
4	0	0.00	0.0		112.85
5	0	0.00	0.0		113.55
6	0	0.00	0.0		114.45

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
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Electropherogram Summary Continued ...

TPP ribo SHAPE seq 9X [MiSeq_262]



Overall Results for sample 2 : TPP ribo SHAPE seq 9X

Number of peaks found: 0 Noise: 0.2

Peak table for sample 2 : TPP ribo SHAPE seq 9X

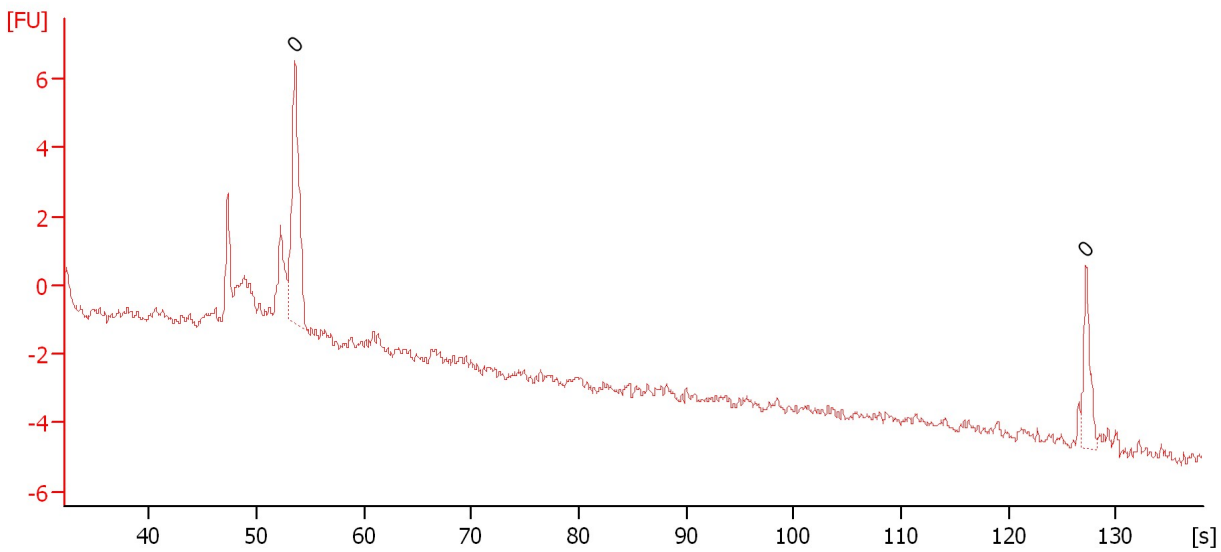
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		58.75

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary Continued ...

TPP ribo SHAPE seq 12X [MiSeq_262]



Overall Results for sample 3 : TPP ribo SHAPE seq 12X

Number of peaks found: 0 Noise: 0.2

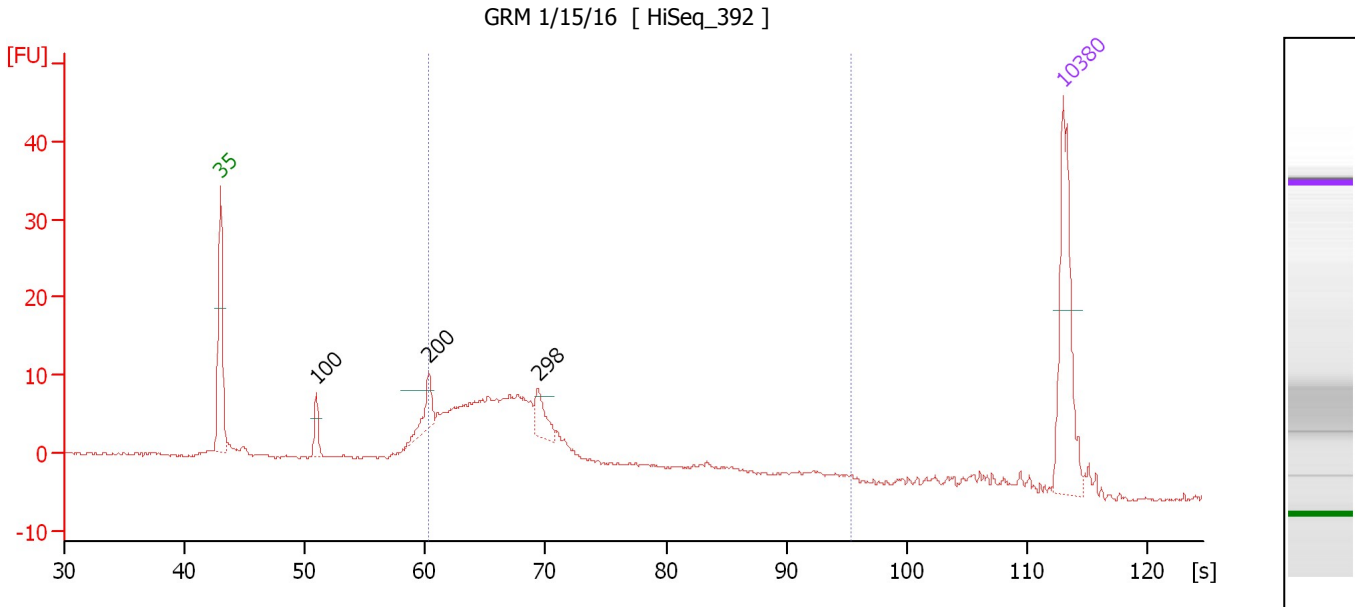
Peak table for sample 3 : TPP ribo SHAPE seq 12X

