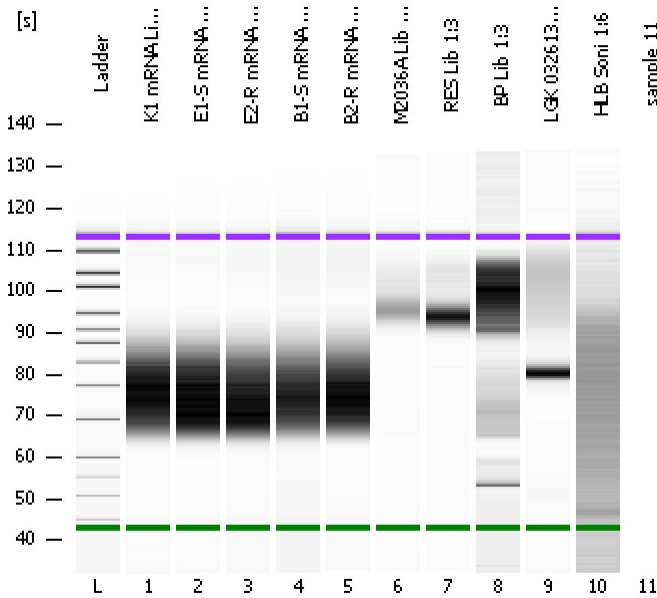


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
Modified: 3/26/2013 5:27:33 PM

Electrophoresis File Run Summary



Instrument Information:

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

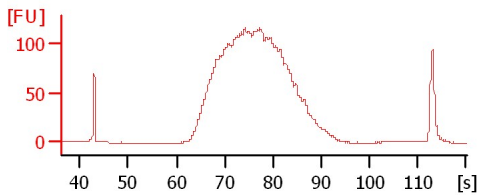
Assay Information:

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

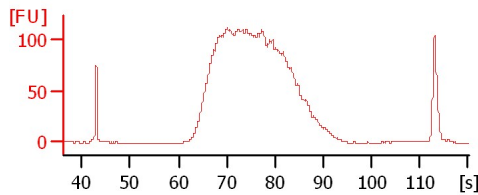
Chip Information:

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

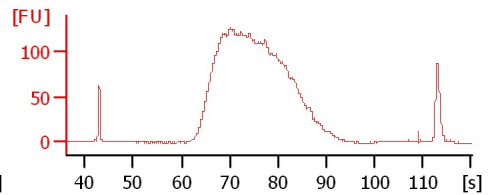
K1 mRNA Lib 1:2



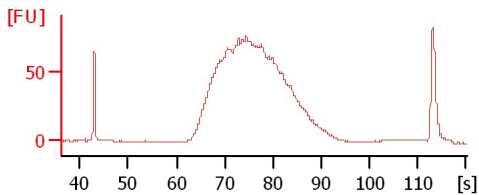
E1-S mRNA Lib 1:2



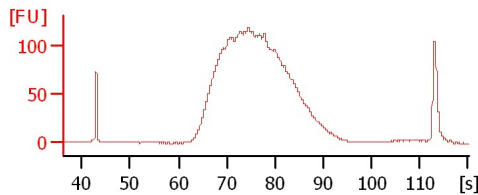
E2-R mRNA Lib 1:2



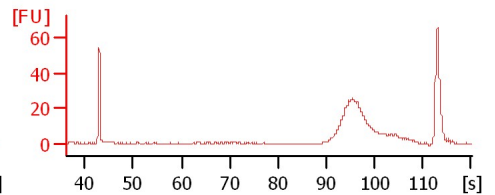
B1-S mRNA Lib 1:5



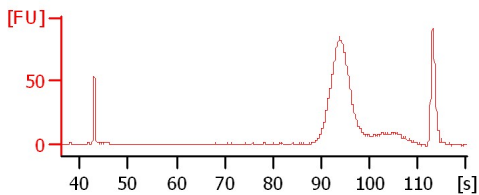
B2-R mRNA Lib 1:4



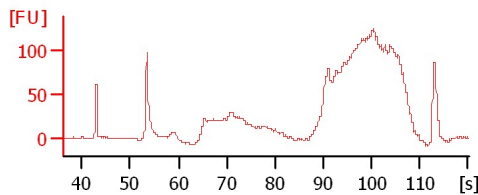
M2036A Lib 1:3



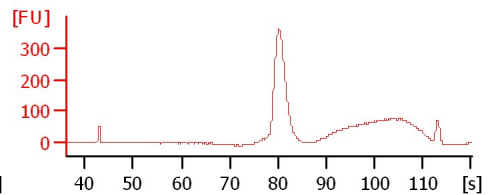
RES Lib 1:3



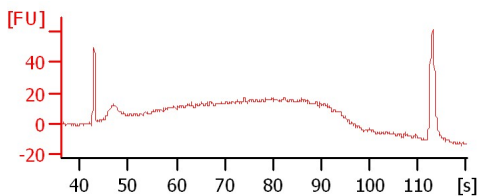
BP Lib 1:3



LGK 032613 1:12



HLB Soni 1:6



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
Modified: 3/26/2013 5:27:33 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
K1 mRNA Lib 1:2		<input type="checkbox"/>	✓			
E1-S mRNA Lib 1:2		<input type="checkbox"/>	✓			
E2-R mRNA Lib 1:2		<input type="checkbox"/>	✓			
B1-S mRNA Lib 1:5		<input type="checkbox"/>	✓			
B2-R mRNA Lib 1:4		<input type="checkbox"/>	✓			
M2036A Lib 1:3		<input type="checkbox"/>	✓			
RES Lib 1:3		<input type="checkbox"/>	✓			
BP Lib 1:3		<input type="checkbox"/>	✓			
LGK 032613 1:12		<input type="checkbox"/>	✓			
HLB Soni 1:6		<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\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 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

