

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...alyzer\2016-01-14\2016-01-14_003_HiSeq268_Libraries_1-5.xad

Created: 1/14/2016 3:08:47 PM
Modified: 1/14/2016 4:56:22 PM

Electrophoresis File Run Summary

Instrument Information:

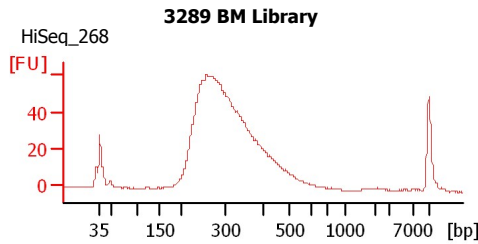
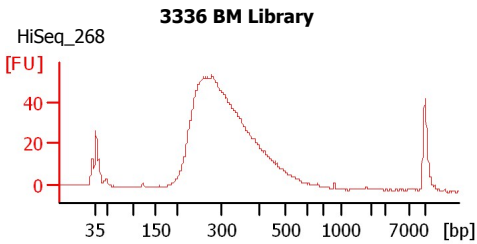
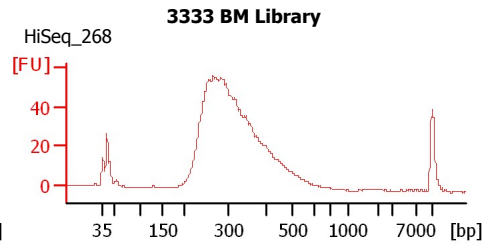
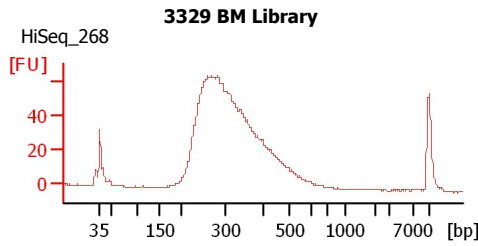
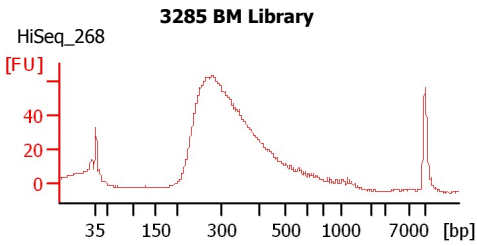
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
3285 BM Library	HiSeq_268	<input type="checkbox"/>	✓			
3329 BM Library	HiSeq_268	<input type="checkbox"/>	✓			
3333 BM Library	HiSeq_268	<input type="checkbox"/>	✓			
3336 BM Library	HiSeq_268	<input type="checkbox"/>	✓			
3289 BM Library	HiSeq_268	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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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