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		53.55
2	0	0.00	0.0		127.20

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 4 : GRM 1/15/16

Number of peaks found: 3 Corr. Area 1: 179.4
 Noise: 0.2

Peak table for sample 4 : GRM 1/15/16

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	100	20.54	312.5		50.94
3	200	28.08	213.2		60.31
4	298	23.74	120.6		69.37
5	10,380	75.00	10.9	Upper Marker	113.00

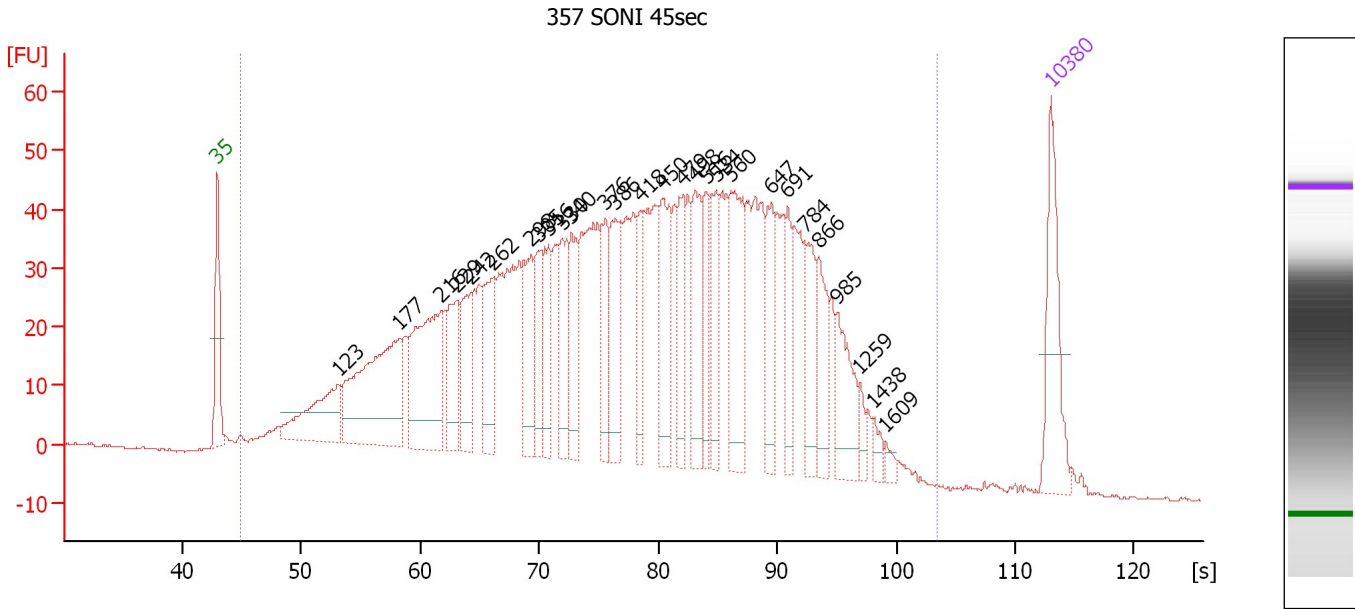
Region table for sample 4 : GRM 1/15/16

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	321	179.4	2,445.4	451.35	71	43.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : 357 SONI 45sec