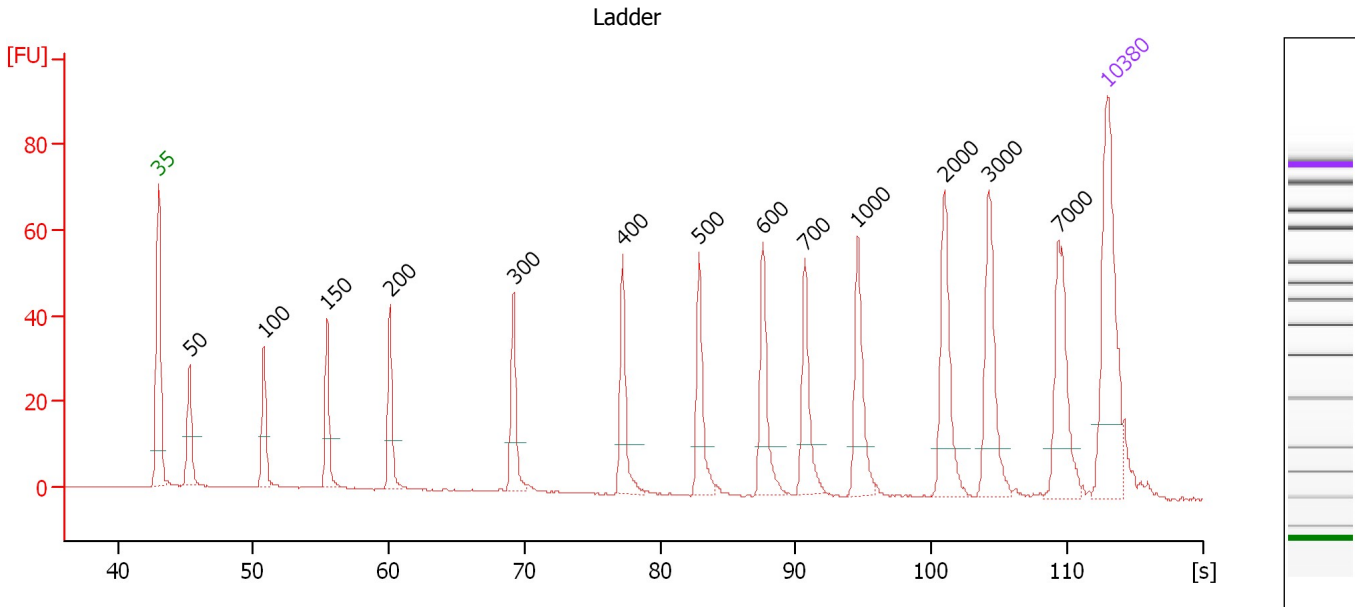
Ladder

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\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

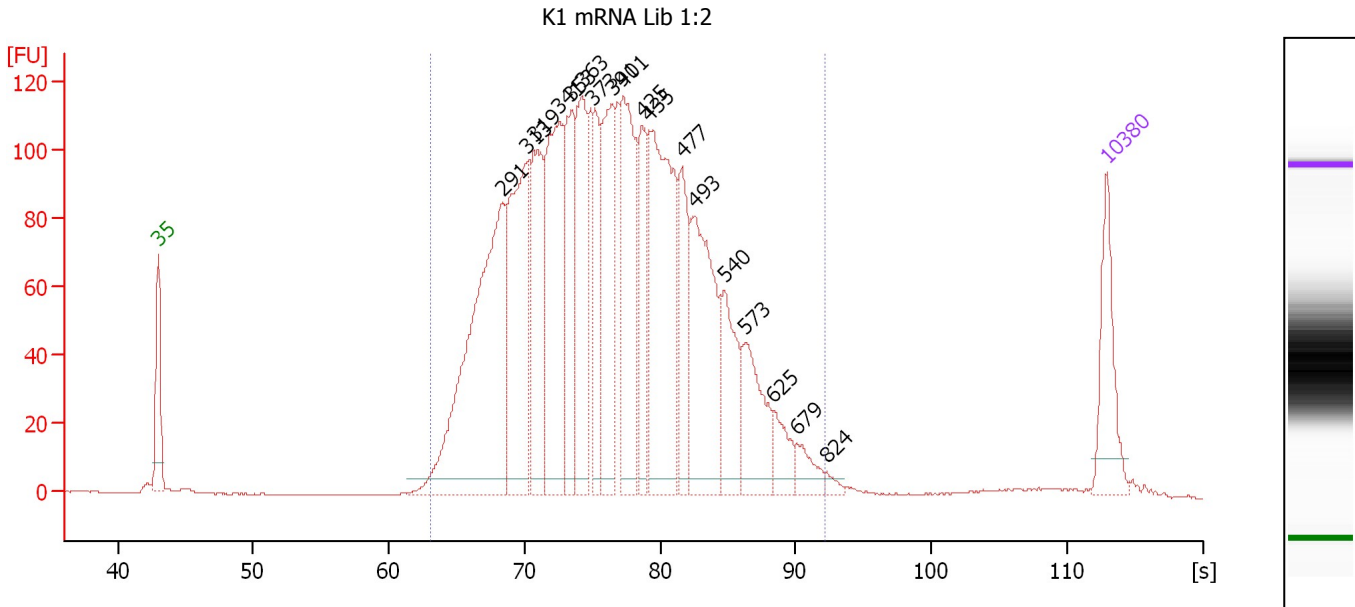
Peak table for Ladder

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\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : K1 mRNA Lib 1:2

Number of peaks found: 18 Corr. Area 1: 2,856.1
 Noise: 0.2

Peak table for sample 1 : K1 mRNA Lib 1:2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	291	529.41	2,761.2	
3	313	301.75	1,461.9	
4	319	199.76	949.4	
5	341	276.92	1,229.5	
6	353	161.91	695.4	
7	363	180.37	753.6	
8	373	122.33	496.8	
9	391	189.91	736.7	
10	401	222.75	840.7	
11	425	100.96	359.5	
12	435	334.96	1,167.2	
13	477	110.53	350.7	
14	493	236.18	725.9	
15	540	108.28	303.7	
16	573	104.55	276.5	
17	625	38.78	93.9	
18	679	30.60	68.2	
19	824	7.32	13.5	
20	10,380	75.00	10.9	Upper Marker