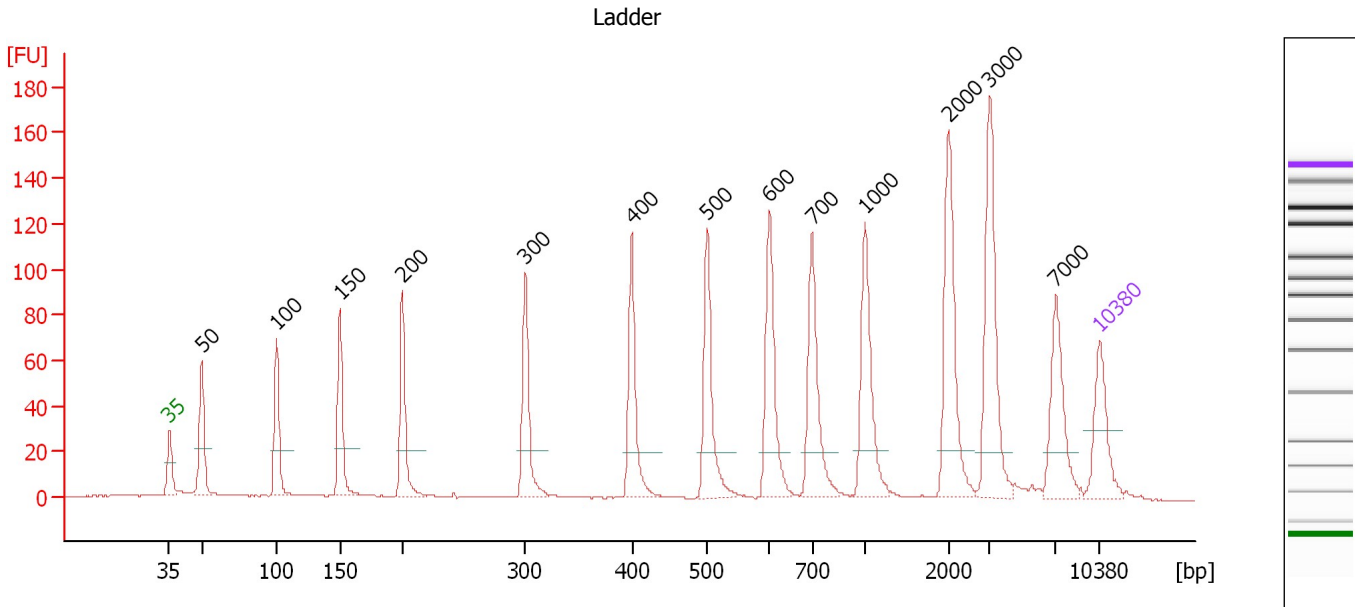
Ladder

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

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Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