Number of peaks found: 28 Corr. Area 1: 2,210.4
 Noise: 0.2

Peak table for sample 5 : 357 SONI 45sec

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	123	129.21	1,590.7		53.15
3	177	262.58	2,247.3		58.20
4	216	202.00	1,417.8		61.81
5	229	88.47	585.1		63.03
6	242	87.40	546.4		64.24
7	262	88.15	509.8		66.05
8	298	91.61	465.3		69.39
9	305	57.29	284.8		69.93
10	316	68.46	328.2		70.83
11	330	69.62	319.6		71.96
12	340	72.85	325.0		72.72
13	376	64.39	259.7		75.61
14	386	87.79	344.9		76.42
15	418	51.76	187.7		78.59
16	450	101.12	340.3		80.44
17	479	64.90	205.5		82.06
18	498	81.07	246.9		83.14
19	516	51.86	152.2		84.04
20	534	62.18	176.4		84.90
21	560	115.27	311.9		86.12
22	647	61.28	143.5		89.50
23	691	43.44	95.2		90.90
24	784	61.91	119.6		92.34
25	866	53.87	94.2		93.47

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 5 : 357 SONI 45sec

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	985	60.41	93.0		95.09
27	1,259	12.18	14.7		96.94
28	1,438	10.67	11.2		98.07
29	1,609	6.62	6.2		99.15
30	10,380	75.00	10.9	Upper Marker	113.00

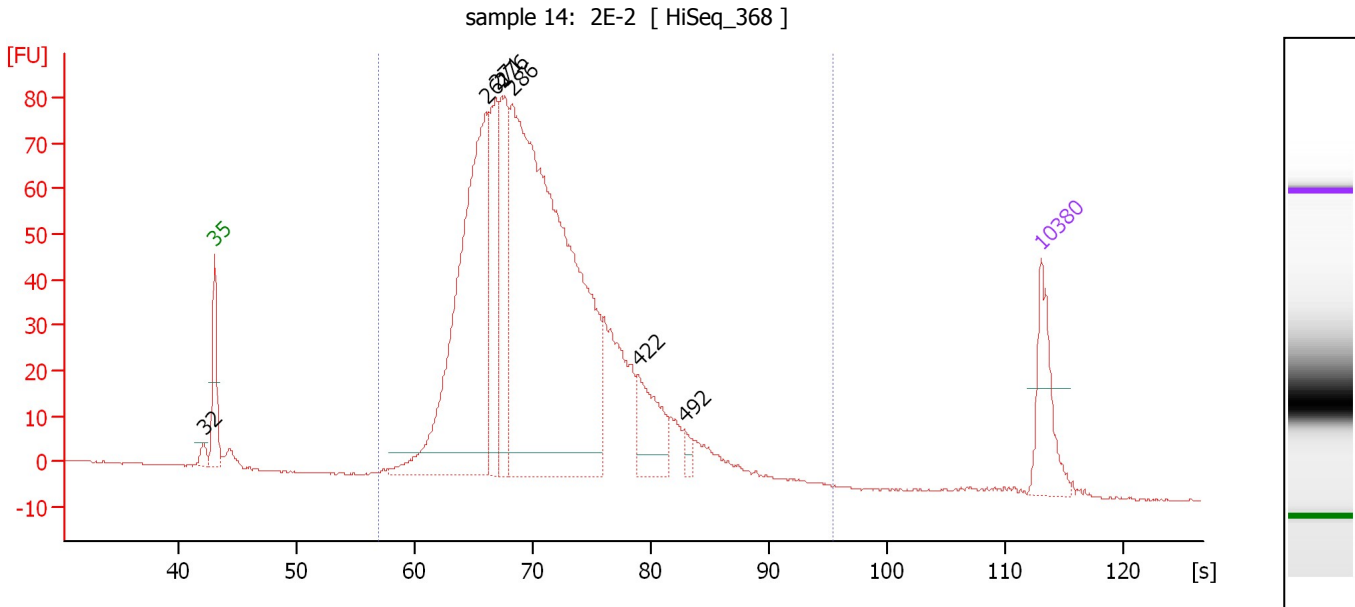
Region table for sample 5 : 357 SONI 45sec

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
47	2,563	470	2,210.4	20,750.3	3,802.76	99	58.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 14: 2E-2

Number of peaks found: 7 Corr. Area 1: 1,509.0
 Noise: 0.2

Peak table for sample 6 : sample 14: 2E-2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.04
2	35	125.00	5,411.3	Lower Marker	43.00
3	261	729.60	4,233.0		65.97
4	271	210.43	1,178.5		66.83
5	276	221.21	1,214.4		67.33
6	286	1,314.84	6,968.2		68.24
7	422	105.56	378.8		78.84
8	492	13.26	40.9		82.80
9	10,380	75.00	10.9	Upper Marker	113.00