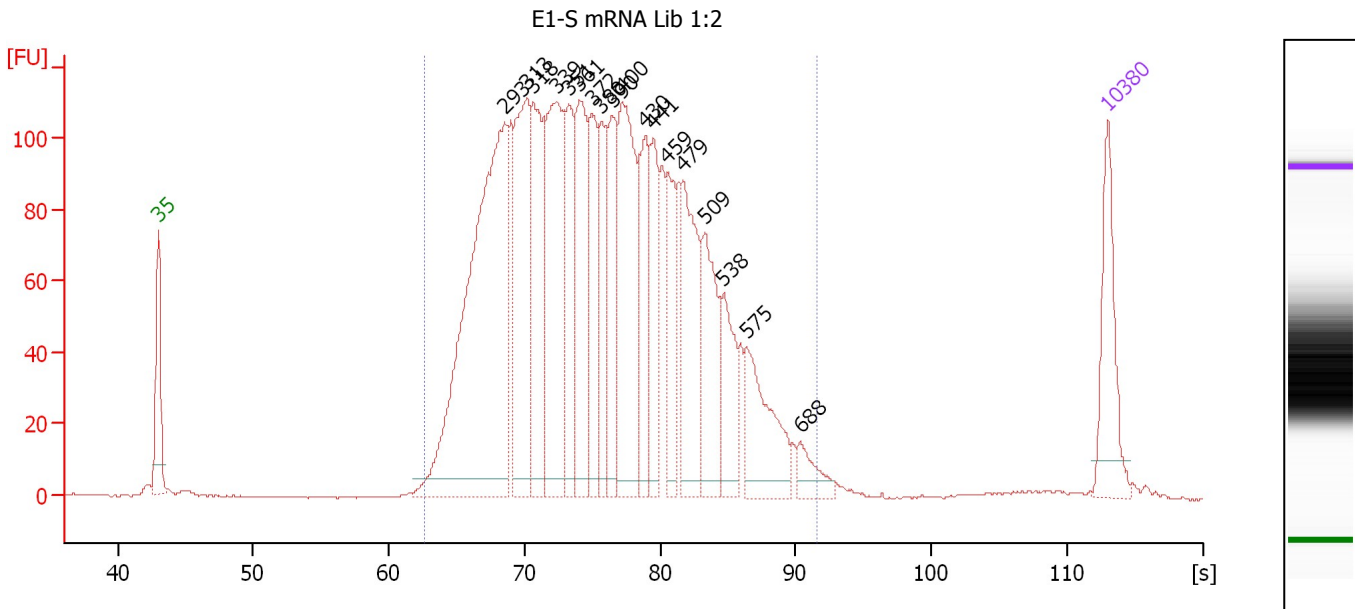
Region table for sample 1 : K1 mRNA Lib 1:2

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color
63.16	92.21	402	14,026.9	3,430.58	2,856.1 98	24.5	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : E1-S mRNA Lib 1:2

Number of peaks found: 18 Corr. Area 1: 2,926.5
 Noise: 0.3

Peak table for sample 2 : E1-S mRNA Lib 1:2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	293	616.04	3,186.8	
3	313	246.78	1,196.3	
4	318	170.07	809.9	
5	339	263.38	1,175.8	
6	351	112.94	488.1	
7	361	162.32	682.0	
8	372	118.80	484.1	
9	380	87.86	350.2	
10	390	113.60	441.1	
11	400	219.36	830.0	
12	430	106.91	377.0	
13	441	93.32	320.8	
14	459	94.74	312.8	
15	479	158.09	500.4	
16	509	119.14	354.9	
17	538	75.44	212.4	
18	575	107.50	283.1	
19	688	29.45	64.8	
20	10,380	75.00	10.9	Upper Marker

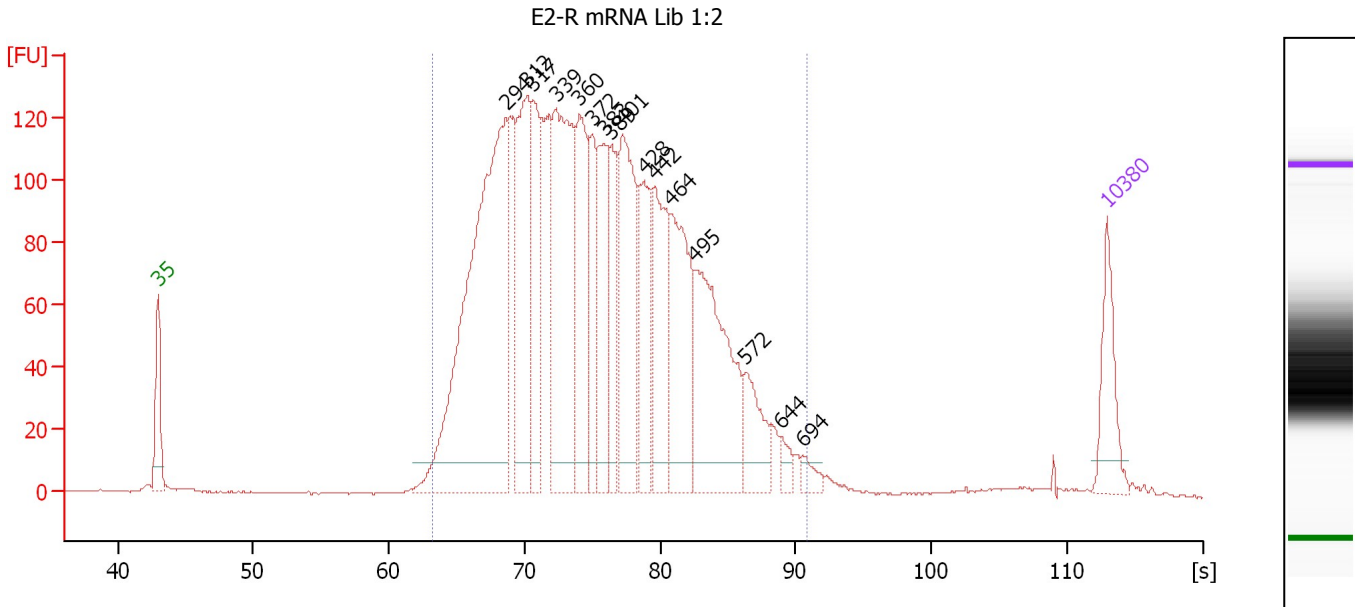
Region table for sample 2 : E1-S mRNA Lib 1:2