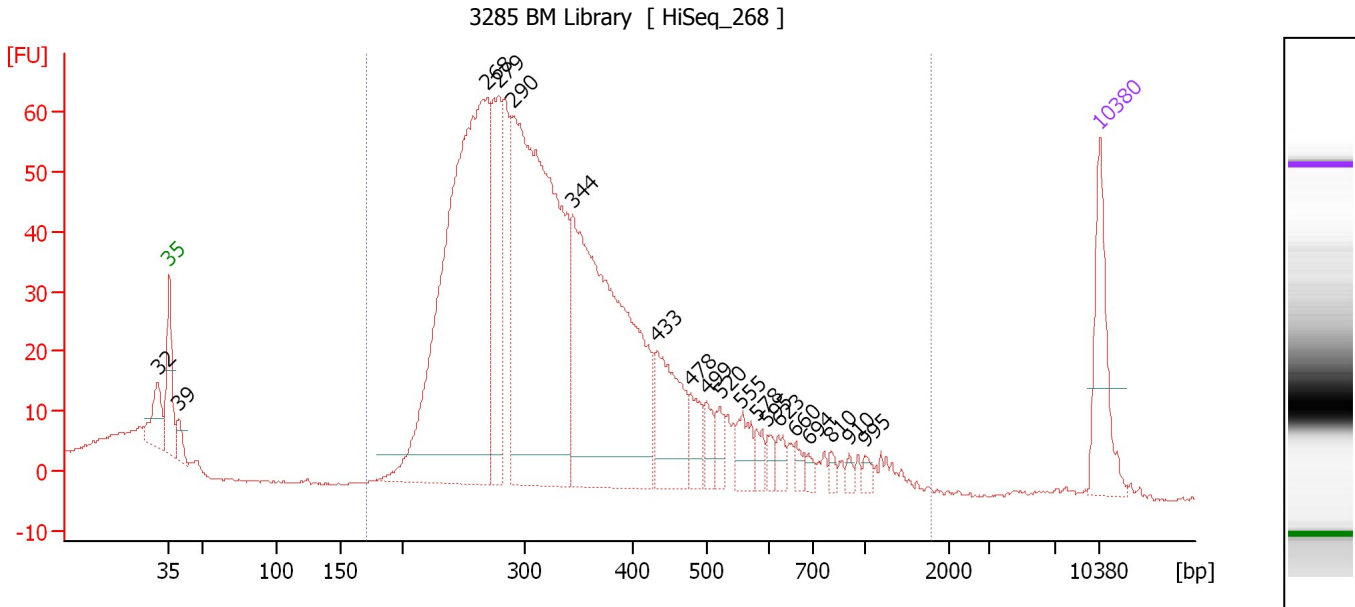
Peak table for Ladder

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	45.43
3	100	150.00	2,272.7	Ladder Peak	51.06
4	150	150.00	1,515.2	Ladder Peak	55.84
5	200	150.00	1,136.4	Ladder Peak	60.51
6	300	150.00	757.6	Ladder Peak	69.77
7	400	150.00	568.2	Ladder Peak	77.79
8	500	150.00	454.5	Ladder Peak	83.46
9	600	150.00	378.8	Ladder Peak	88.14
10	700	150.00	324.7	Ladder Peak	91.34
11	1,000	150.00	227.3	Ladder Peak	95.34
12	2,000	150.00	113.6	Ladder Peak	101.60
13	3,000	150.00	75.8	Ladder Peak	104.70
14	7,000	150.00	32.5	Ladder Peak	109.66
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 7 : 3285 BM Library

Number of peaks found: 19 Corr. Area 1: 1,300.0
 Noise: 0.2

Peak table for sample 7 : 3285 BM Library

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.11
2	35	125.00	5,411.3	Lower Marker	43.00
3	39	24.69	952.8		43.69
4	268	858.36	4,845.6		66.84
5	279	177.03	963.0		67.78
6	290	754.67	3,939.3		68.87
7	344	578.73	2,547.7		73.31
8	433	120.70	422.0		79.68
9	478	36.01	114.2		82.20
10	499	24.76	75.3		83.38
11	520	20.08	58.5		84.42
12	555	34.91	95.3		86.05
13	578	15.27	40.0		87.13
14	595	12.45	31.7		87.92
15	623	15.75	38.3		88.86
16	660	11.62	26.7		90.05
17	694	8.93	19.5		91.13
18	810	7.53	14.1		92.81
19	910	6.56	10.9		94.14
20	995	7.78	11.8		95.28
21	10,380	75.00	10.9	Upper Marker	113.00

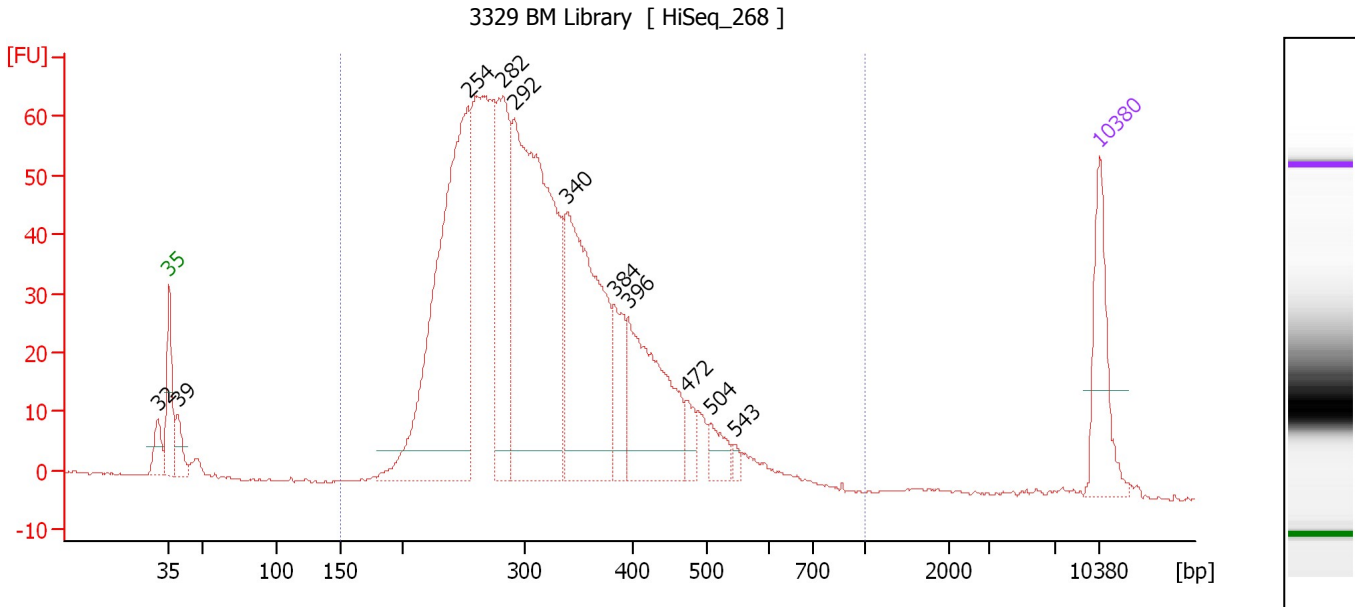
Region table for sample 7 : 3285 BM Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
171	1,786	359	1,300.0	13,582.3	2,800.44	89	43.5

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Electropherogram Summary Continued ...



