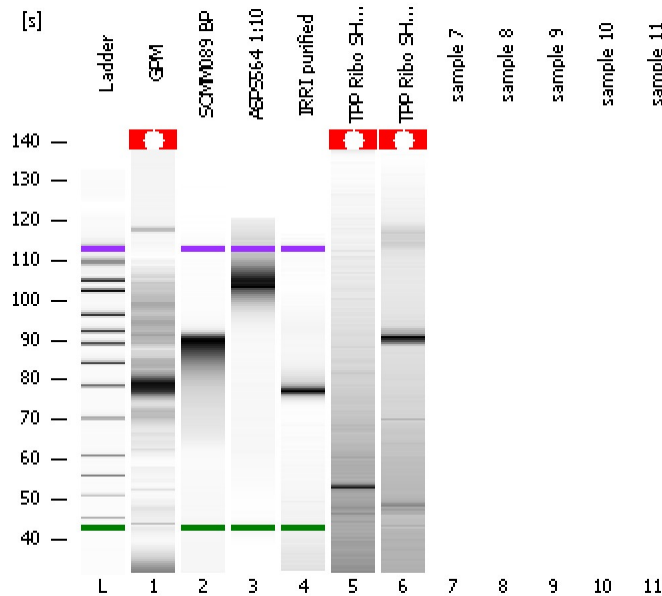


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 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

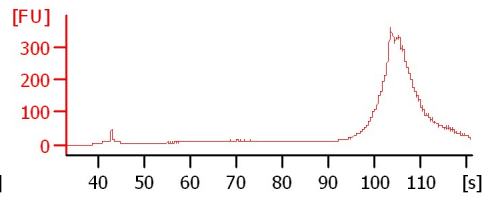
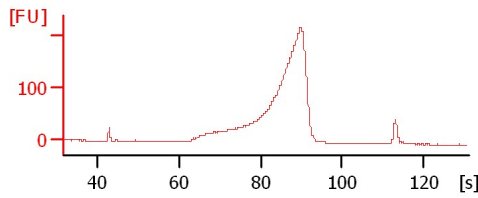
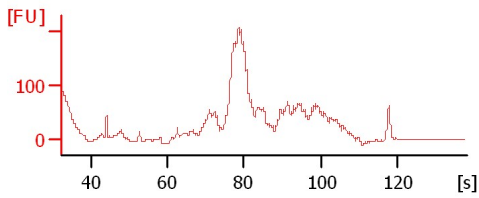
Chip Information:

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

GPM

SOMM089 BP

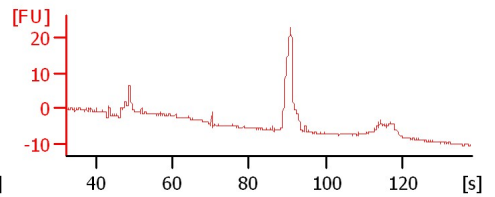
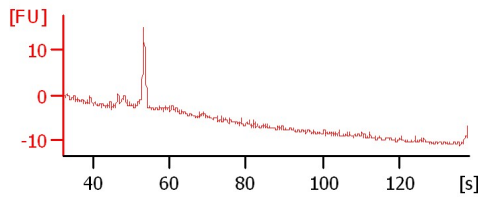
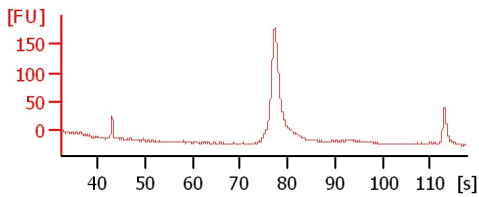
ASP564 1:10



IIRI purified

TPP Ribo SHAPE Seq 9X

TPP Ribo SHAPE Seq 12X



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
GPM		<input type="checkbox"/>	✓			
SOMM089 BP		<input type="checkbox"/>	✓			
ASPS564 1:10		<input type="checkbox"/>	✓			
IRRI purified		<input type="checkbox"/>	✓			
TPP Ribo SHAPE Seq 9X		<input type="checkbox"/>	✓			
TPP Ribo SHAPE Seq 12X		<input type="checkbox"/>	✓			
sample 7		<input type="checkbox"/>				
sample 8		<input type="checkbox"/>				
sample 9		<input type="checkbox"/>				
sample 10		<input type="checkbox"/>				
sample 11		<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\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 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

