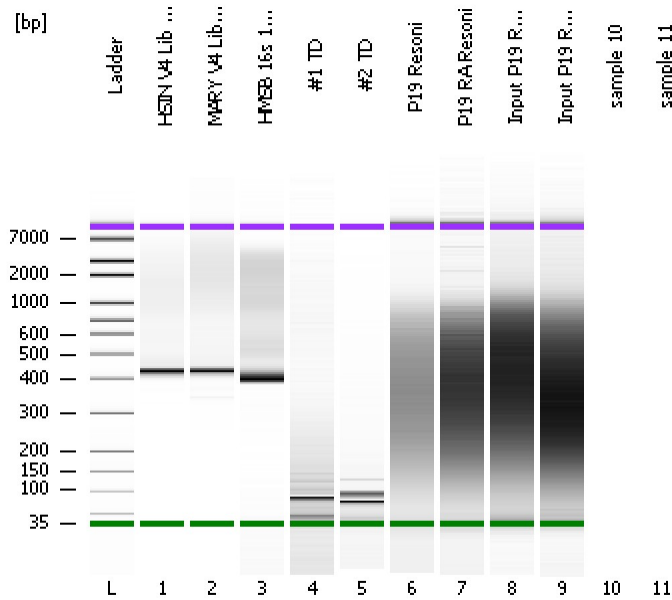


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
Modified: 1/29/2014 5:39:44 PM

Electrophoresis File Run Summary



Instrument Information:

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

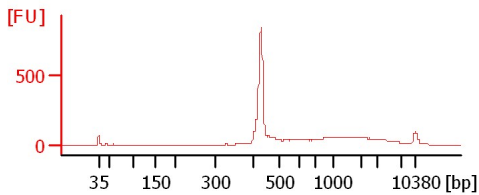
Assay Information:

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

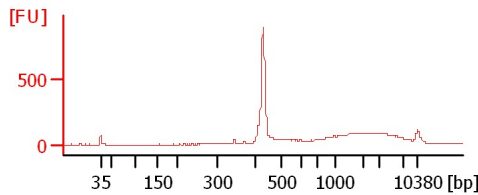
Chip Information:

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

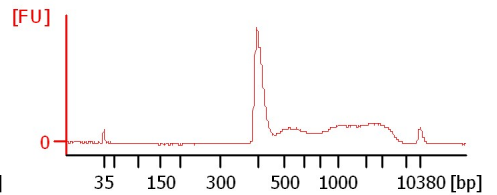
HSIN V4 Lib 1:45



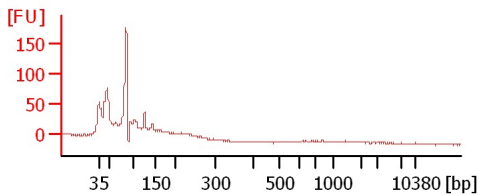
MARY V4 Lib 1:46



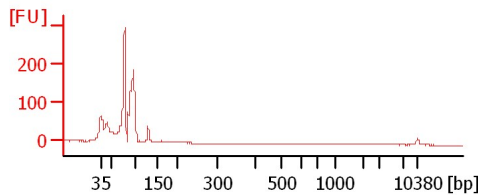
HMSB 16s 138 1:12



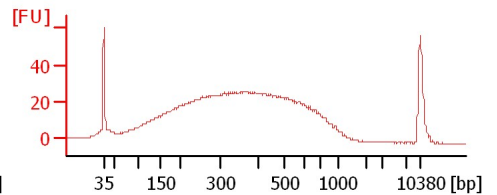
#1 TD



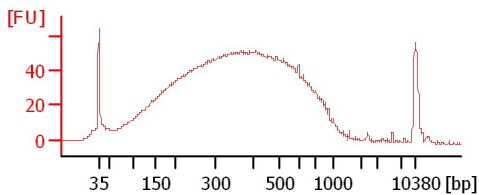
#2 TD



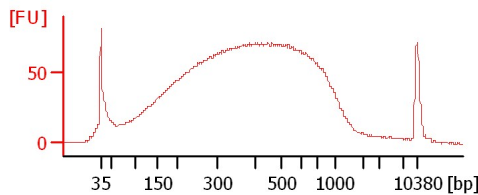
P19 Resoni



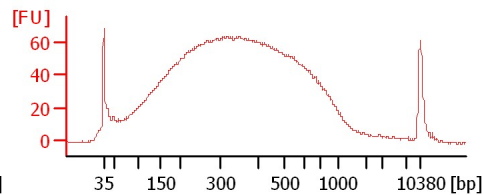
P19 RA Resoni



Input P19 Resoni 1:2



Input P19 RA Resoni 1:2



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
HSIN V4 Lib 1:45		<input type="checkbox"/>	✓			
MARY V4 Lib 1:46		<input type="checkbox"/>	✓			
HMSB 16s 138 1:12		<input type="checkbox"/>	✓			
#1 TD		<input type="checkbox"/>	✓			
#2 TD		<input type="checkbox"/>	✓			
P19 Resoni		<input type="checkbox"/>	✓			
P19 RA Resoni		<input type="checkbox"/>	✓			
Input P19 Resoni 1:2		<input type="checkbox"/>	✓			
Input P19 RA Resoni 1:2		<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\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
Modified: 1/29/2014 5:39:44 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