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color	
62.69	91.64	393	13,089.3	3,120.48	2,926.5	98	25.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 3 : E2-R mRNA Lib 1:2

Height Threshold [FU] : 10

Overall Results for sample 3 : E2-R mRNA Lib 1:2

Number of peaks found: 16 Corr. Area 1: 3,098.7
 Noise: 0.3

Peak table for sample 3 : E2-R mRNA Lib 1:2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	294	917.78	4,728.4	
3	312	303.14	1,472.3	
4	317	202.87	969.4	
5	339	439.93	1,968.5	
6	360	216.15	909.3	
7	372	124.90	508.6	
8	383	175.11	692.0	
9	389	117.13	456.1	
10	401	267.13	1,010.3	
11	428	146.21	517.7	
12	442	201.62	691.8	
13	464	250.85	819.1	
14	495	331.16	1,013.0	
15	572	87.97	233.2	
16	644	21.56	50.7	
17	694	19.93	43.5	
18	10,380	75.00	10.9	Upper Marker

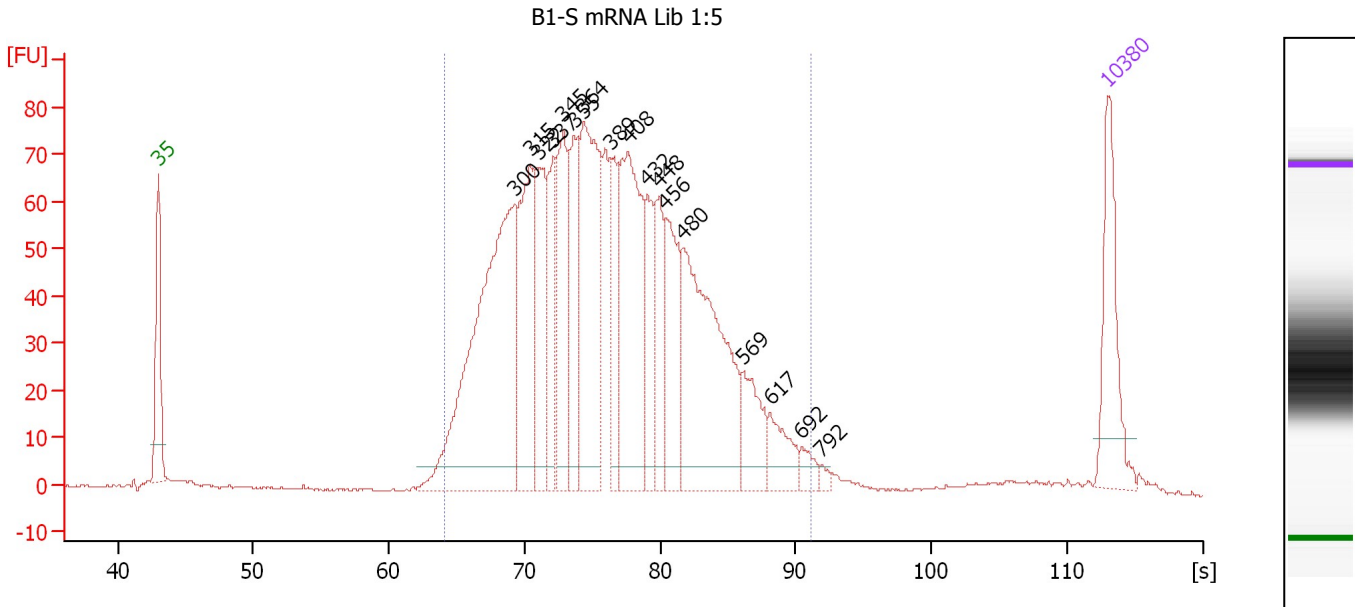
Region table for sample 3 : E2-R mRNA Lib 1:2

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color
63.26	90.80	385	17,988.7	4,226.77	3,098.7 97	24.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : B1-S mRNA Lib 1:5

Number of peaks found: 17 Corr. Area 1: 1,755.9
 Noise: 0.3

Peak table for sample 4 : B1-S mRNA Lib 1:5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	300	444.89	2,244.4	
3	315	163.58	787.4	
4	322	131.20	616.8	
5	337	79.79	359.0	
6	345	121.16	531.5	
7	355	100.76	429.8	
8	364	224.48	934.7	
9	389	71.65	278.9	
10	408	211.46	785.8	
11	432	78.42	274.9	
12	448	67.35	227.9	
13	456	104.07	345.8	
14	480	256.68	810.8	
15	569	57.85	154.0	
16	617	37.49	92.0	
17	692	14.21	31.1	
18	792	4.88	9.3	
19	10,380	75.00	10.9	Upper Marker

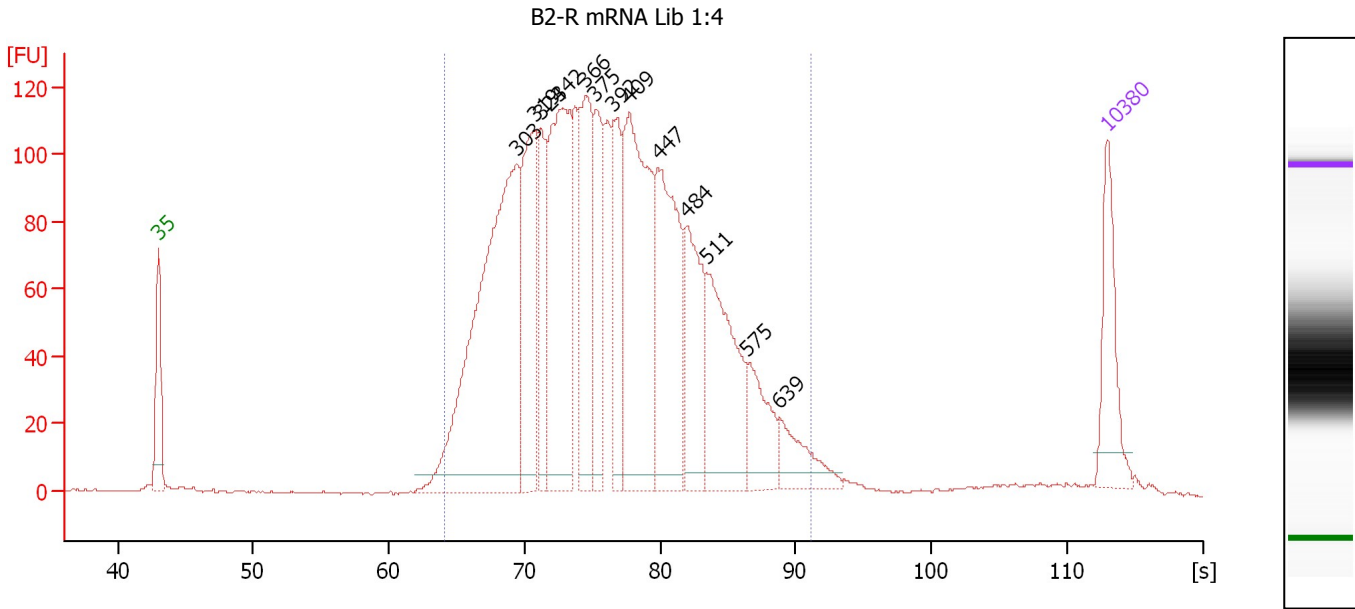
Region table for sample 4 : B1-S mRNA Lib 1:5