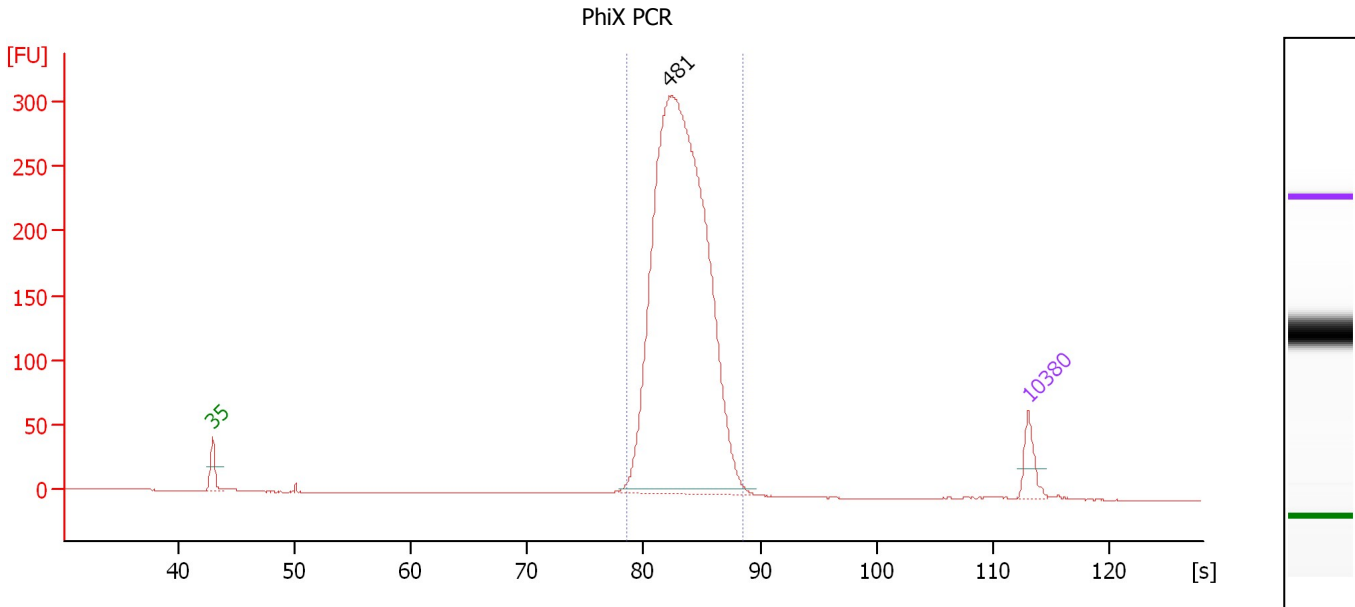
Region table for sample 6 : sample 14: 2E-2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
162	1,000	324	1,509.0	14,243.0	2,831.12	98	26.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : PhiX PCR

Number of peaks found: 1 Corr. Area 1: 2,067.7
 Noise: 0.2

Peak table for sample 7 : PhiX PCR

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	481	3,648.87	11,505.0		82.17
3	10,380	75.00	10.9	Upper Marker	113.00