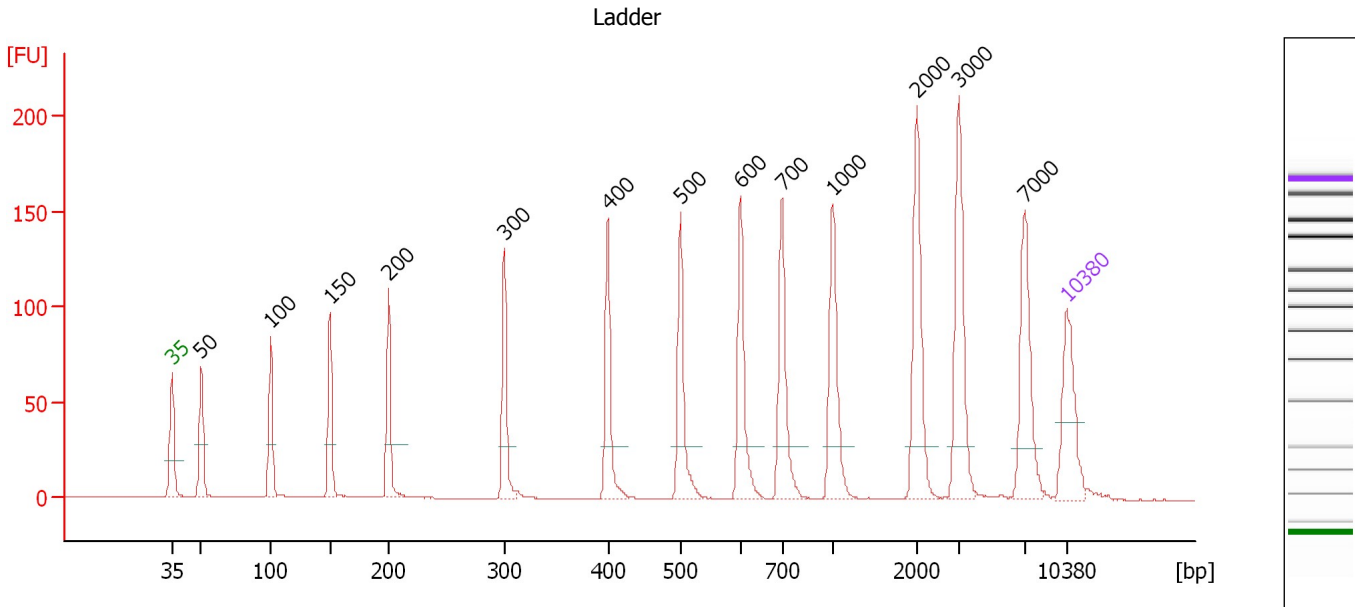
Ladder

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\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

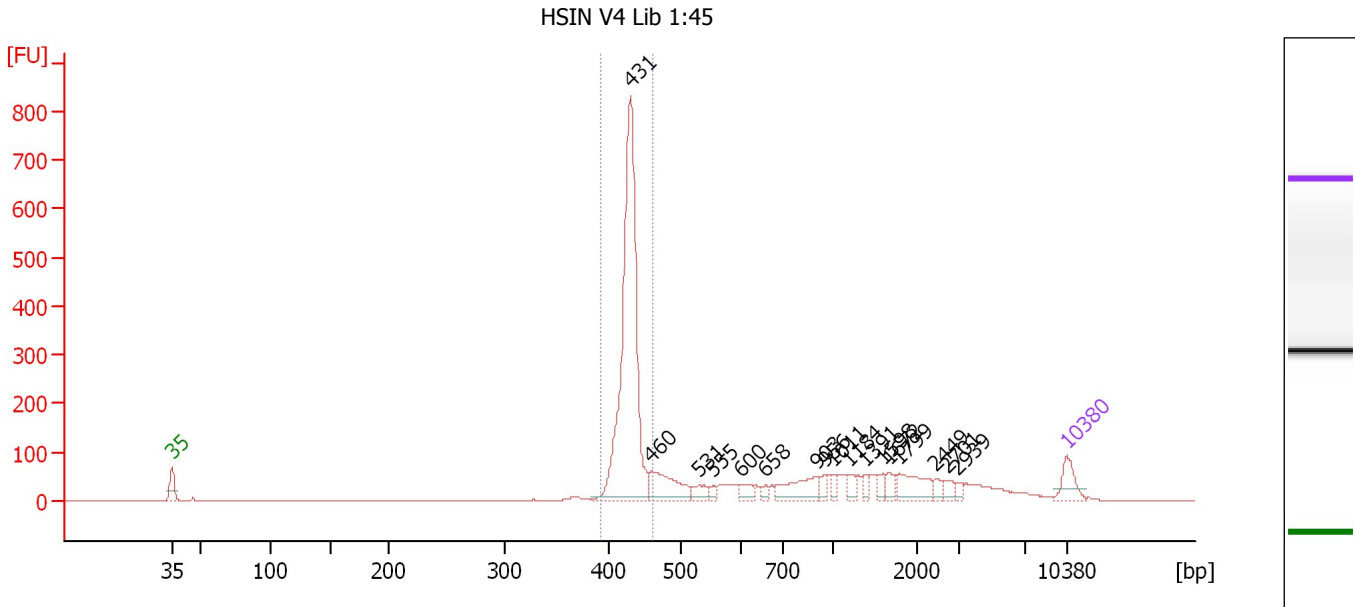
Peak table for Ladder

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	45.30	50	150.00	4,545.5	Ladder Peak
3	50.72	100	150.00	2,272.7	Ladder Peak
4	55.37	150	150.00	1,515.2	Ladder Peak
5	59.97	200	150.00	1,136.4	Ladder Peak
6	69.00	300	150.00	757.6	Ladder Peak
7	77.11	400	150.00	568.2	Ladder Peak
8	82.80	500	150.00	454.5	Ladder Peak
9	87.48	600	150.00	378.8	Ladder Peak
10	90.74	700	150.00	324.7	Ladder Peak
11	94.73	1,000	150.00	227.3	Ladder Peak
12	101.28	2,000	150.00	113.6	Ladder Peak
13	104.58	3,000	150.00	75.8	Ladder Peak
14	109.79	7,000	150.00	32.5	Ladder Peak
15	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 1 : HSIN V4 Lib 1:45

Height Threshold [FU] : 10

Overall Results for sample 1 : HSIN V4 Lib 1:45

Number of peaks found: 17 Corr. Area 1: 1,388.4
 Noise: 0.2

Peak table for sample 1 : HSIN V4 Lib 1:45

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	78.88	431	1,503.56	5,285.2	
3	80.54	460	210.00	691.3	
4	84.26	531	52.59	150.0	
5	85.35	555	23.67	64.7	
6	87.49	600	43.74	110.4	
7	89.38	658	22.12	50.9	
8	93.44	903	142.80	239.5	
9	94.14	956	36.78	58.3	
10	94.80	1,011	30.24	45.3	
11	95.94	1,184	41.81	53.5	
12	97.29	1,391	27.33	29.8	
13	98.65	1,598	36.04	34.2	
14	99.13	1,672	38.52	34.9	
15	99.96	1,799	117.84	99.3	
16	102.76	2,449	29.23	18.1	
17	103.59	2,701	28.61	16.1	
18	104.38	2,939	18.36	9.5	
19	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...