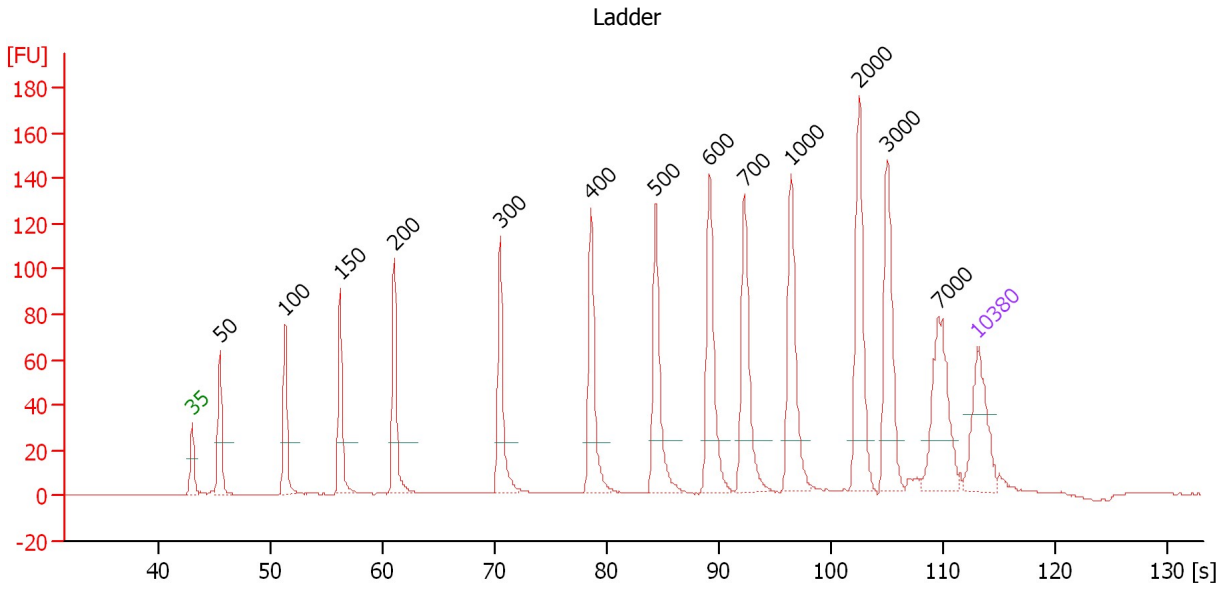
Ladder

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\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