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color
64.15	91.17	397	9,346.3	2,269.81	1,755.9	95	23.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : B2-R mRNA Lib 1:4

Number of peaks found: 13 Corr. Area 1: 2,764.2
 Noise: 0.3

Peak table for sample 5 : B2-R mRNA Lib 1:4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	303	614.53	3,071.8	
3	319	204.20	968.4	
4	324	116.20	543.1	
5	342	316.67	1,404.2	
6	366	179.88	745.3	
7	375	124.10	501.4	
8	392	116.66	450.9	
9	409	338.31	1,252.3	
10	447	244.75	830.4	
11	484	149.72	468.9	
12	511	195.31	578.7	
13	575	79.07	208.3	
14	639	58.01	137.5	
15	10,380	75.00	10.9	Upper Marker

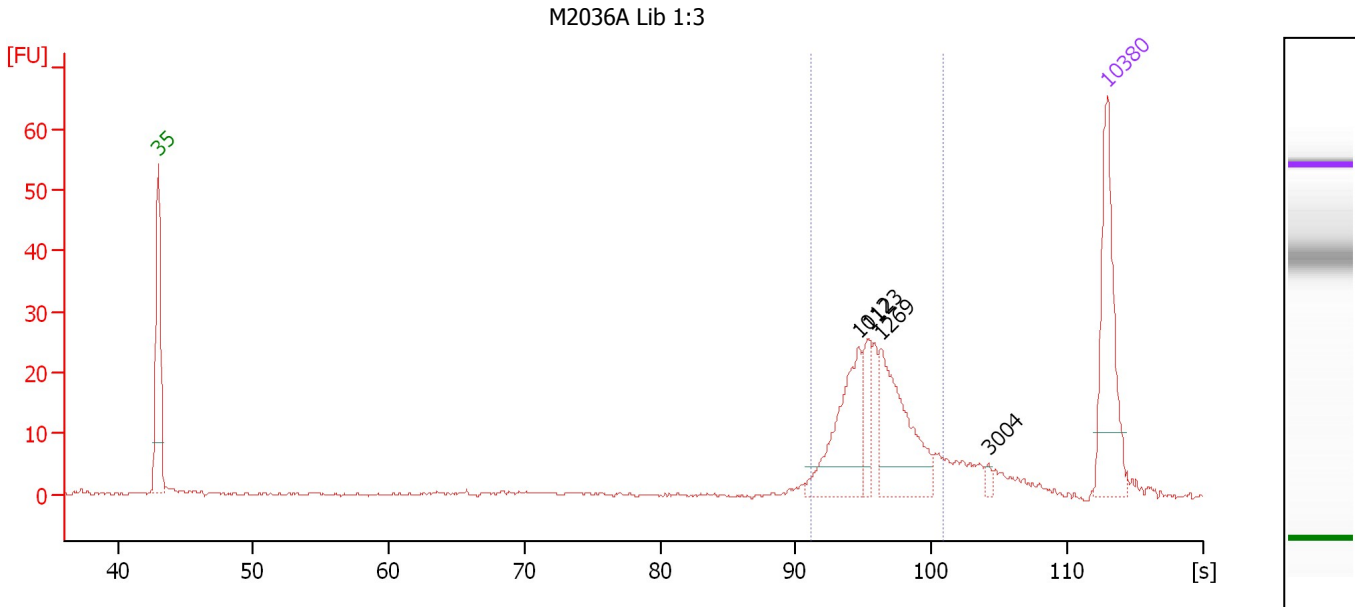
Region table for sample 5 : B2-R mRNA Lib 1:4

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color
64.10	91.09	398	12,358.3	3,008.79	2,764.2	95	23.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : M2036A Lib 1:3

Number of peaks found: 4 Corr. Area 1: 150.5
 Noise: 0.2

Peak table for sample 6 : M2036A Lib 1:3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	1,012	77.54	116.1	
3	1,123	26.02	35.1	
4	1,269	81.30	97.0	
5	3,004	3.53	1.8	
6	10,380	75.00	10.9	Upper Marker