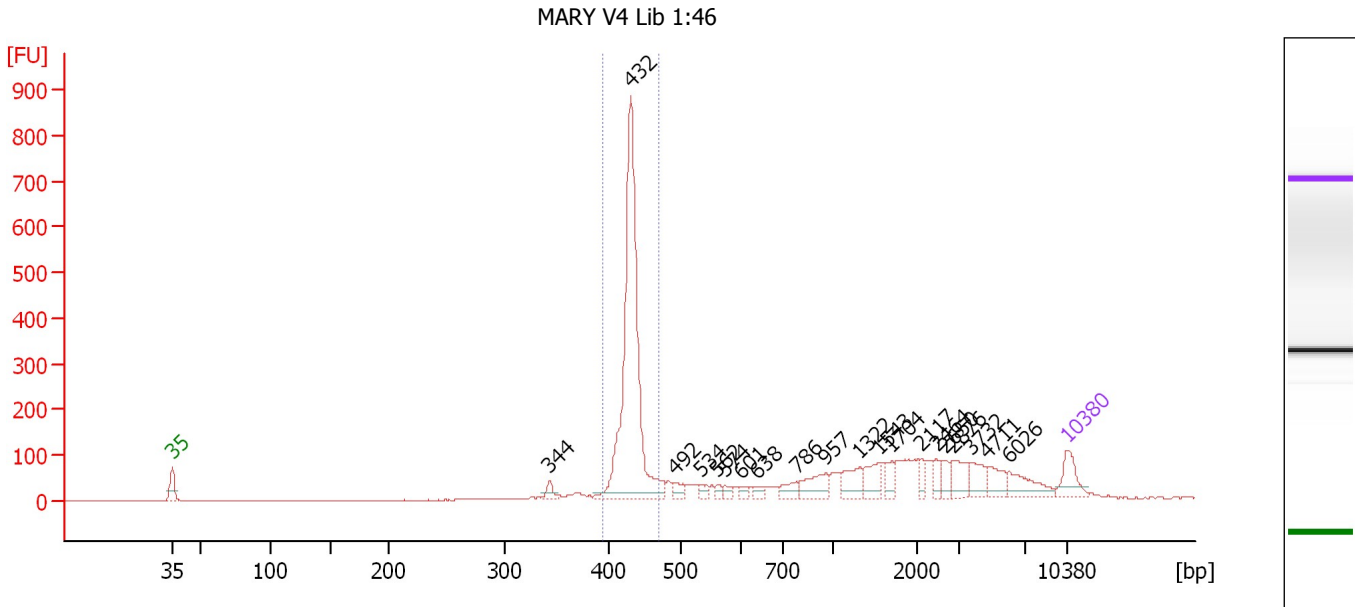
... Region table for sample 1 : HSIN V4 Lib 1:45

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
393	461	429	5,340.7	1,511.68	1,388.4	52	2.6	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 2 : MARY V4 Lib 1:46

Height Threshold [FU] : 15

Overall Results for sample 2 : MARY V4 Lib 1:46

Number of peaks found: 20 Corr. Area 1: 1,350.3
 Noise: 0.8

Peak table for sample 2 : MARY V4 Lib 1:46

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	72.55	344	33.63	148.2	
3	78.92	432	1,236.29	4,337.3	
4	82.35	492	34.54	106.3	
5	84.41	534	25.61	72.6	
6	85.69	562	17.73	47.8	
7	86.26	574	21.14	55.8	
8	87.53	601	16.12	40.6	
9	88.72	638	22.25	52.9	
10	91.88	786	41.71	80.4	
11	94.16	957	91.78	145.3	
12	96.84	1,322	82.58	94.6	
13	98.29	1,543	76.12	74.7	
14	99.34	1,704	36.71	32.6	
15	101.67	2,117	32.05	22.9	
16	102.81	2,464	38.68	23.8	
17	103.43	2,650	37.17	21.3	
18	104.17	2,876	72.48	38.2	
19	105.53	3,732	68.93	28.0	
20	106.81	4,711	62.37	20.1	
21	108.52	6,026	93.80	23.6	
22	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...

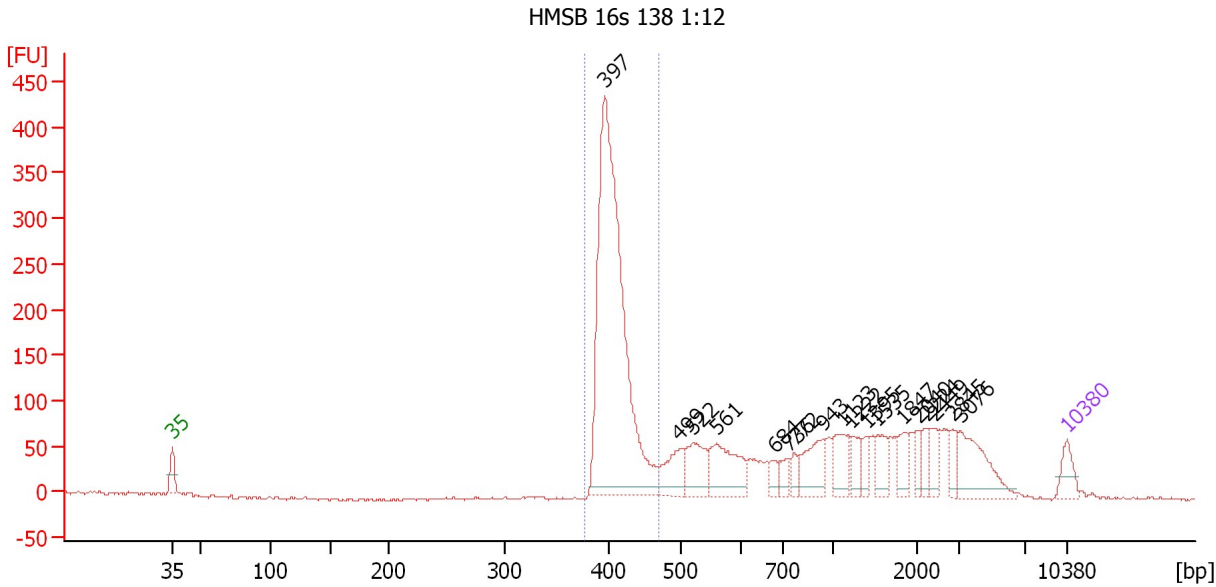
... Region table for sample 2 : MARY V4 Lib 1:46