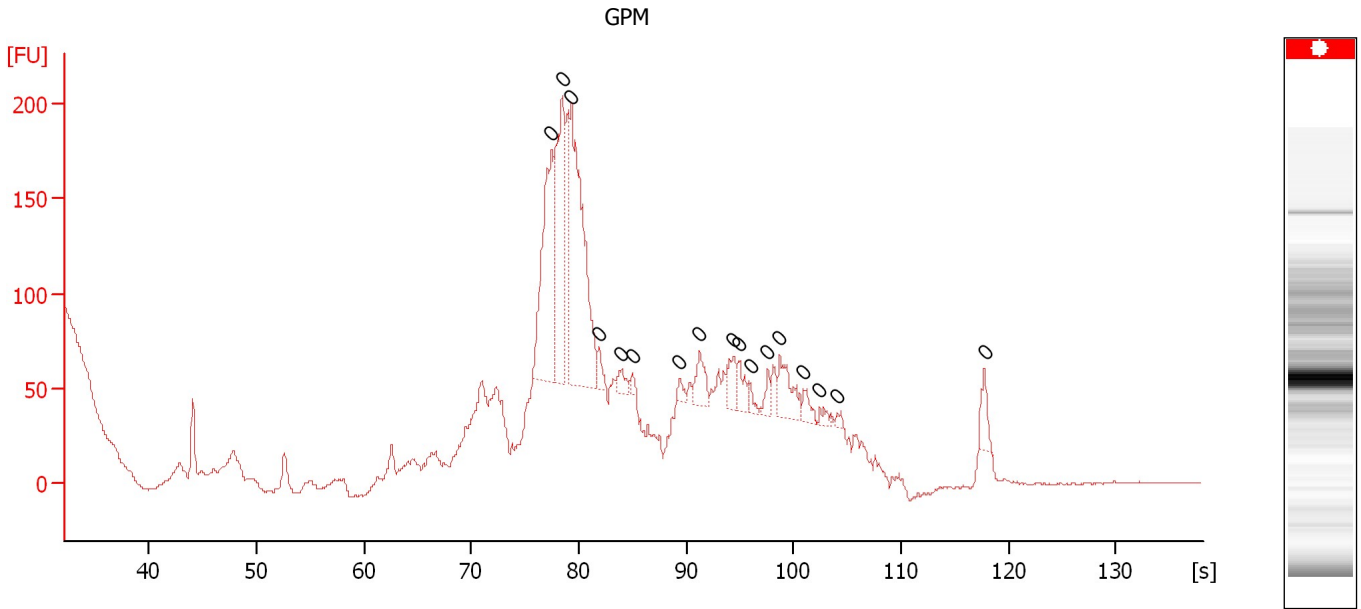
Peak table for Ladder

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	45.49
3	100	150.00	2,272.7	Ladder Peak	51.28
4	150	150.00	1,515.2	Ladder Peak	56.21
5	200	150.00	1,136.4	Ladder Peak	61.05
6	300	150.00	757.6	Ladder Peak	70.48
7	400	150.00	568.2	Ladder Peak	78.57
8	500	150.00	454.5	Ladder Peak	84.32
9	600	150.00	378.8	Ladder Peak	89.16
10	700	150.00	324.7	Ladder Peak	92.27
11	1,000	150.00	227.3	Ladder Peak	96.39
12	2,000	150.00	113.6	Ladder Peak	102.47
13	3,000	150.00	75.8	Ladder Peak	104.96
14	7,000	150.00	32.5	Ladder Peak	109.50
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : GPM

Number of peaks found: 0 Noise: 0.6

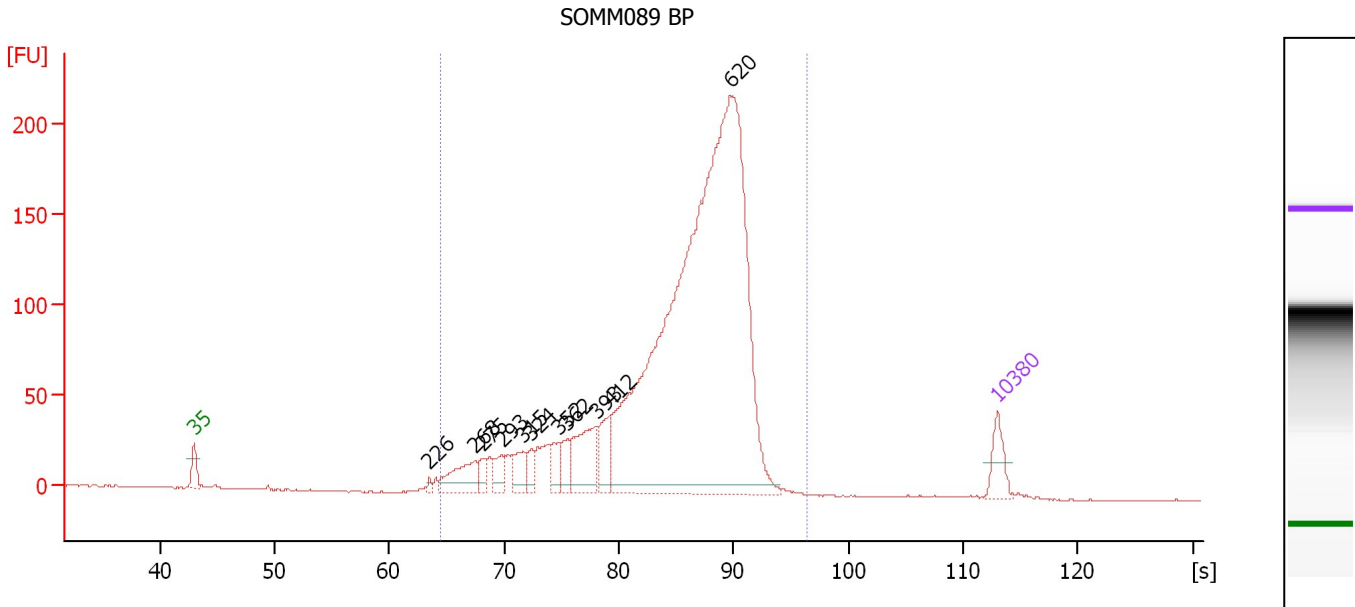
Peak table for sample 1 : GPM

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		77.45
2	0	0.00	0.0		78.50
3	0	0.00	0.0		79.25
4	0	0.00	0.0		81.80
5	0	0.00	0.0		83.85
6	0	0.00	0.0		85.00
7	0	0.00	0.0		89.40
8	0	0.00	0.0		91.25
9	0	0.00	0.0		94.40
10	0	0.00	0.0		94.85
11	0	0.00	0.0		95.95
12	0	0.00	0.0		97.60
13	0	0.00	0.0		98.70
14	0	0.00	0.0		100.90
15	0	0.00	0.0		102.45
16	0	0.00	0.0		104.10
17	0	0.00	0.0		117.75