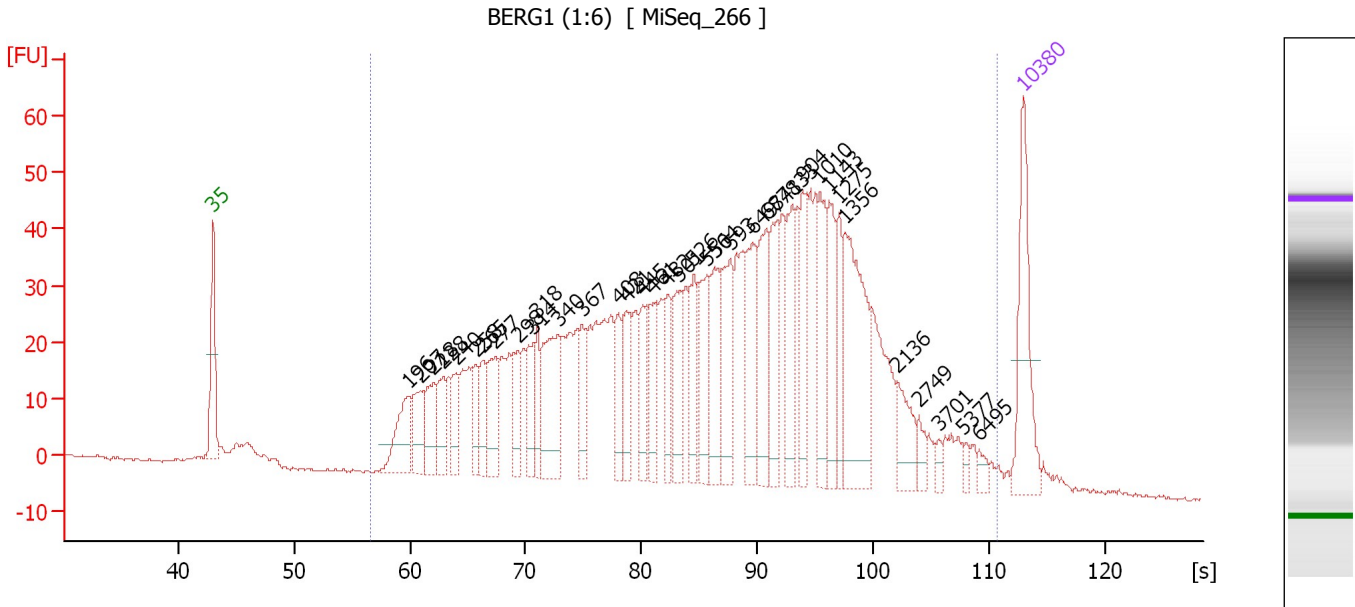
Region table for sample 7 : PhiX PCR

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
416	617	502	2,067.7	10,778.3	3,547.75	98	7.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
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Electropherogram Summary Continued ...



Overall Results for sample 8 : BERG1 (1:6)

Number of peaks found: 37 Corr. Area 1: 1,680.6
 Noise: 0.3

Peak table for sample 8 : BERG1 (1:6)

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	196	66.49	515.2		59.93
3	207	44.60	327.2		60.95
4	218	53.60	372.2		62.02
5	228	39.38	262.0		62.90
6	240	37.82	238.9		64.01
7	258	34.28	201.3		65.68
8	265	34.86	199.6		66.29
9	277	61.86	338.7		67.40
10	298	40.81	207.6		69.35
11	314	38.67	186.7		70.65
12	318	40.12	190.9		71.02
13	340	101.61	452.3		72.78
14	367	39.56	163.3		74.92
15	408	36.57	135.8		78.02
16	421	42.75	153.9		78.77
17	445	43.50	148.0		80.16
18	461	49.81	163.8		81.04
19	482	39.14	123.1		82.24
20	501	54.65	165.4		83.31
21	526	52.31	150.6		84.52
22	550	63.75	175.7		85.63
23	564	65.13	174.8		86.33
24	593	70.14	179.2		87.67
25	649	69.35	161.8		89.57

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
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Electropherogram Summary Continued ...

... Peak table for sample 8 : BERG1 (1:6)

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	687	76.49	168.6		90.78
27	741	64.20	131.2		91.75
28	833	60.11	109.4		93.01
29	904	57.56	96.5		93.98
30	1,010	57.10	85.6		95.37
31	1,143	54.10	71.7		96.21
32	1,275	38.38	45.6		97.04
33	1,356	121.66	136.0		97.55
34	2,136	32.90	23.3		102.05
35	2,749	11.70	6.4		104.00
36	3,701	6.23	2.6		105.67
37	5,377	5.91	1.7		107.76
38	6,495	7.68	1.8		109.15
39	10,380	75.00	10.9	Upper Marker	113.00

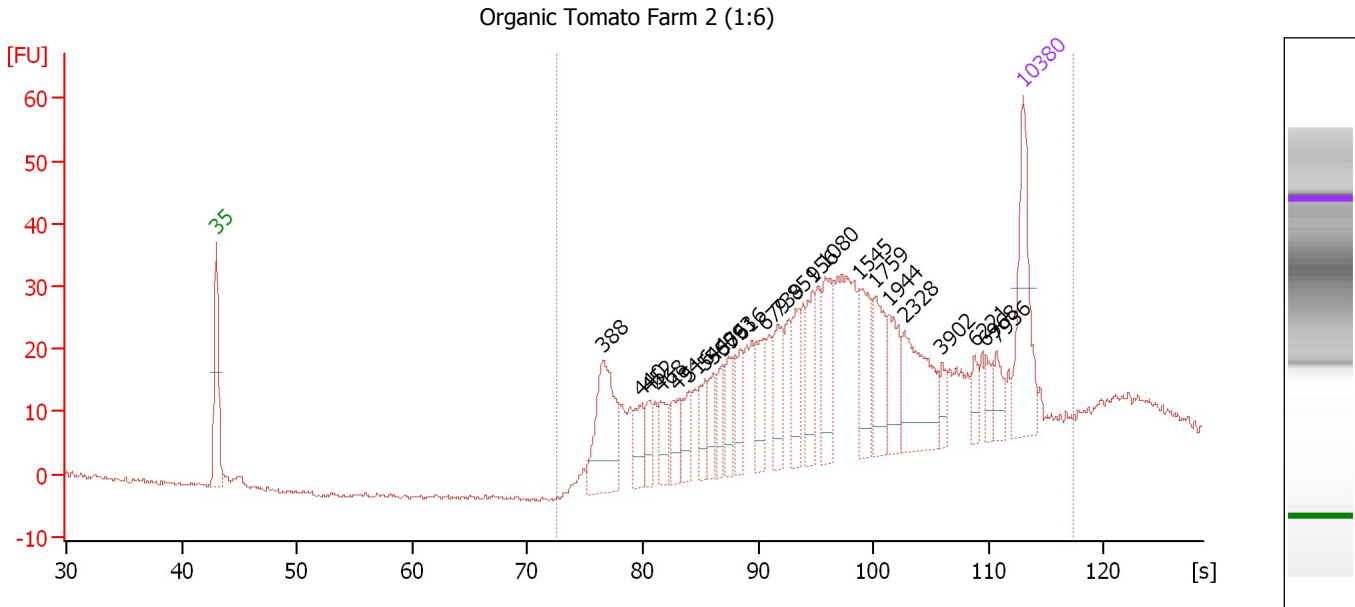
Region table for sample 8 : BERG1 (1:6)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
161	7,905	886	1,680.6	9,020.7	2,657.83	98	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : Organic Tomato Farm 2 (1:6)