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
396	470	432	4,229.2	1,205.04	1,350.3	43	2.7	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 3 : HMSB 16s 138 1:12

Height Threshold [FU] : 10

Overall Results for sample 3 : HMSB 16s 138 1:12

Number of peaks found: 18 Corr. Area 1: 1,297.6
 Noise: 1.7

Peak table for sample 3 : HMSB 16s 138 1:12

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	76.89	397	2,289.10	8,730.3	
3	82.73	499	184.87	561.6	
4	83.82	522	208.84	606.5	
5	85.64	561	294.13	794.7	
6	90.22	684	53.89	119.4	
7	91.22	736	58.11	119.6	
8	91.70	772	49.64	97.4	
9	93.96	943	191.06	307.1	
10	95.53	1,123	146.10	197.2	
11	96.19	1,222	82.41	102.2	
12	97.32	1,395	58.84	63.9	
13	98.23	1,535	96.14	94.9	
14	100.28	1,847	89.86	73.7	
15	101.41	2,040	54.31	40.3	
16	102.02	2,224	55.48	37.8	
17	102.76	2,449	72.10	44.6	
18	104.07	2,845	61.87	32.9	
19	104.68	3,076	236.88	116.7	
20	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...

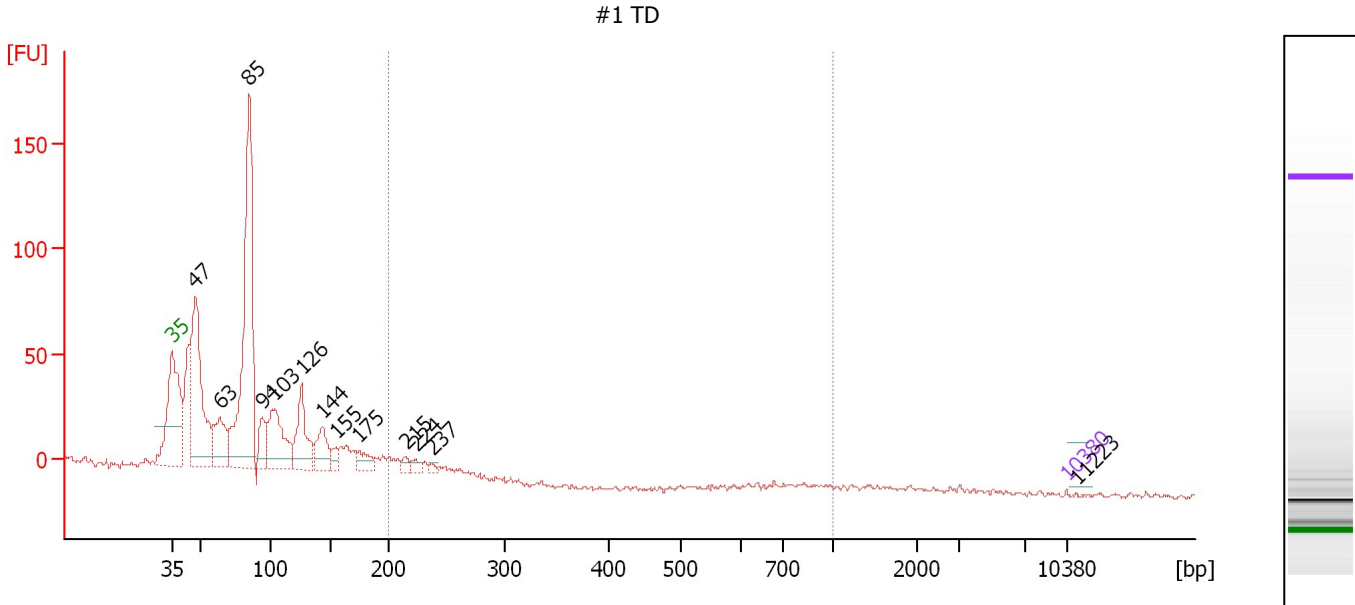
... Region table for sample 3 : HMSB 16s 138 1:12

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
378	472	410	8,388.8	2,263.81	1,297.6	44	4.3	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : #1 TD

Number of peaks found: 13 Corr. Area 1: 31.6
 Noise: 1.3

Peak table for sample 4 : #1 TD

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.82	47	75,498.78	2,439,515.3	
3	46.74	63	22,084.82	529,187.3	
4	49.05	85	120,575.83	2,160,793.5	
5	50.12	94	12,854.13	206,344.0	
6	51.05	103	32,229.80	471,827.7	
7	53.14	126	25,934.29	311,849.2	
8	54.79	144	14,284.36	150,588.6	
9	55.85	155	4,245.46	41,429.4	
10	57.68	175	6,984.87	60,446.2	
11	61.32	215	2,665.02	18,780.6	
12	62.12	224	2,635.02	17,833.6	
13	63.32	237	1,484.59	9,484.0	
14	113.00	10,380	75.00	10.9	Upper Marker
15	113.80	11,223	0.00	0.0	

