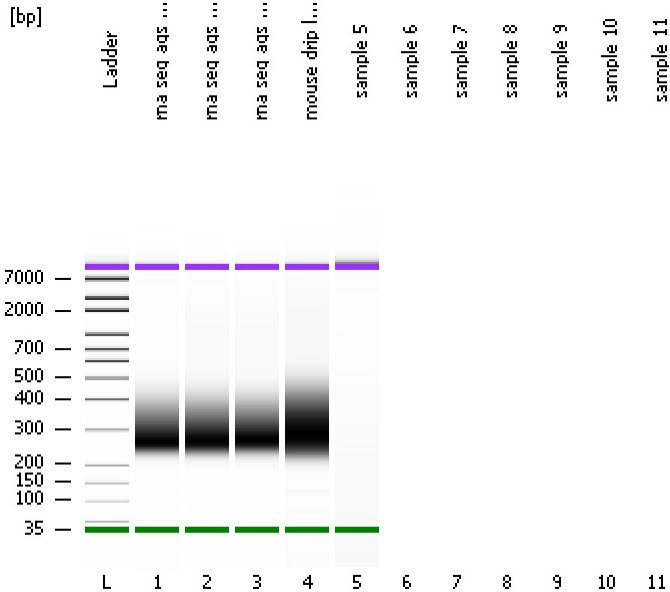


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
Modified: 9/4/2013 3:32:42 PM

Electrophoresis File Run Summary



Instrument Information:

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

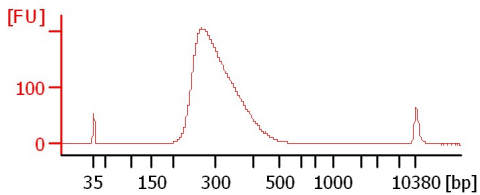
Assay Information:

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

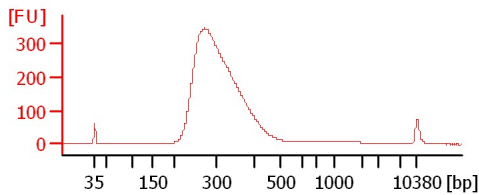
Chip Information:

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

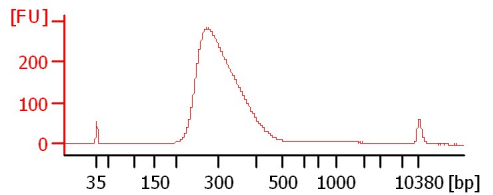
rna seq ags control i20



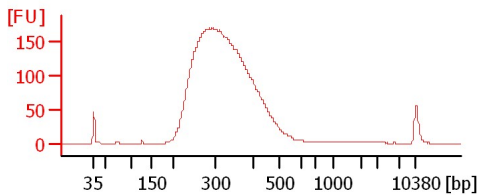
rna seq ags h2a i21



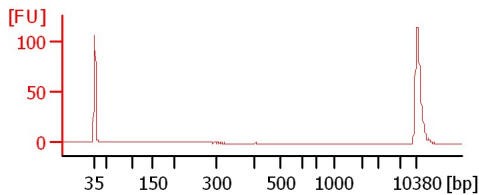
rna seq ags h2b i22



mouse drip lib i23



sample 5



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
Modified: 9/4/2013 3:32:42 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
rna seq ags control i20		<input type="checkbox"/>	✓			
rna seq ags h2a i21		<input type="checkbox"/>	✓			
rna seq ags h2b i22		<input type="checkbox"/>	✓			
mouse drip lib i23		<input type="checkbox"/>	✓			
sample 5		<input type="checkbox"/>	✓			
sample 6		<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\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
Modified: 9/4/2013 3:32:42 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