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : SOMM089 BP

Number of peaks found: 11 Corr. Area 1: 2,326.4
 Noise: 0.3

Peak table for sample 2 : SOMM089 BP

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	226	14.49	97.2		63.49
3	268	175.96	994.5		67.47
4	275	43.15	238.2		68.08
5	293	74.99	388.0		69.81
6	315	96.49	464.3		71.68
7	324	56.58	265.0		72.39
8	352	70.22	301.9		74.72
9	362	68.69	287.3		75.52
10	393	228.36	880.6		78.00
11	412	135.71	499.1		79.26
12	620	3,606.97	8,808.7		89.79
13	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 2 : SOMM089 BP

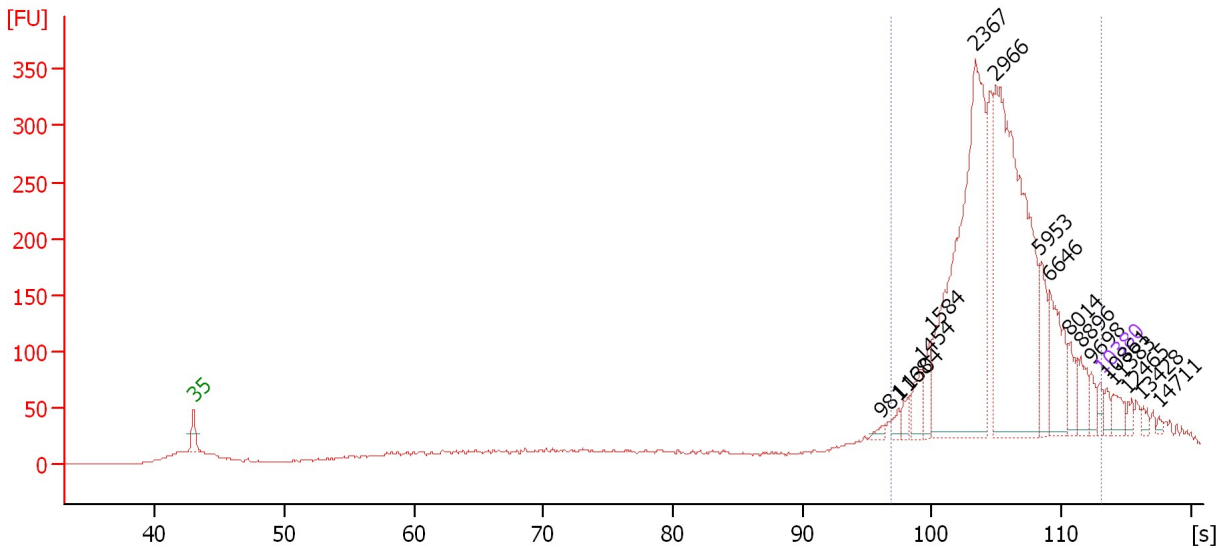
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
236	1,000	524	2,326.4	16,015.0	5,048.16	99	21.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...

ASPS564 1:10



Overall Results for sample 3 : ASPS564 1:10

Number of peaks found: 17 Corr. Area 1: 2,497.4
 Noise: 0.2

Peak table for sample 3 : ASPS564 1:10

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	981	43.58	67.3		96.12
3	1,168	68.72	89.1		97.41
4	1,304	98.52	114.4		98.24
5	1,454	212.28	221.1		99.15
6	1,584	203.86	195.0		99.94
7	2,367	3,121.61	1,998.6		103.38
8	2,966	3,125.17	1,596.4		104.87
9	5,953	357.96	91.1		108.31
10	6,646	566.32	129.1		109.10
11	8,014	201.61	38.1		110.55
12	8,896	189.58	32.3		111.47
13	9,698	93.76	14.6		112.30
14	10,380	75.00	10.9	Upper Marker	113.00
15	10,861	0.00	0.0		113.50
16	11,383	0.00	0.0		114.04
17	12,465	0.00	0.0		115.16
18	13,428	0.00	0.0		116.15
19	14,711	0.00	0.0		117.48