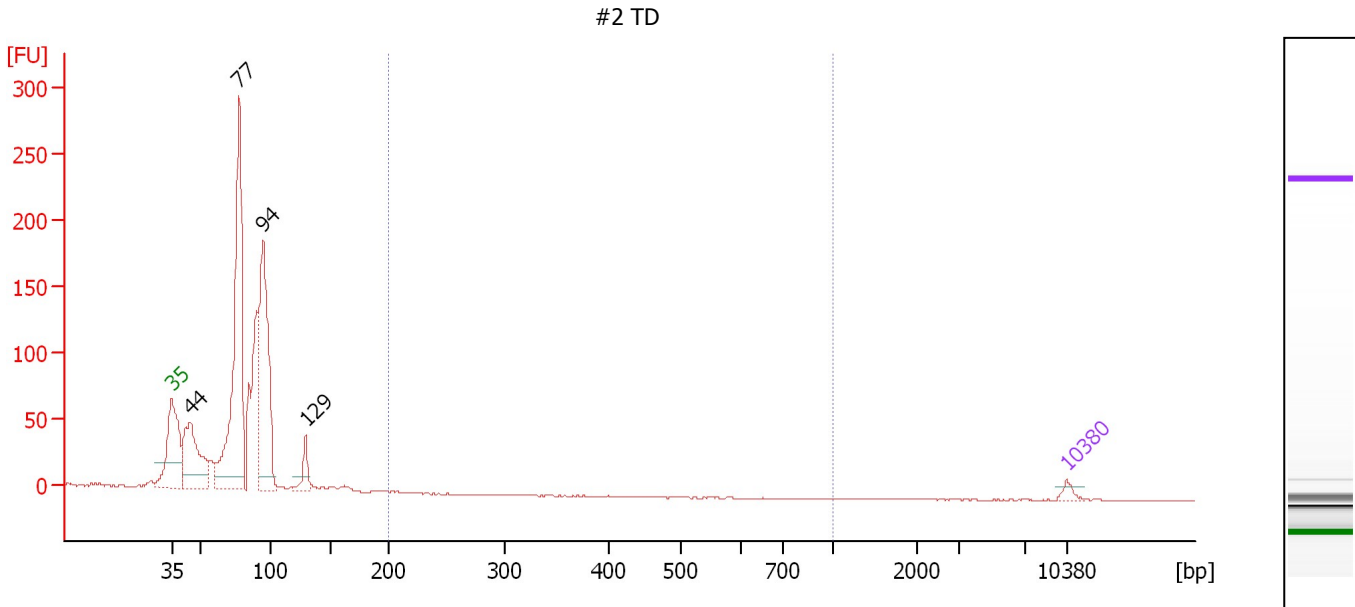
Region table for sample 4 : #1 TD

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	223	71,748.5	10,528.36	31.6	4	11.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 5 : #2 TD

Height Threshold [FU] : 10

Overall Results for sample 5 : #2 TD

Number of peaks found: 4 Corr. Area 1: 0.0
 Noise: 0.2

Peak table for sample 5 : #2 TD

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.44	44	2,144.81	73,181.0	
3	48.27	77	5,814.85	113,817.2	
4	50.12	94	4,044.39	64,885.1	
5	53.46	129	429.77	5,031.3	
6	113.00	10,380	75.00	10.9	Upper Marker

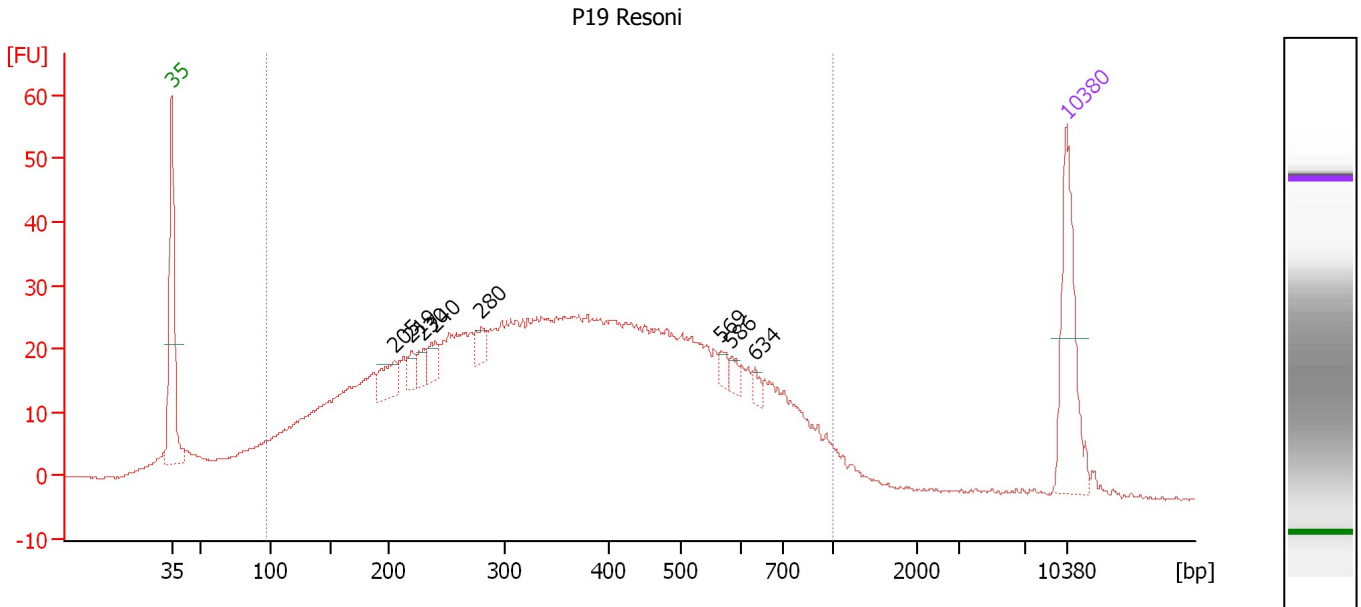
Region table for sample 5 : #2 TD

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	0	0.0	0.00	0.0	0	0.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : P19 Resoni