Overall Results for sample 8 : 3329 BM Library

Number of peaks found: 11 Corr. Area 1: 1,258.3
 Noise: 0.2

Peak table for sample 8 : 3329 BM Library

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.11
2	35	125.00	5,411.3	Lower Marker	43.00
3	39	37.90	1,473.9		43.64
4	254	595.08	3,554.8		65.48
5	282	236.41	1,270.6		68.10
6	292	605.44	3,146.7		68.98
7	340	354.87	1,583.7		72.94
8	384	66.81	263.7		76.49
9	396	202.40	774.0		77.48
10	472	21.76	69.8		81.88
11	504	28.55	85.8		83.66
12	543	6.93	19.3		85.48
13	10,380	75.00	10.9	Upper Marker	113.00

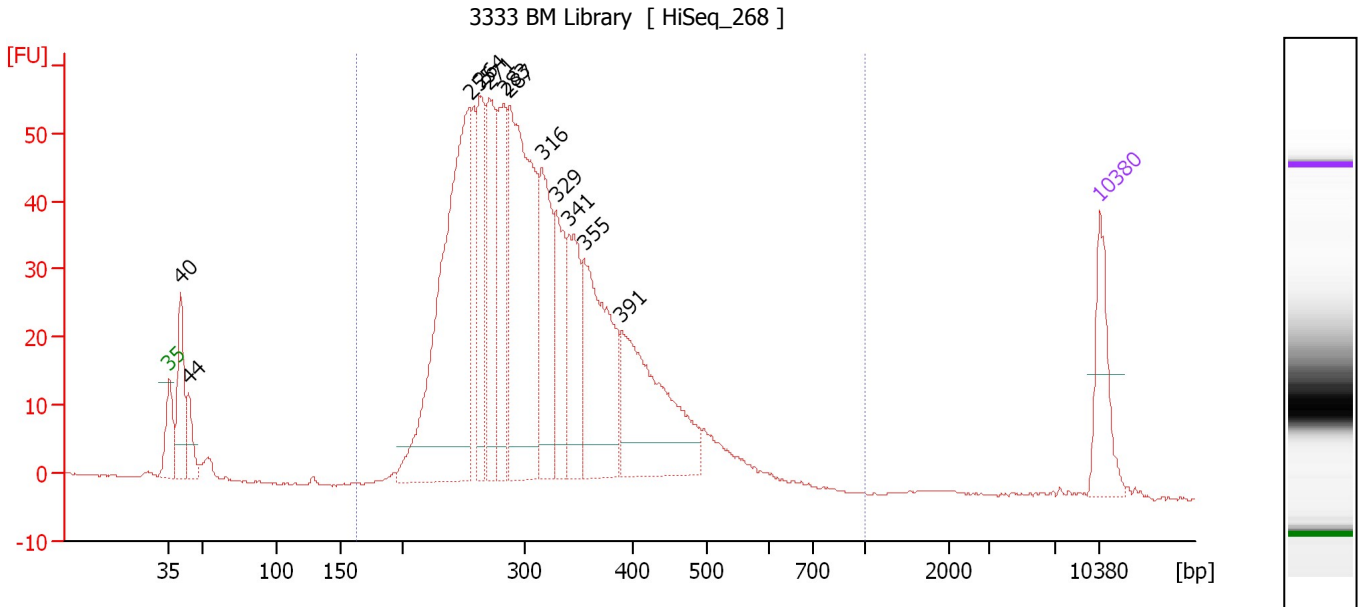
Region table for sample 8 : 3329 BM Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
150	1,000	323	1,258.3	12,868.1	2,547.25	97	25.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...alyzer\2016-01-14\2016-01-14_003_HiSeq268_Libraries_1-5.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : 3333 BM Library