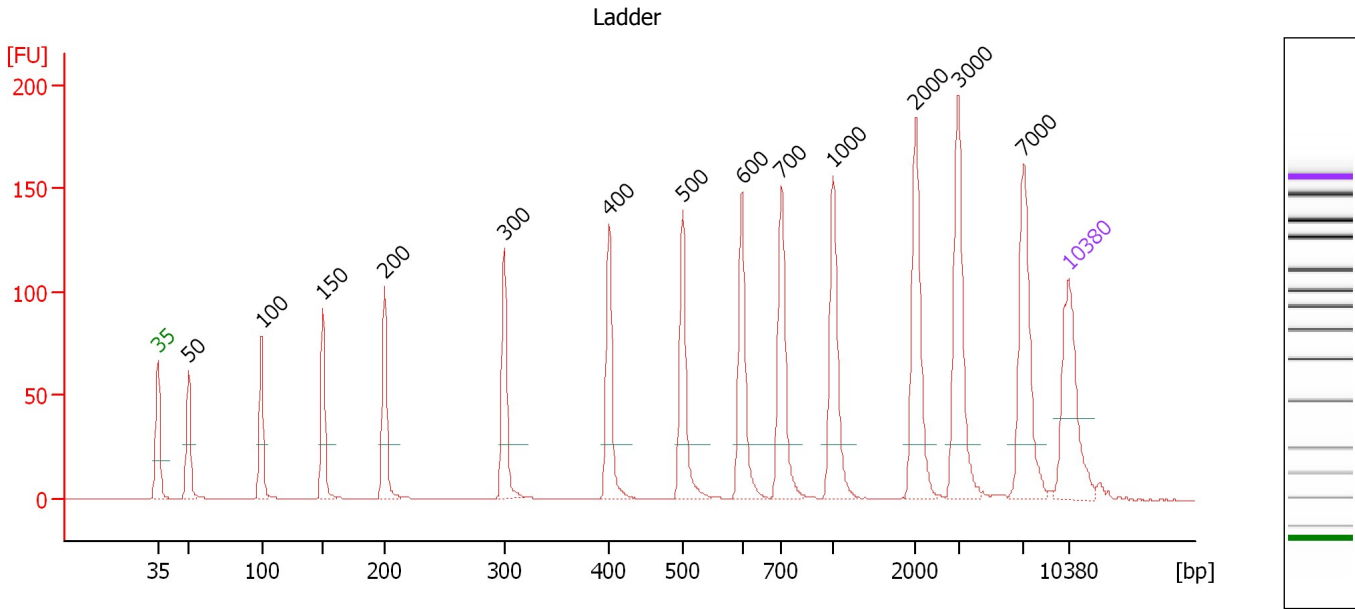
Ladder

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\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
 Modified: 9/4/2013 3:32:42 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

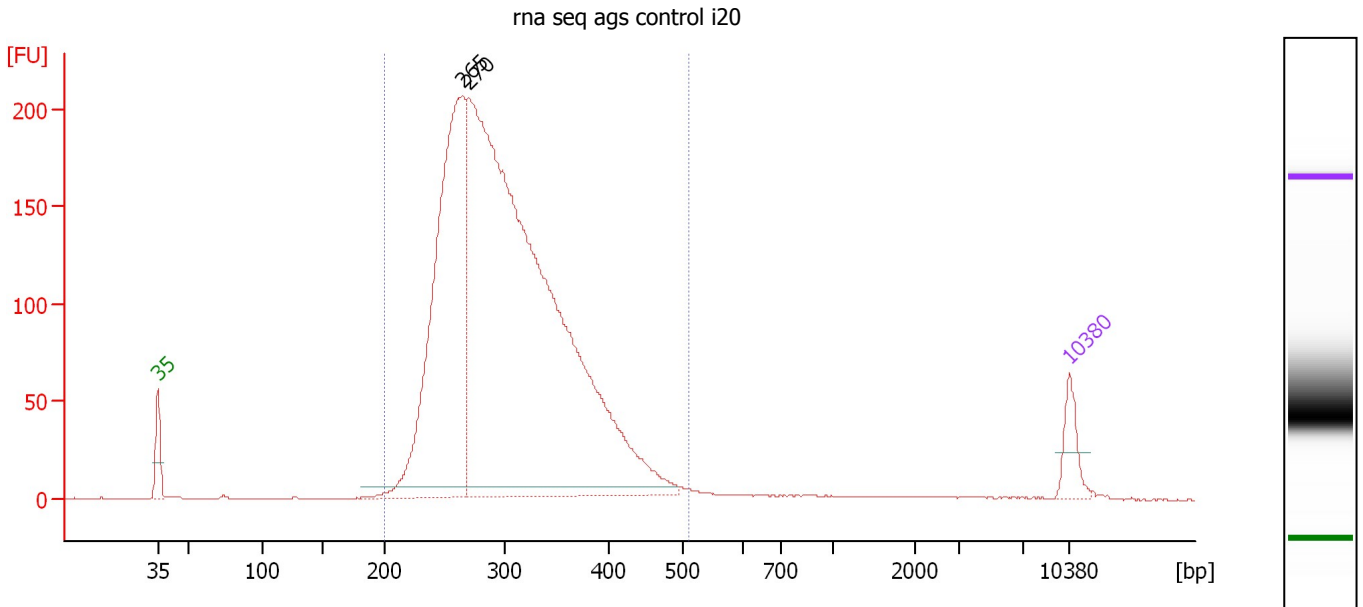
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	Ladder Peak
9	600	150.00	378.8	Ladder Peak
10	700	150.00	324.7	Ladder Peak
11	1,000	150.00	227.3	Ladder Peak
12	2,000	150.00	113.6	Ladder Peak
13	3,000	150.00	75.8	Ladder Peak
14	7,000	150.00	32.5	Ladder Peak
15	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 1 : rna seq ags control i20