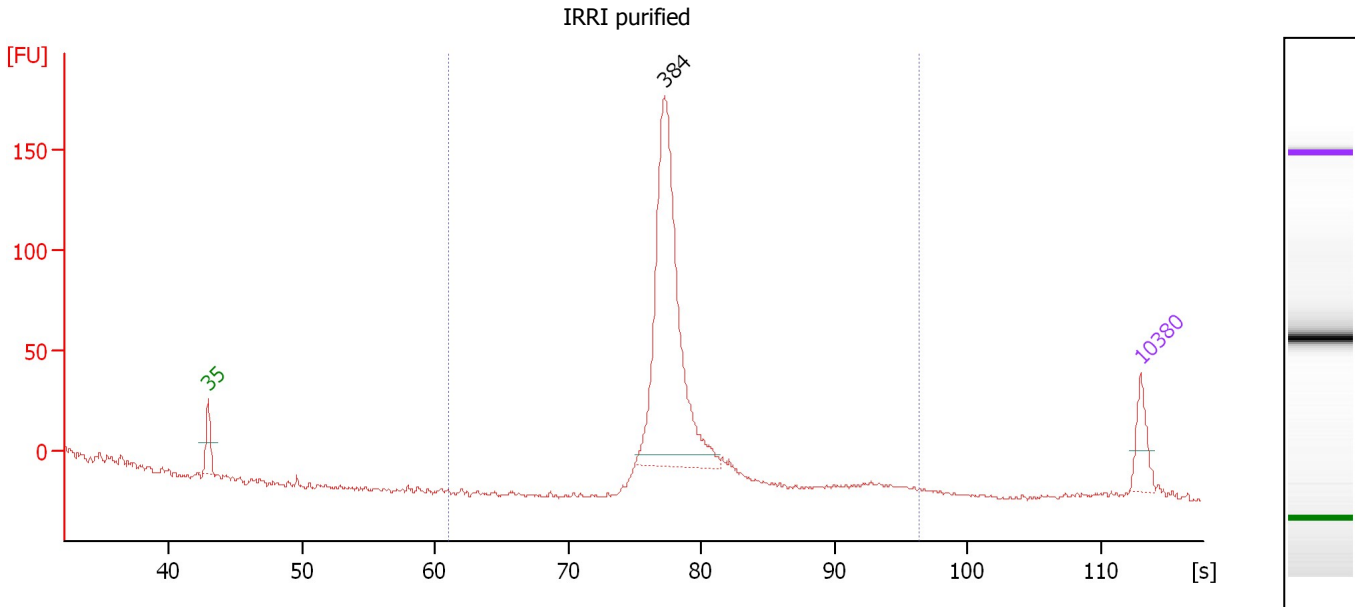
Region table for sample 3 : ASPS564 1:10

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
1,085	10,468	3,812	2,497.4	4,698.4	9,022.27	90	56.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : IRRI purified

Number of peaks found: 1 Corr. Area 1: 593.9
 Noise: 1.0

Peak table for sample 4 : IRRI purified

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	384	1,152.36	4,547.7		77.27
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 4 : IRRI purified