Number of peaks found: 12 Corr. Area 1: 1,046.5
 Noise: 0.2

Peak table for sample 9 : 3333 BM Library

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	40	134.66	5,087.7		43.83
3	44	56.08	1,944.3		44.41
4	255	664.72	3,952.3		65.59
5	264	162.99	936.3		66.41
6	271	208.05	1,164.9		67.05
7	283	188.18	1,008.6		68.17
8	287	493.65	2,607.0		68.56
9	316	195.07	935.8		71.04
10	329	128.44	592.3		72.06
11	341	147.84	656.2		73.08
12	355	266.91	1,138.2		74.20
13	391	278.76	1,081.5		77.03
14	10,380	75.00	10.9	Upper Marker	113.00

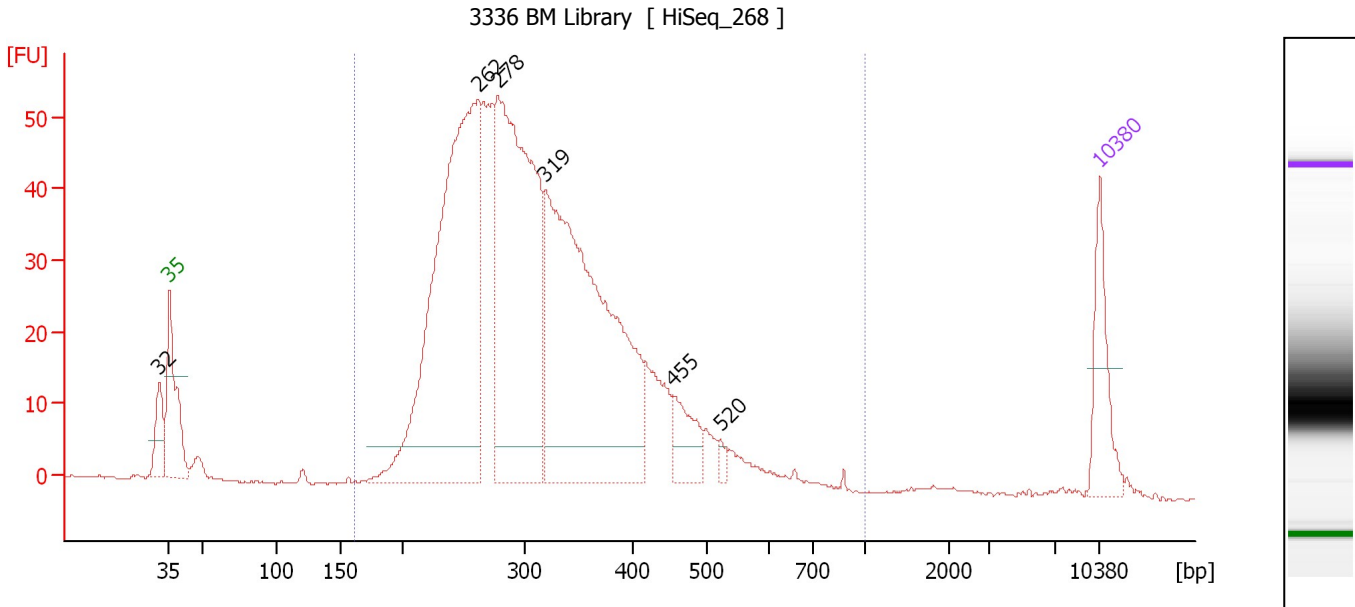
Region table for sample 9 : 3333 BM Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
164	1,000	321	1,046.5	15,425.2	3,054.19	95	24.2

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Electropherogram Summary Continued ...