Number of peaks found: 2 Corr. Area 1: 2,996.7
 Noise: 0.1

Peak table for sample 1 : rna seq ags control i20

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	265	1,695.91	9,680.7	
3	270	4,229.81	23,710.1	
4	10,380	75.00	10.9	Upper Marker

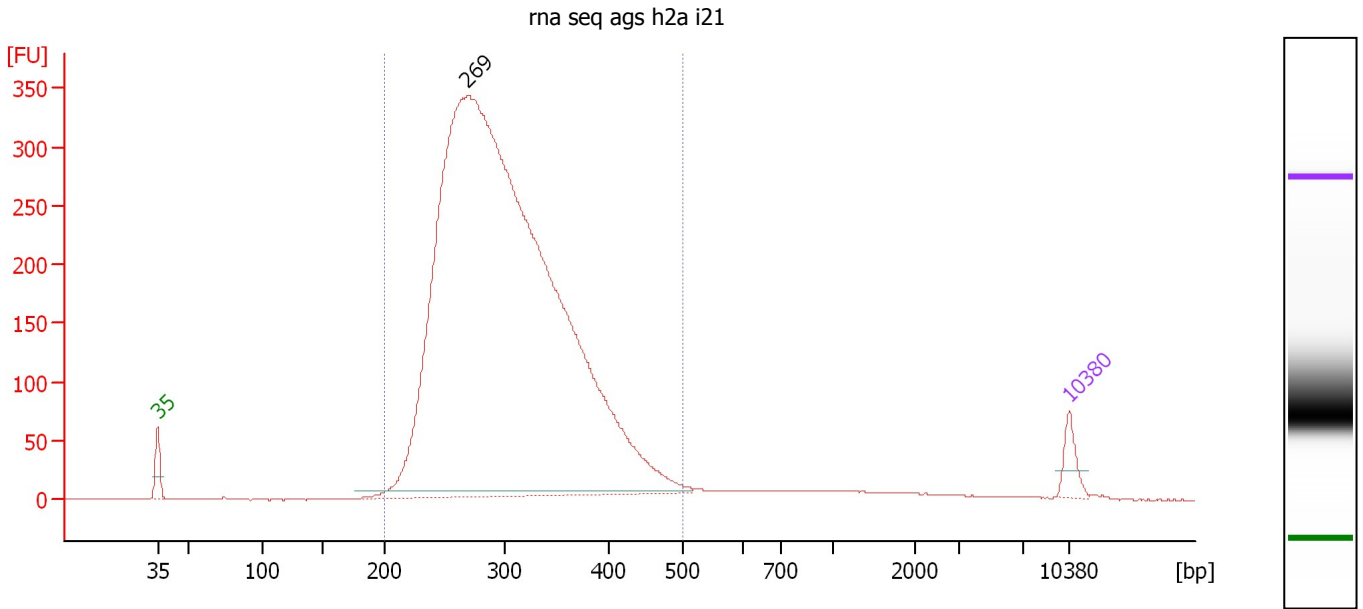
Region table for sample 1 : rna seq ags control i20

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	509	305	28,997.9	5,640.50	2,996.7	97	17.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
 Modified: 9/4/2013 3:32:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : rna seq ags h2a i21

Number of peaks found: 1 Corr. Area 1: 5,184.2
 Noise: 0.2

Peak table for sample 2 : rna seq ags h2a i21

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	269	9,652.29	54,343.7	
3	10,380	75.00	10.9	Upper Marker