Number of peaks found: 8 Corr. Area 1: 1,275.7
 Noise: 0.1

Peak table for sample 6 : P19 Resoni

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	60.40	205	30.94	229.0	
3	61.72	219	13.26	91.6	
4	62.69	230	14.31	94.2	
5	63.60	240	18.28	115.3	
6	67.22	280	15.57	84.2	
7	86.03	569	7.97	21.2	
8	86.81	586	7.65	19.8	
9	88.60	634	6.95	16.6	
10	113.00	10,380	75.00	10.9	Upper Marker

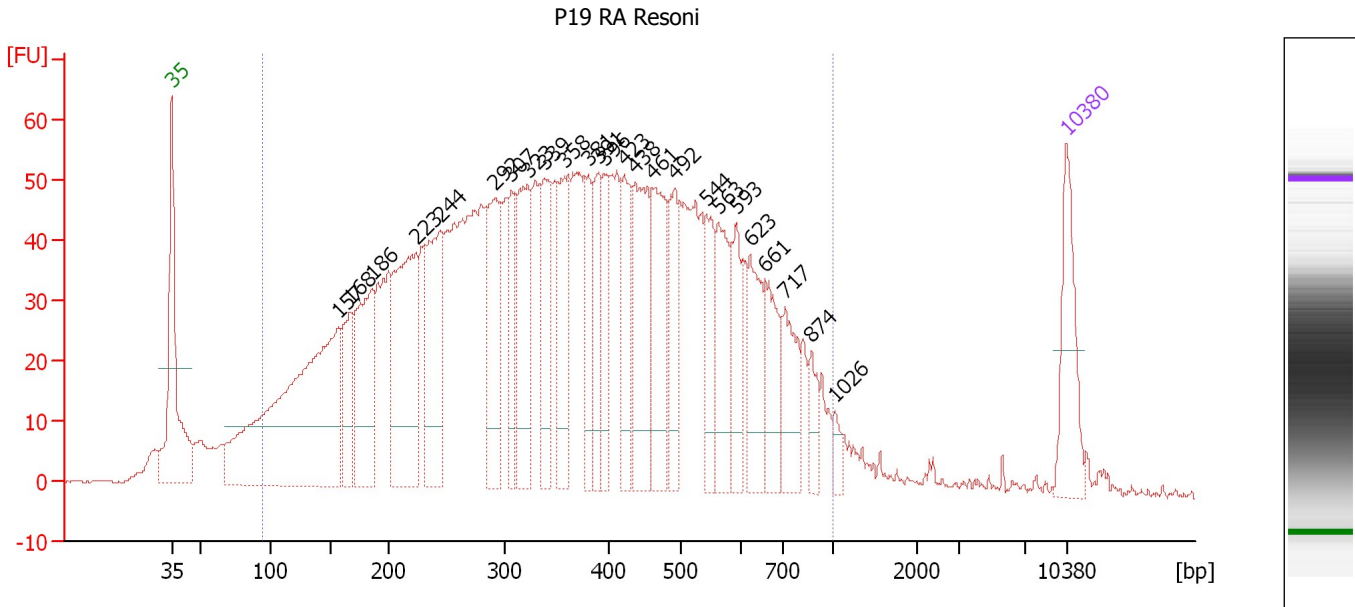
Region table for sample 6 : P19 Resoni

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
96	1,000	385	13,287.1	2,371.40	1,275.7	93	45.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 7 : P19 RA Resoni

Height Threshold [FU] : 10

Overall Results for sample 7 : P19 RA Resoni

Number of peaks found: 25 Corr. Area 1: 2,424.5
 Noise: 0.2

Peak table for sample 7 : P19 RA Resoni

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	56.06	157	601.01	5,782.8	
3	56.98	168	94.04	850.5	
4	58.65	186	191.17	1,560.2	
5	62.07	223	266.06	1,805.1	
6	63.97	244	192.63	1,194.4	
7	68.28	292	152.37	790.5	
8	69.57	307	82.20	405.5	
9	70.87	323	154.12	722.7	
10	72.17	339	115.84	517.6	
11	73.74	358	111.58	471.6	
12	75.55	381	72.79	289.7	
13	76.38	391	77.15	299.0	
14	76.80	396	71.22	272.4	
15	78.42	423	94.21	337.5	
16	79.25	438	156.73	542.6	
17	80.59	461	141.91	466.1	
18	82.35	492	85.18	262.2	
19	84.85	544	68.82	191.7	
20	85.73	563	101.29	272.8	
21	87.17	593	62.83	160.5	
22	88.23	623	93.16	226.6	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...

... Peak table for sample 7 : P19 RA Resoni

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
23	89.48	661	71.85	164.6	
24	90.96	717	74.55	157.5	
25	93.05	874	29.80	51.7	
26	94.90	1,026	14.10	20.8	
27	113.00	10,380	75.00	10.9	Upper Marker

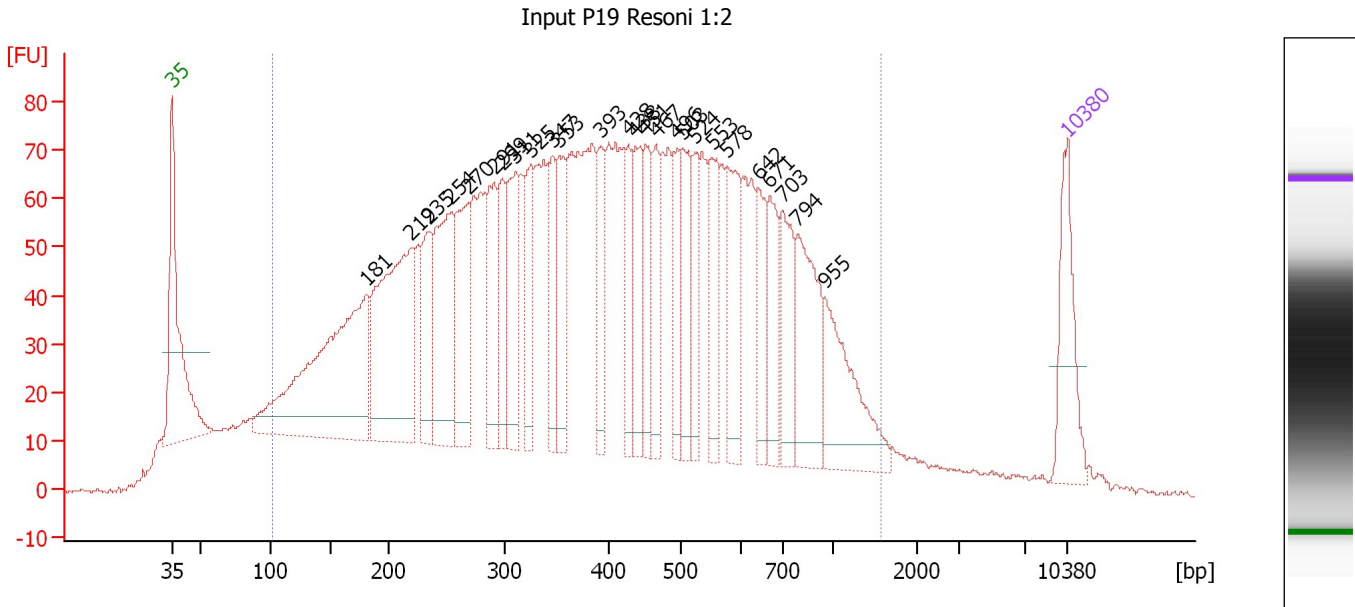
Region table for sample 7 : P19 RA Resoni