Number of peaks found: 24 Corr. Area 1: 558.1
 Noise: 0.3

Peak table for sample 9 : Organic Tomato Farm 2 (1:6)

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	388	113.76	444.6		76.58
3	440	32.04	110.4		79.84
4	452	21.98	73.7		80.54
5	468	28.70	92.9		81.47
6	494	26.75	82.0		82.96
7	516	27.10	79.6		84.03
8	544	25.74	71.6		85.38
9	560	25.17	68.1		86.13
10	576	24.43	64.3		86.87
11	593	27.50	70.3		87.66
12	616	34.09	83.9		88.50
13	679	43.10	96.2		90.50
14	739	41.00	84.1		91.72
15	851	39.80	70.9		93.25
16	956	36.10	57.2		94.70
17	1,080	47.15	66.1		95.81
18	1,545	49.37	48.4		98.75
19	1,759	46.54	40.1		100.10
20	1,944	40.70	31.7		101.26
21	2,328	72.92	47.5		102.66
22	3,902	10.96	4.3		105.92
23	6,221	12.55	3.1		108.81
24	6,968	10.90	2.4		109.74
25	7,936	17.37	3.3		110.67

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
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Electropherogram Summary Continued ...**... Peak table for sample 9 : Organic Tomato Farm 2 (1:6)**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00

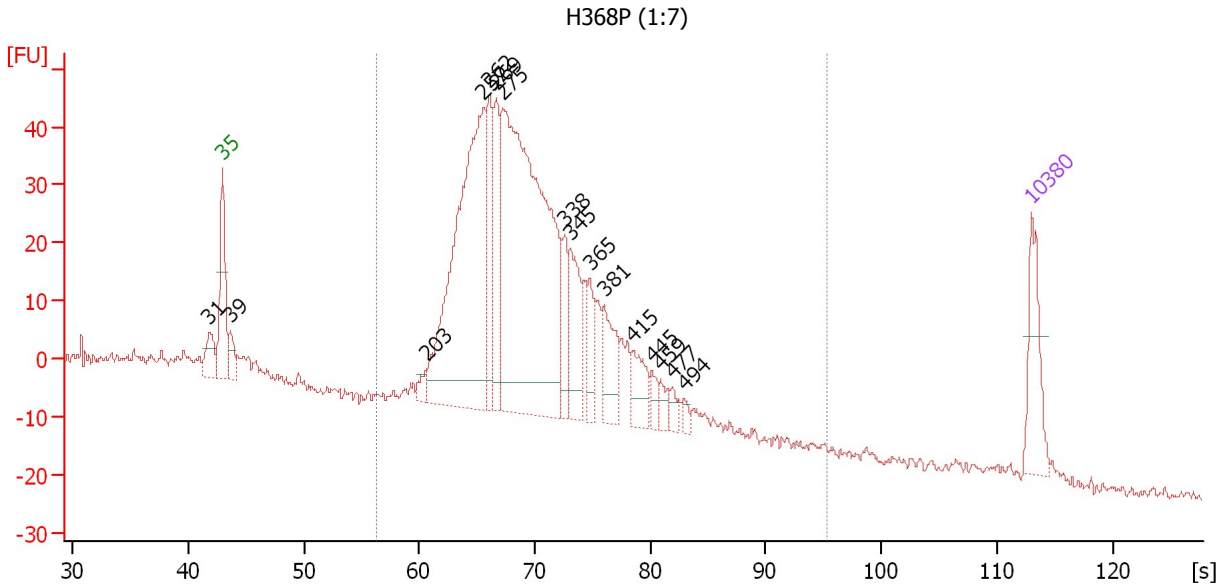
Region table for sample 9 : Organic Tomato Farm 2 (1:6)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
337	14,858	2,807	558.1	1,718.7	965.27	95	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : H368P (1:7)