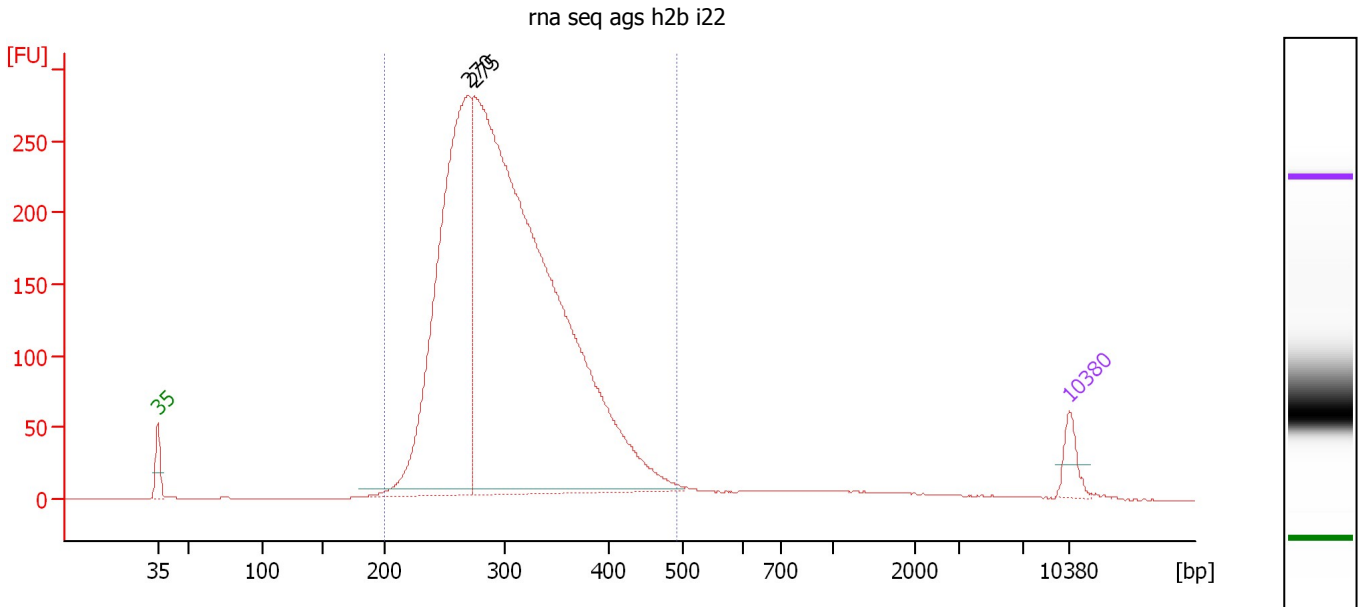
Region table for sample 2 : rna seq ags h2a i21

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	502	306	46,931.6	9,134.35	5,184.2	96	17.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
 Modified: 9/4/2013 3:32:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : rna seq ags h2b i22

Number of peaks found: 2 Corr. Area 1: 4,107.9
 Noise: 0.1

Peak table for sample 3 : rna seq ags h2b i22

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	270	2,552.86	14,348.8	
3	275	5,732.22	31,628.7	
4	10,380	75.00	10.9	Upper Marker

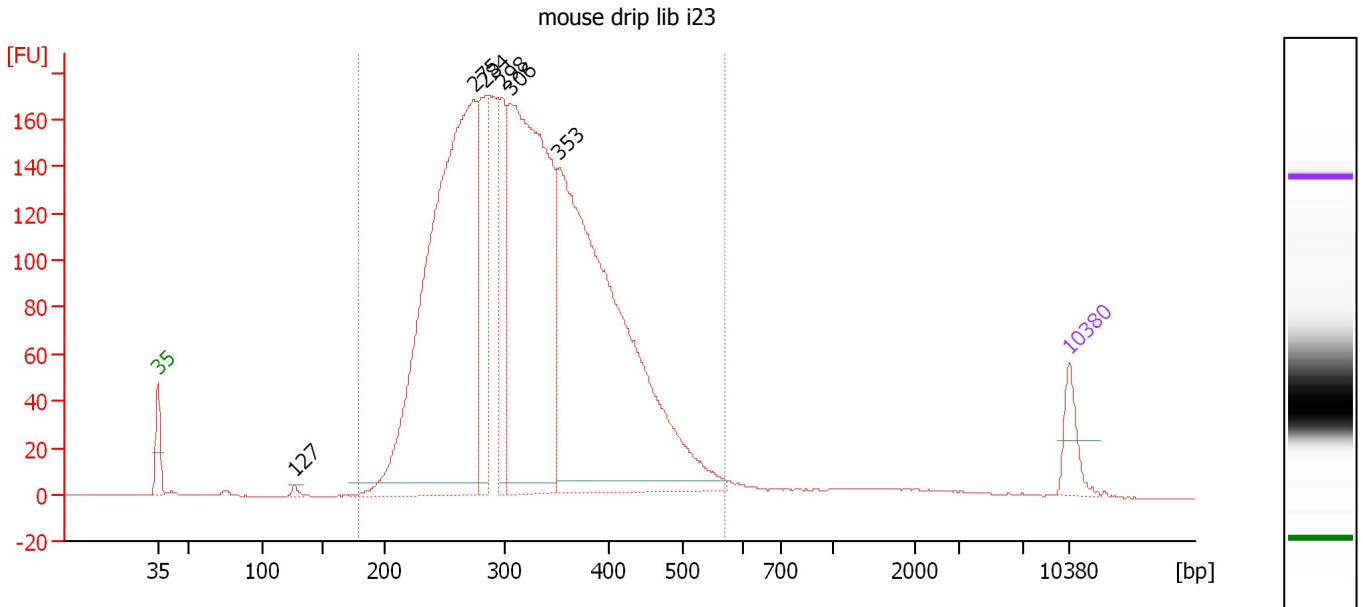
Region table for sample 3 : rna seq ags h2b i22