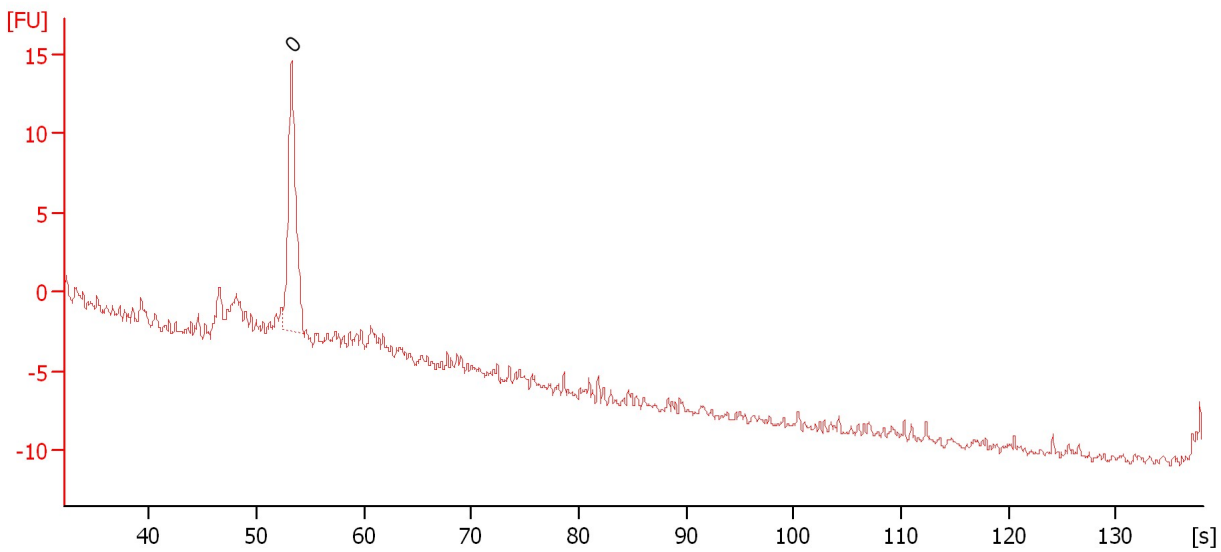
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	397	593.9	5,125.3	1,329.49	98	11.7

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...

TPP Ribo SHAPE Seq 9X



Overall Results for sample 5 : TPP Ribo SHAPE Seq 9X

Number of peaks found: 0 Noise: 0.4

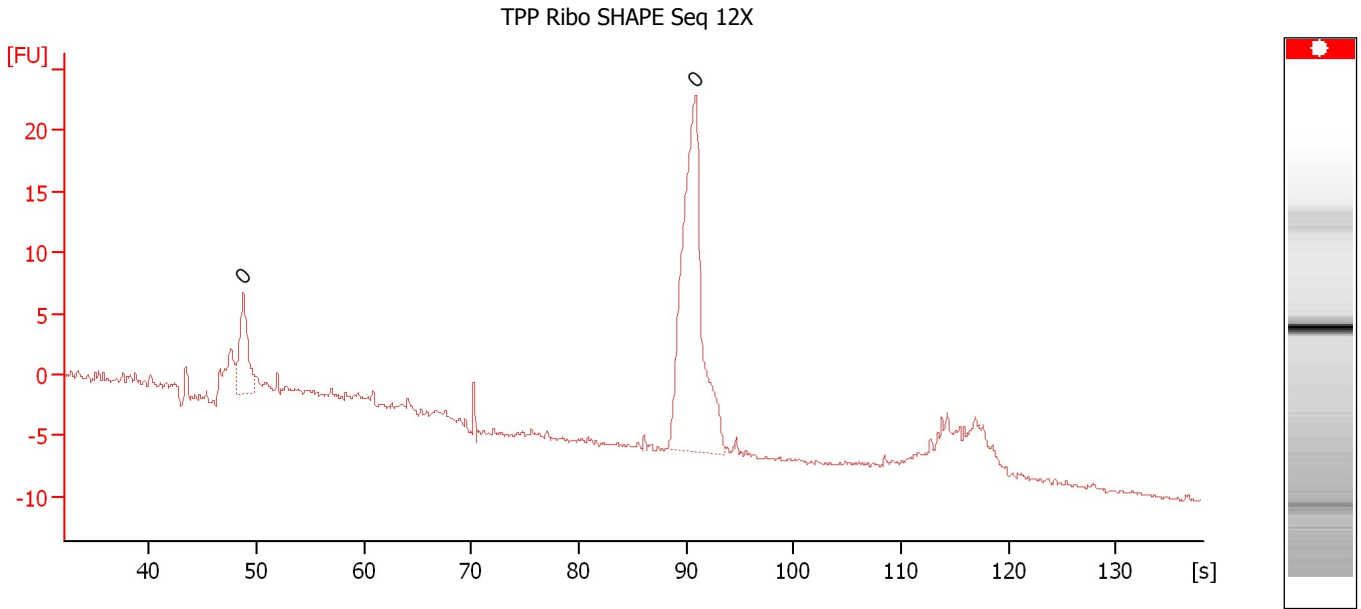
Peak table for sample 5 : TPP Ribo SHAPE Seq 9X