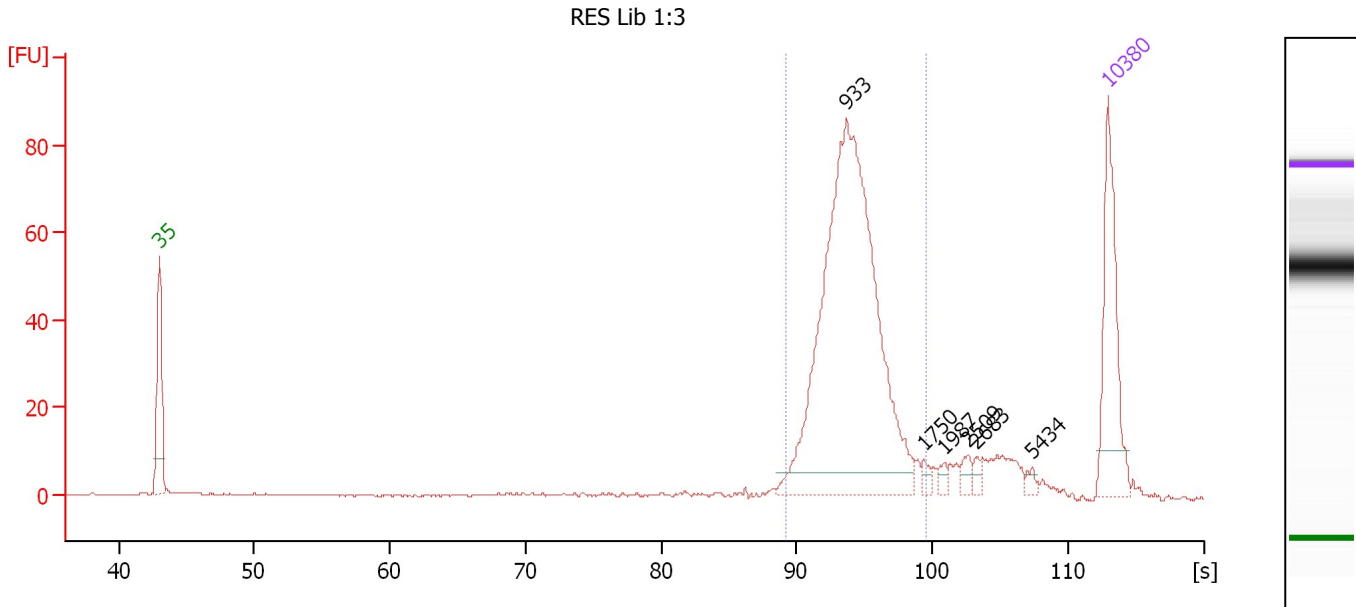
Region table for sample 6 : M2036A Lib 1:3

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
91.20	100.89	1,235	261.9	204.06	150.5	76	24.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : RES Lib 1:3

Number of peaks found: 6 Corr. Area 1: 433.0
 Noise: 0.2

Peak table for sample 7 : RES Lib 1:3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	933	454.75	738.8	
3	1,750	5.62	4.9	
4	1,987	5.05	3.9	
5	2,509	6.80	4.1	
6	2,683	4.64	2.6	
7	5,434	4.00	1.1	
8	10,380	75.00	10.9	Upper Marker

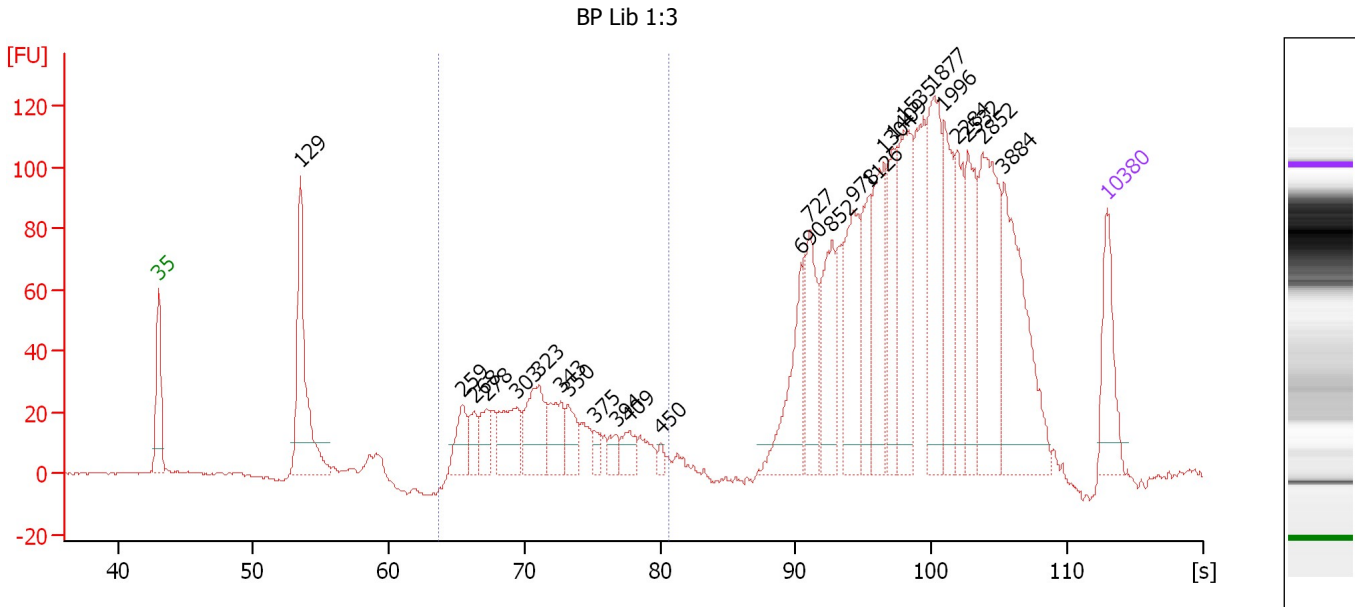
Region table for sample 7 : RES Lib 1:3

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
89.23	99.58	1,003	720.1	457.35	433.0	83	22.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 8 : BP Lib 1:3

Height Threshold [FU] : 10

Overall Results for sample 8 : BP Lib 1:3

Number of peaks found: 26 Corr. Area 1: 407.5
 Noise: 0.3

Peak table for sample 8 : BP Lib 1:3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	129	269.39	3,170.4	
3	259	57.27	335.0	
4	268	33.73	190.8	
5	278	46.24	251.7	
6	303	88.79	443.7	
7	323	105.56	494.6	
8	343	65.84	290.9	
9	350	44.54	192.8	
10	375	16.96	68.5	
11	394	20.30	78.1	
12	409	29.82	110.4	
13	450	10.27	34.6	
14	690	126.79	278.5	
15	727	116.06	241.9	
16	852	116.78	207.6	
17	978	136.06	210.9	
18	1,126	83.59	112.5	
19	1,304	128.19	148.9	
20	1,409	88.39	95.1	
21	1,535	145.42	143.5	
22	1,877	147.69	119.2	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...