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
95	1,000	387	25,375.6	4,545.95	2,424.5	94	45.3	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

Created: 1/29/2014 4:52:27 PM
 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Input P19 Resoni 1:2

Number of peaks found: 26 Corr. Area 1: 3,527.7
 Noise: 0.6

Peak table for sample 8 : Input P19 Resoni 1:2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	58.18	181	432.77	3,632.2	
3	61.68	219	327.99	2,269.7	
4	63.13	235	97.49	628.6	
5	64.85	254	189.69	1,131.0	
6	66.26	270	134.50	755.7	
7	68.22	291	106.14	551.9	
8	68.87	299	78.62	398.9	
9	69.90	311	122.57	596.9	
10	71.02	325	77.71	362.4	
11	72.79	347	71.45	312.2	
12	73.31	353	81.90	351.4	
13	76.58	393	72.51	279.3	
14	78.68	428	65.34	231.6	
15	79.28	438	79.78	275.8	
16	80.03	451	74.28	249.3	
17	80.92	467	68.37	221.8	
18	82.55	496	62.06	189.7	
19	82.93	503	80.45	242.4	
20	83.91	524	59.29	171.5	
21	85.26	553	81.55	223.6	
22	86.43	578	103.83	272.4	
23	88.86	642	61.44	145.0	
24	89.79	671	69.62	157.2	
25	90.77	703	81.58	175.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad


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 Modified: 1/29/2014 5:39:44 PM

Electropherogram Summary Continued ...

... Peak table for sample 8 : Input P19 Resoni 1:2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations
26	91.99	794	120.59	230.2	
27	94.13	955	116.88	185.4	
28	113.00	10,380	75.00	10.9	Upper Marker

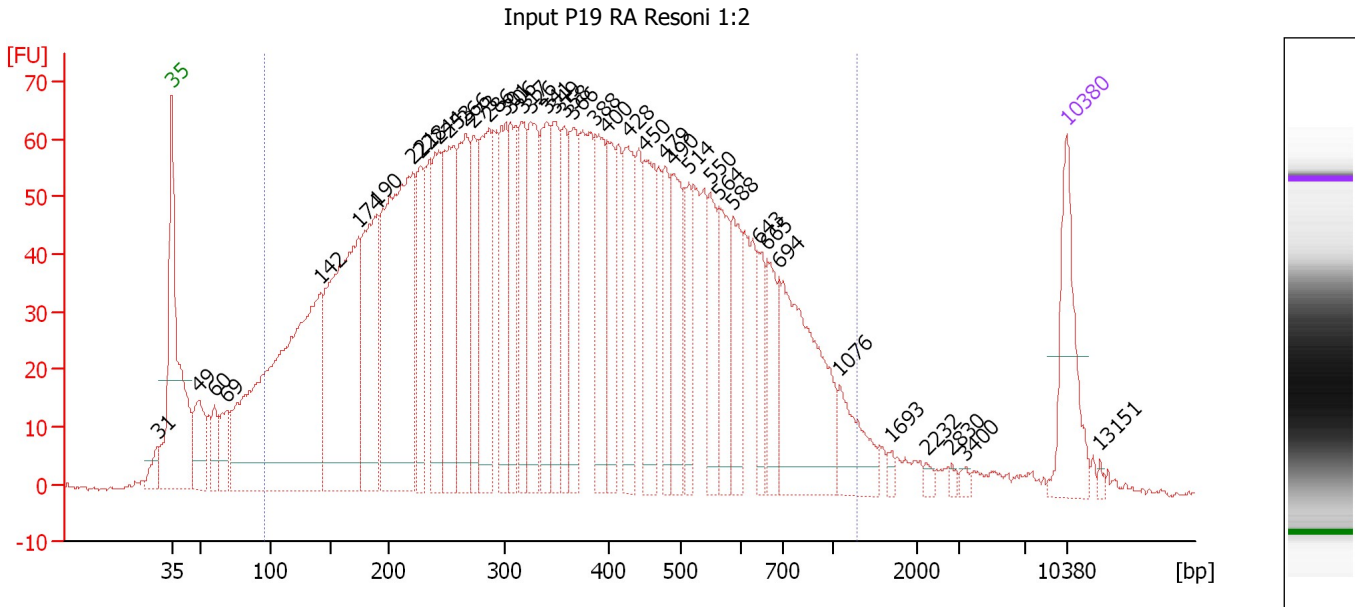
Region table for sample 8 : Input P19 Resoni 1:2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
101	1,583	442	26,136.7	5,012.90	3,527.7	93	54.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : Input P19 RA Resoni 1:2