Number of peaks found: 16 Corr. Area 1: 917.1
 Noise: 0.9

Peak table for sample 10 : H368P (1:7)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.79
2	35	125.00	5,411.3	Lower Marker	43.00
3	39	37.56	1,454.2		43.65
4	203	20.75	154.9		60.62
5	257	657.88	3,886.0		65.54
6	262	135.43	782.9		66.05
7	269	132.14	743.9		66.71
8	275	918.43	5,056.1		67.26
9	338	65.55	293.7		72.61
10	345	109.13	479.9		73.12
11	365	50.09	207.7		74.79
12	381	85.90	341.6		76.05
13	415	57.76	211.0		78.42
14	445	19.08	65.0		80.14
15	459	20.91	69.1		80.93
16	477	16.56	52.6		81.95
17	494	10.97	33.7		82.93
18	10,380	75.00	10.9	Upper Marker	113.00

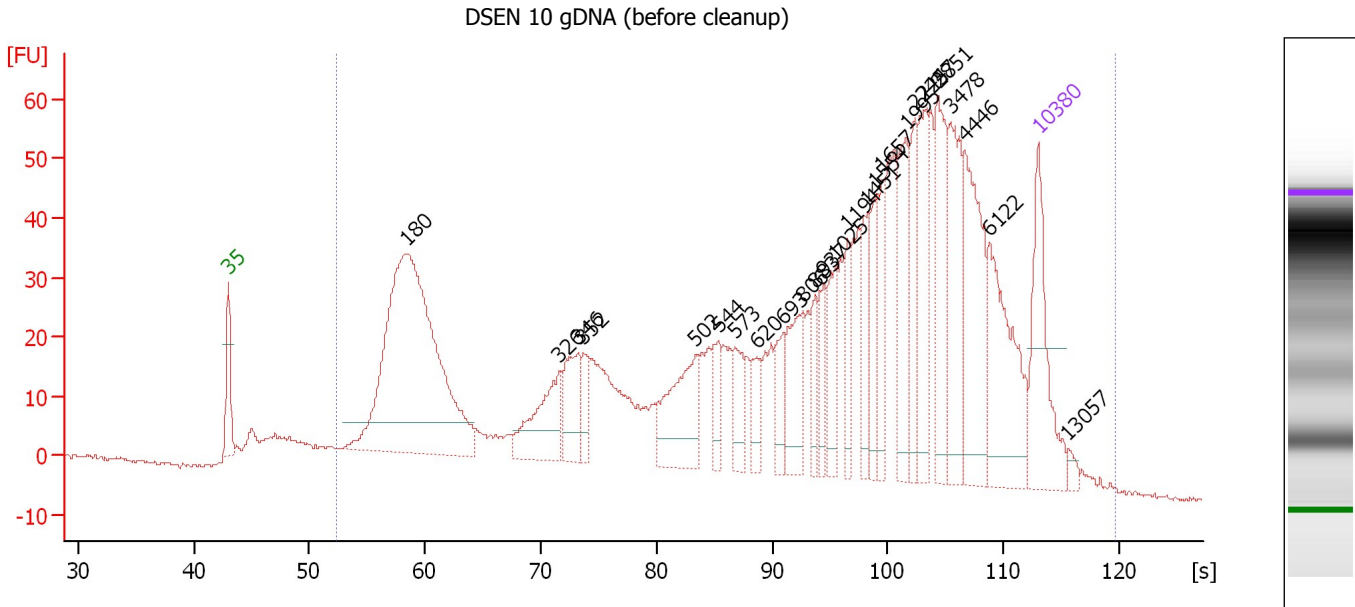
Region table for sample 10 : H368P (1:7)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
156	1,000	314	917.1	12,827.8	2,478.53	90	26.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : DSEN 10 gDNA (before cleanup)

Number of peaks found: 25 Corr. Area 1: 1,801.9
 Noise: 0.4

Peak table for sample 11 : DSEN 10 gDNA (before cleanup)

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	180	542.33	4,573.2		58.45
3	326	79.67	370.4		71.62
4	346	57.33	251.4		73.20
5	352	25.31	109.0		73.71
6	502	90.12	272.2		83.36
7	544	27.79	77.4		85.35
8	573	30.94	81.8		86.74
9	620	23.28	56.9		88.65
10	693	29.32	64.1		90.97
11	806	57.44	108.0		92.64
12	883	23.28	39.9		93.70
13	937	26.08	42.2		94.44
14	1,025	38.69	57.2		95.47
15	1,194	27.96	35.5		96.53
16	1,451	39.25	41.0		98.16
17	1,554	33.67	32.8		98.81
18	1,657	32.37	29.6		99.45
19	1,995	58.93	44.8		101.59
20	2,224	42.14	28.7		102.33
21	2,457	60.99	37.6		103.07
22	2,851	72.09	38.3		104.33
23	3,478	75.70	33.0		105.39
24	4,446	101.22	34.5		106.60
25	6,122	99.89	24.7		108.69