... Peak table for sample 8 : BP Lib 1:3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
23	1,996	99.03	75.2	
24	2,284	82.13	54.5	
25	2,532	99.63	59.6	
26	2,852	177.29	94.2	
27	3,884	200.88	78.4	
28	10,380	75.00	10.9	Upper Marker

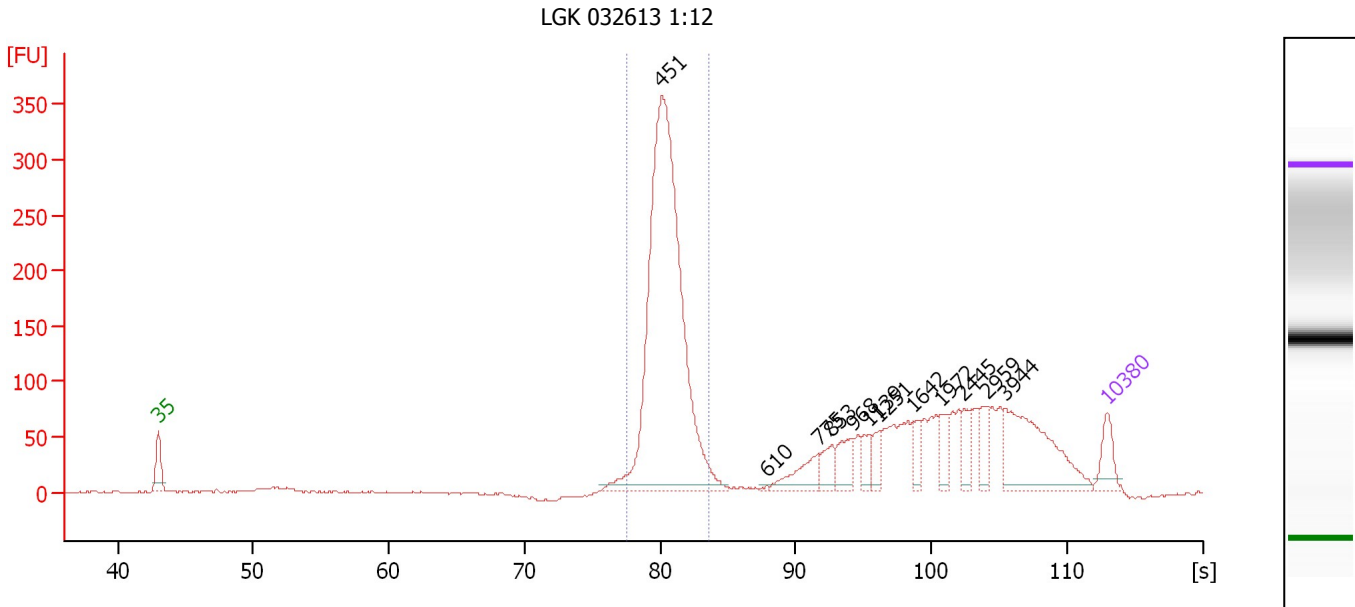
Region table for sample 8 : BP Lib 1:3

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area % of Total	Size distribution in CV [%]	Color
63.64	80.68	332	2,971.2	630.43	407.5	17	15.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : LGK 032613 1:12

Number of peaks found: 12 Corr. Area 1: 1,222.7
 Noise: 0.9

Peak table for sample 9 : LGK 032613 1:12

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	451	2,134.32	7,166.9	
3	610	5.05	12.5	
4	775	116.06	226.9	
5	853	72.03	127.9	
6	968	96.20	150.5	
7	1,139	53.87	71.7	
8	1,251	58.87	71.3	
9	1,642	53.08	49.0	
10	1,972	62.91	48.3	
11	2,445	65.52	40.6	
12	2,959	66.63	34.1	
13	3,944	362.17	139.1	
14	10,380	75.00	10.9	Upper Marker

Region table for sample 9 : LGK 032613 1:12

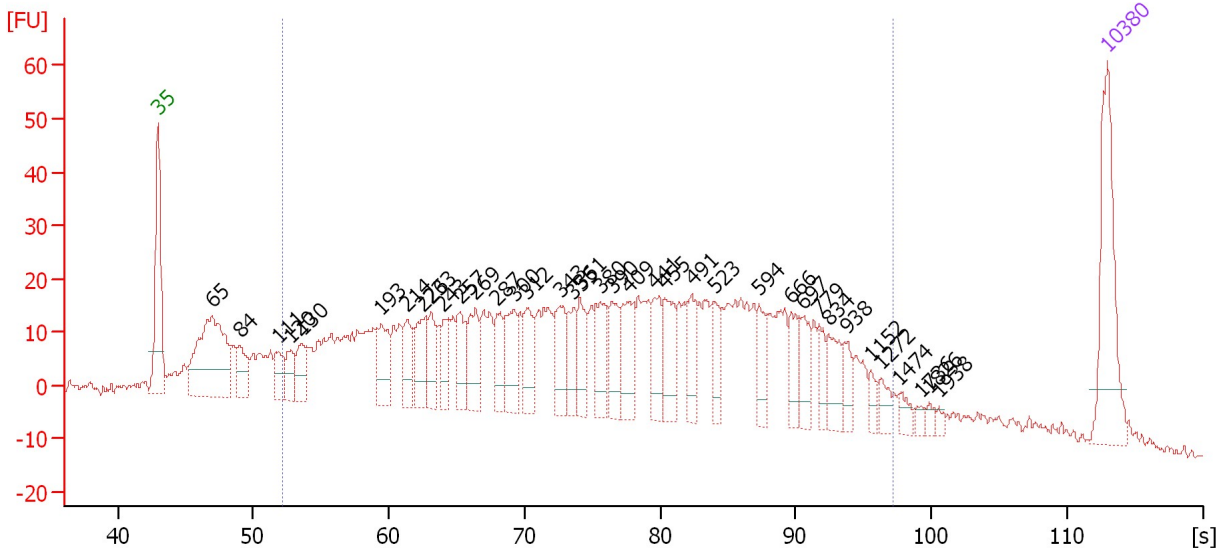
From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
77.49	83.68	457	6,838.9	2,063.50	1,222.7	49	4.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...