Overall Results for sample 10 : 3336 BM Library

Number of peaks found: 6 Corr. Area 1: 1,027.9
 Noise: 0.1

Peak table for sample 10 : 3336 BM Library

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.22
2	35	125.00	5,411.3	Lower Marker	43.00
3	262	907.22	5,249.9		66.24
4	278	750.24	4,093.1		67.71
5	319	807.87	3,838.3		71.28
6	455	67.82	226.0		80.89
7	520	10.10	29.4		84.42
8	10,380	75.00	10.9	Upper Marker	113.00

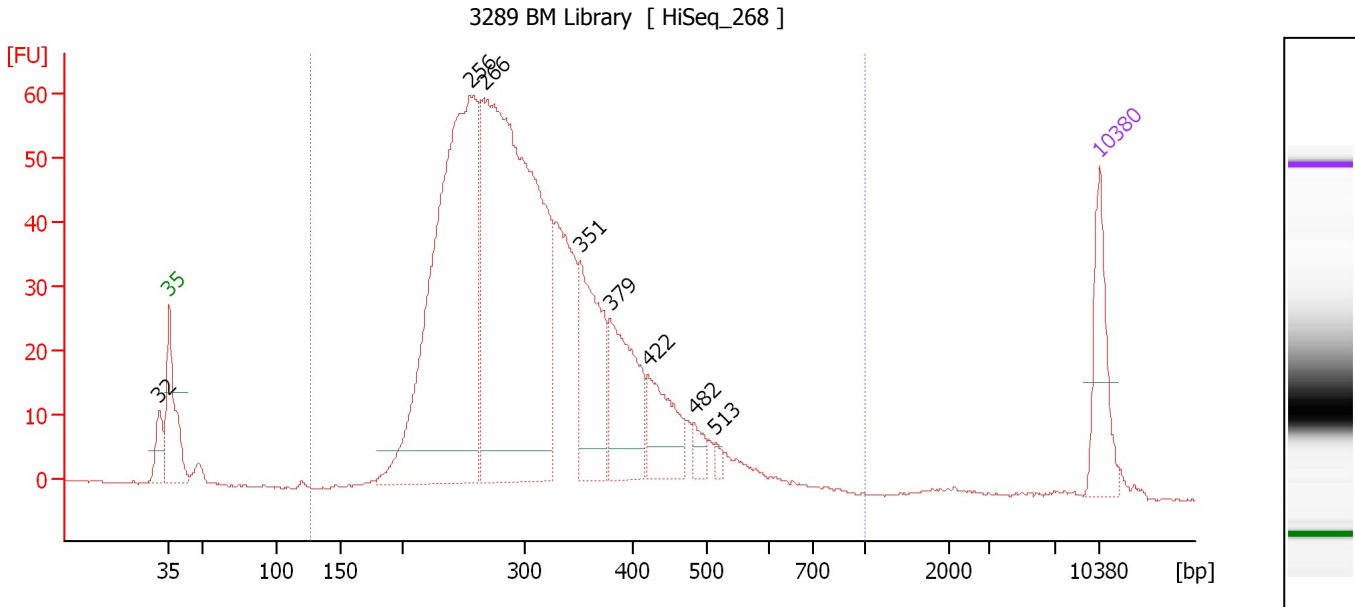
Region table for sample 10 : 3336 BM Library

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	319	1,027.9	14,870.2	2,906.98	97	26.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...alyzer\2016-01-14\2016-01-14_003_HiSeq268_Libraries_1-5.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : 3289 BM Library

Number of peaks found: 8 Corr. Area 1: 1,135.0
 Noise: 0.2

Peak table for sample 11 : 3289 BM Library

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.22
2	35	125.00	5,411.3	Lower Marker	43.00
3	256	879.75	5,213.8		65.66
4	266	1,065.18	6,062.0		66.64
5	351	195.59	843.4		73.89
6	379	169.44	676.5		76.14
7	422	98.14	352.5		79.03
8	482	22.30	70.1		82.45
9	513	7.73	22.8		84.07
10	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : 3289 BM Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
127	1,000	315	1,135.0	14,438.6	2,788.20	98	25.6

Assay Class: High Sensitivity DNA Assay
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Gel Image

Assay Class: High Sensitivity DNA Assay
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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		1/14/2016 3:50:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyser\2016-01-14\2016-01-14_003.xad)		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/14/2016 3:08:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1