Number of peaks found: 41 Corr. Area 1: 3,160.8
 Noise: 0.5

Peak table for sample 9 : Input P19 RA Resoni 1:2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	41.87	31	0.00	0.0	
2	43.00	35	125.00	5,411.3	Lower Marker
3	45.12	49	74.78	2,321.8	
4	46.34	60	39.87	1,014.3	
5	47.33	69	44.10	973.0	
6	54.61	142	607.27	6,486.3	
7	57.57	174	380.68	3,315.5	
8	59.03	190	188.91	1,508.0	
9	61.85	221	409.99	2,812.6	
10	62.46	228	103.06	686.0	
11	63.92	244	125.19	778.1	
12	64.72	253	176.13	1,056.3	
13	65.94	266	159.96	910.6	
14	66.60	273	93.28	516.9	
15	67.78	286	155.12	820.4	
16	69.09	301	117.50	591.1	
17	69.47	306	89.49	443.4	
18	70.41	317	93.98	448.6	
19	71.07	326	131.32	611.2	
20	72.34	341	114.03	506.4	
21	72.95	349	93.63	406.8	
22	73.70	358	78.44	332.0	
23	74.36	366	94.44	390.9	
24	76.14	388	104.73	408.9	
25	77.08	400	79.22	300.3	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

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Electropherogram Summary Continued ...

... Peak table for sample 9 : Input P19 RA Resoni 1:2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
26	78.68	428	111.51	395.1	
27	79.95	450	105.80	356.3	
28	81.60	479	59.69	188.8	
29	82.25	490	93.07	287.5	
30	83.43	514	54.85	161.8	
31	85.12	550	64.64	178.2	
32	85.78	564	74.92	201.4	
33	86.91	588	62.99	162.4	
34	88.88	643	37.65	88.7	
35	89.59	665	54.57	124.4	
36	90.53	694	194.81	425.6	
37	95.23	1,076	50.94	71.7	
38	99.27	1,693	5.32	4.8	
39	102.05	2,232	4.34	3.0	
40	104.02	2,830	3.33	1.8	
41	105.10	3,400	3.94	1.8	
42	113.00	10,380	75.00	10.9	Upper Marker
43	115.63	13,151	0.00	0.0	

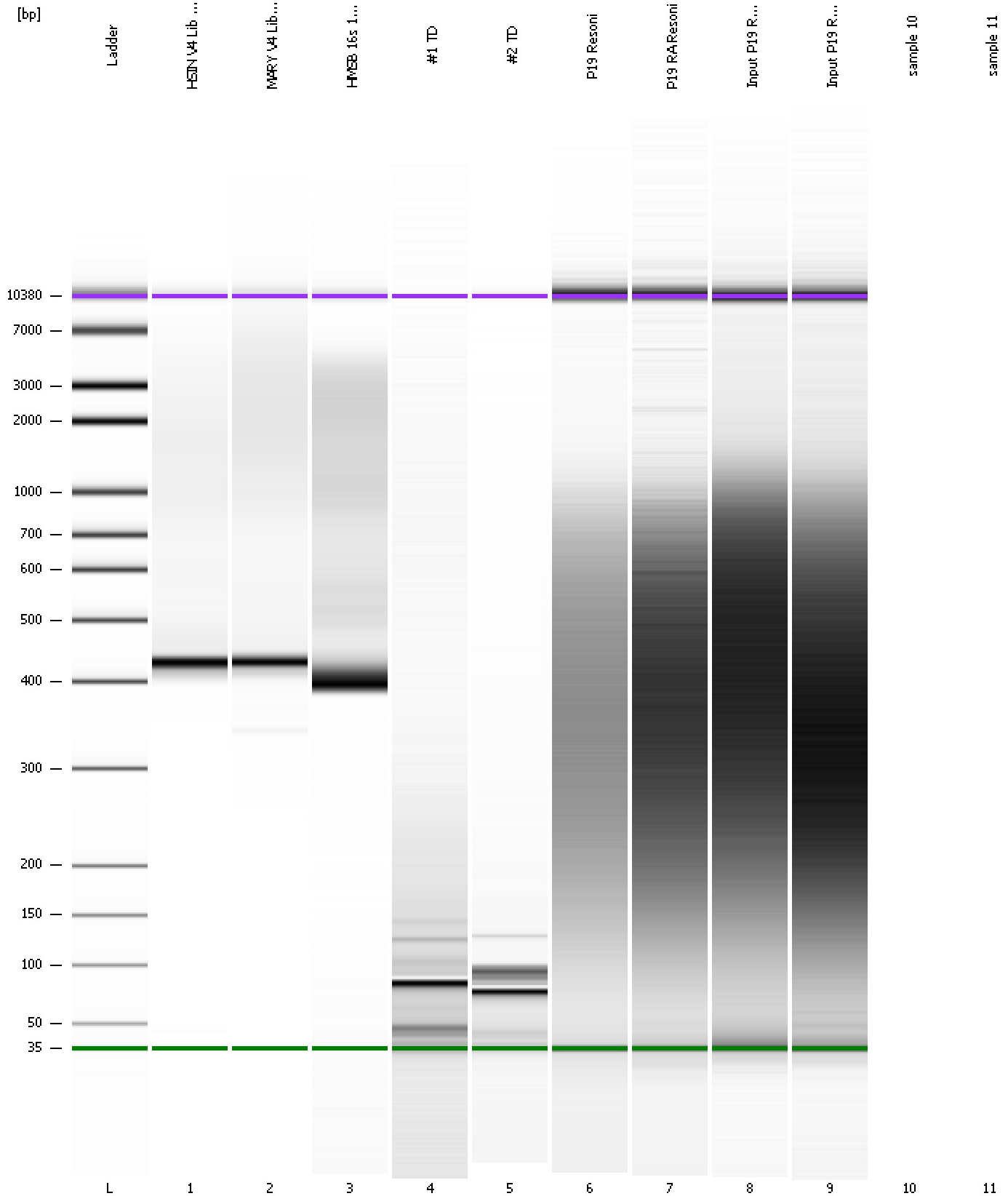
Region table for sample 9 : Input P19 RA Resoni 1:2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
96	1,297	387	28,837.4	4,981.87	3,160.8	92	51.7	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

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Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-29\2014-01-29_004.xad

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Modified: 1/29/2014 5:39:44 PM

Invalid Samples

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.