HLB Soni 1:6



Overall Results for sample 10 : HLB Soni 1:6

Number of peaks found: 37 Corr. Area 1: 1,092.1
 Noise: 0.8

Peak table for sample 10 : HLB Soni 1:6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	65	160.15	3,745.5	
3	84	34.94	633.4	
4	111	23.68	323.2	
5	120	26.99	340.1	
6	130	33.59	391.4	
7	193	43.01	337.3	
8	214	32.87	232.7	
9	226	41.83	280.3	
10	233	31.56	204.8	
11	243	30.22	188.1	
12	257	35.26	208.3	
13	269	43.20	243.2	
14	287	31.49	166.3	
15	300	42.35	213.9	
16	312	37.78	183.5	
17	343	38.87	172.0	
18	355	29.55	126.1	
19	361	34.21	143.5	
20	380	37.97	151.5	
21	390	35.33	137.1	
22	409	40.58	150.2	
23	441	31.70	108.8	
24	455	42.50	141.5	
25	491	25.38	78.4	
26	523	23.42	67.8	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad


Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Electropherogram Summary Continued ...

... Peak table for sample 10 : HLB Soni 1:6

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27	594	21.96	56.1	
28	666	25.75	58.6	
29	697	23.71	51.5	
30	779	17.83	34.7	
31	834	26.25	47.7	
32	938	14.91	24.1	
33	1,152	9.32	12.3	
34	1,272	10.76	12.8	
35	1,474	8.68	8.9	
36	1,736	4.70	4.1	
37	1,826	4.00	3.3	
38	1,938	3.15	2.5	
39	▶ 10,380	75.00	10.9	Upper Marker

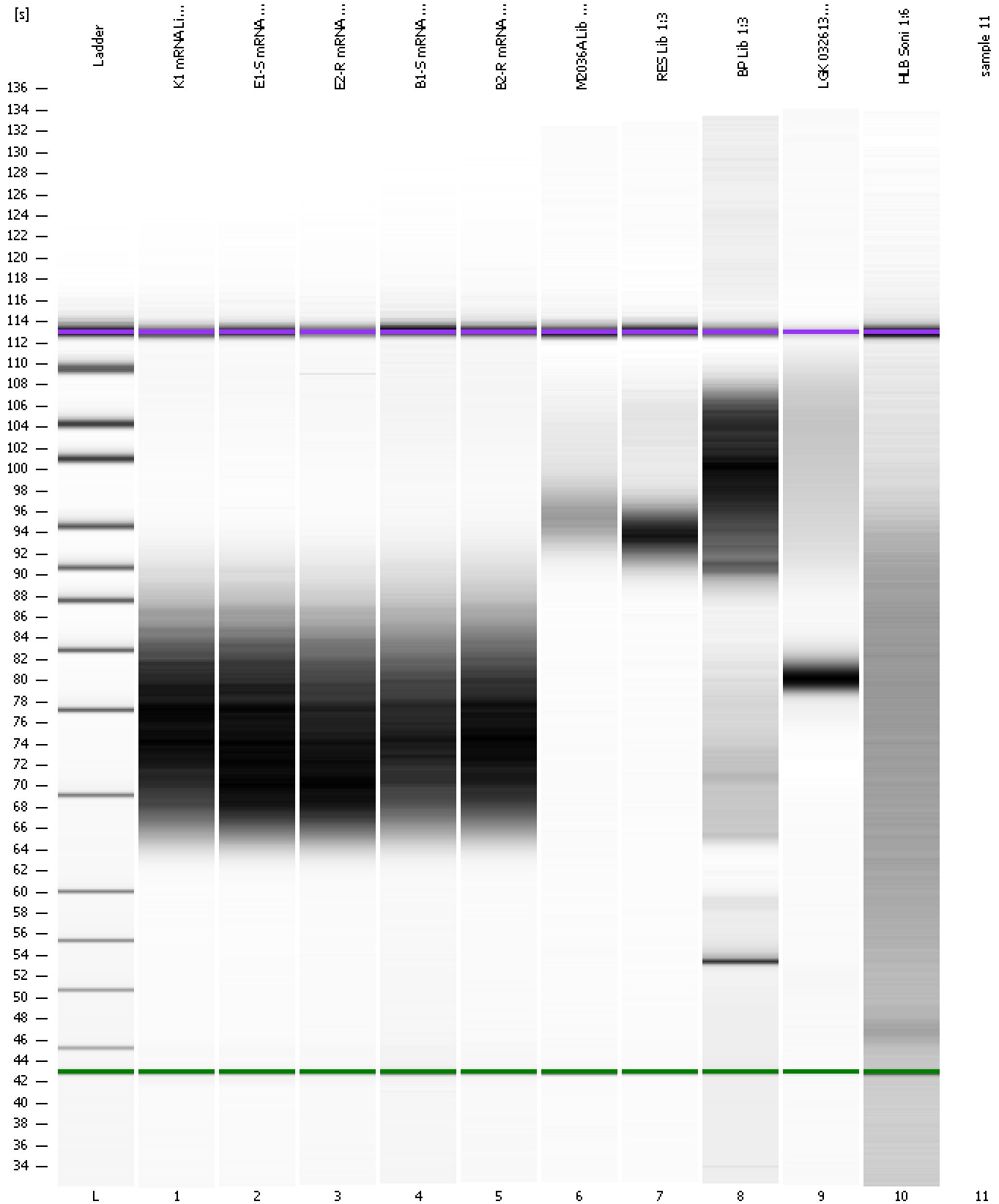
Region table for sample 10 : HLB Soni 1:6

From [s]	To [s]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area % of Total	Size distribution in CV [%]	Color
52.10	97.22	445	8,678.9	1,680.08	1,092.1 83	53.5	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
Modified: 3/26/2013 5:27:33 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
Modified: 3/26/2013 5:27:33 PM

Invalid Samples

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad

Created: 3/26/2013 4:44:28 PM
 Modified: 3/26/2013 5:27:33 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 11)		Instrument	Run		3/26/2013 5:22:55 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-03-26\2013-03-26_002.xad)		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/26/2013 4:44:33 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1