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		53.25

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : TPP Ribo SHAPE Seq 12X

Number of peaks found: 0 Noise: 0.2

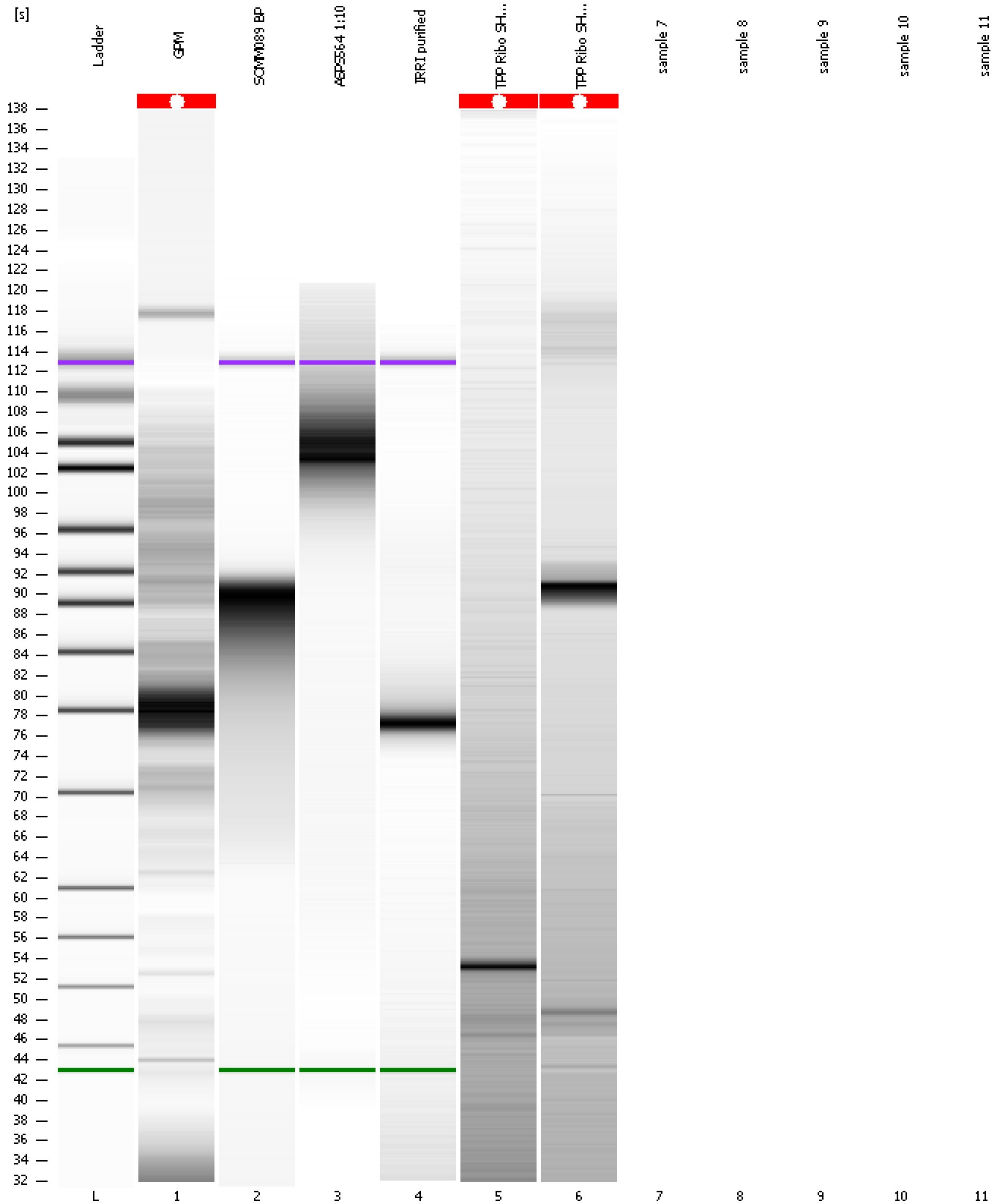
Peak table for sample 6 : TPP Ribo SHAPE Seq 12X

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		48.75
2	0	0.00	0.0		90.85

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
Modified: 1/22/2016 2:14:51 PM

Invalid Samples

Sample 7 has not been run, no results available.

Sample 8 has not been run, no results available.

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad

Created: 1/22/2016 1:24:16 PM
 Modified: 1/22/2016 2:14:51 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 7)		Instrument	Run		1/22/2016 1:51:16 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2016-01-22\2016-01-22_002.xad)		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/22/2016 1:24:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1