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 11 : DSEN 10 gDNA (before cleanup)

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00
27	13,057	0.00	0.0		115.55

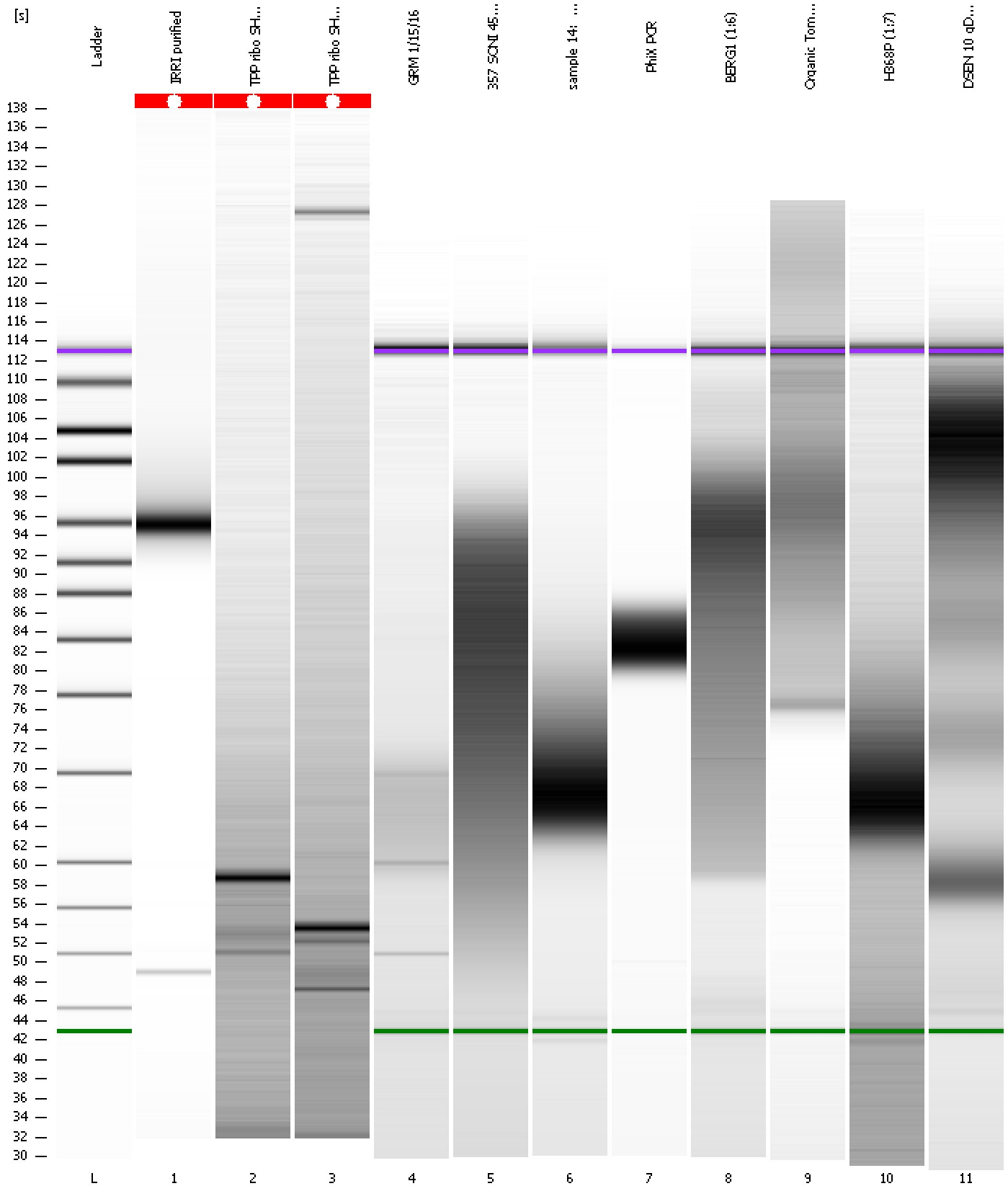
Region table for sample 11 : DSEN 10 gDNA (before cleanup)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
115	17,330	2,471	1,801.9	9,036.1	2,428.39	96	100.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
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Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad

Created: 1/27/2016 3:05:35 PM
 Modified: 1/27/2016 3:47:39 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		1/27/2016 3:46:54 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2016-01-27\2016-01-27_003.xad)		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/27/2016 3:05:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1