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	493	307	40,545.2	7,926.34	4,107.9	96	17.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
 Modified: 9/4/2013 3:32:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : mouse drip lib i23

Number of peaks found: 6 Corr. Area 1: 3,570.9
 Noise: 0.1

Peak table for sample 4 : mouse drip lib i23

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	127	13.03	155.8	
3	275	2,331.52	12,839.8	
4	284	470.87	2,510.3	
5	298	345.19	1,752.8	
6	306	1,753.61	8,686.1	
7	353	2,123.67	9,119.8	
8	10,380	75.00	10.9	Upper Marker

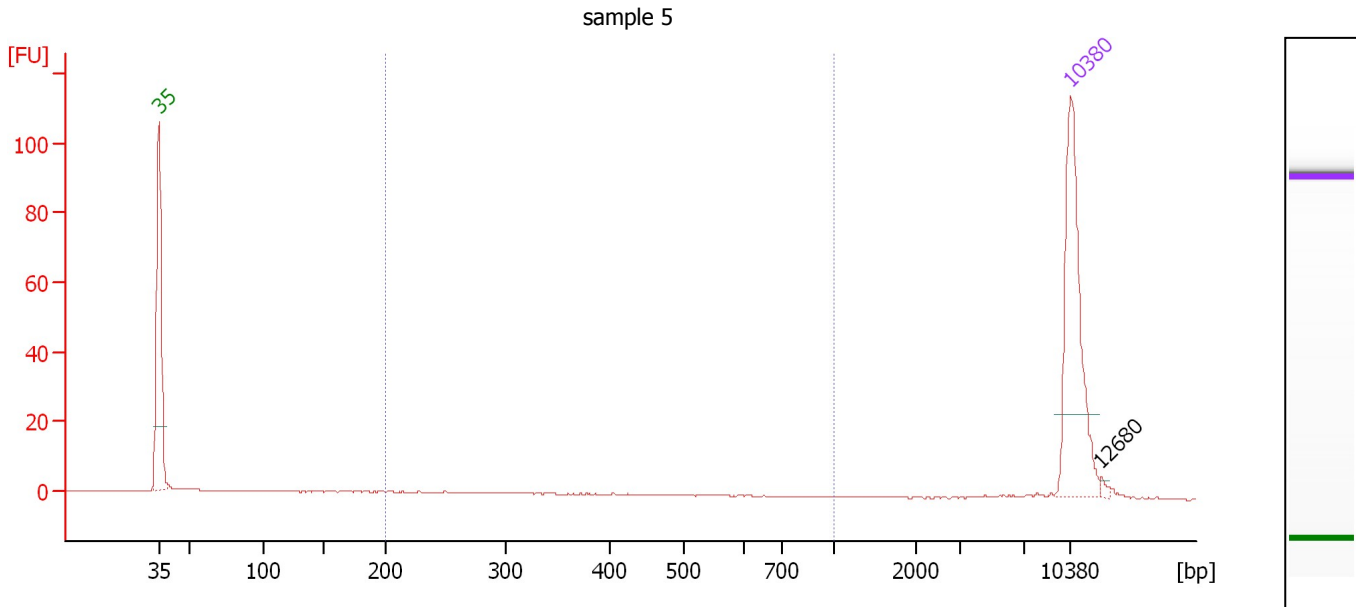
Region table for sample 4 : mouse drip lib i23

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
178	571	325	36,576.4	7,394.68	3,570.9	97	21.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
 Modified: 9/4/2013 3:32:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 1 Corr. Area 1: 7.4
 Noise: 0.1

Peak table for sample 5 : sample 5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	10,380	75.00	10.9	Upper Marker
3	12,680	0.00	0.0	

Region table for sample 5 : sample 5

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	309	34.5	6.40	7.4	24	29.2	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-09-04\2013-09-04_003.xad

Created: 9/4/2013 3:08:32 PM
Modified: 9/4/2013 3:32:42 PM

Invalid Samples

Sample 6 has not been run, no results available.

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.