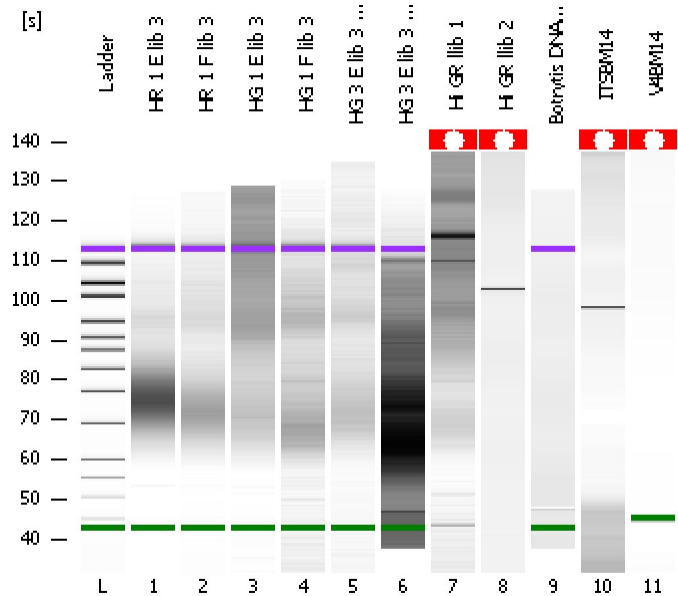


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 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

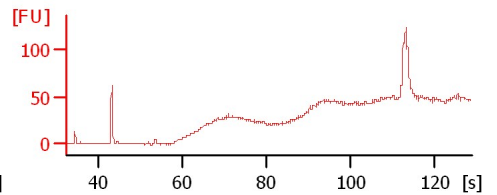
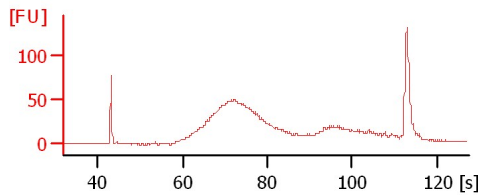
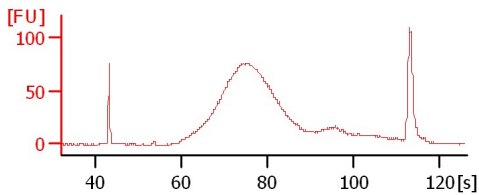
Chip Information:

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

HR 1 E lib 3

HR 1 F lib 3

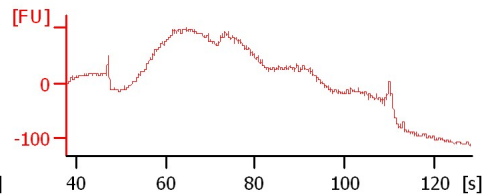
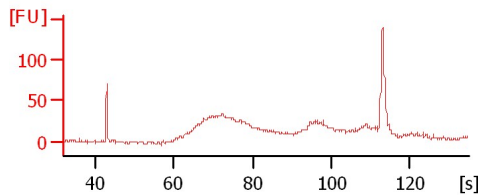
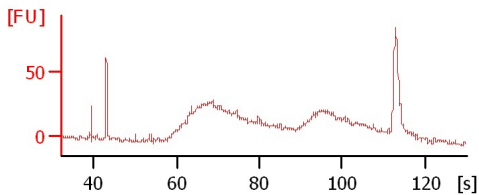
HG 1 E lib 3



HG 1 F lib 3

HG 3 E lib 3 (5)

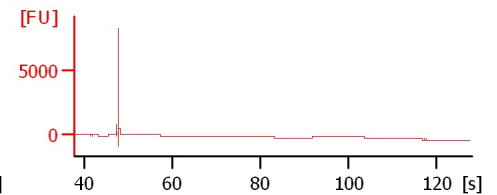
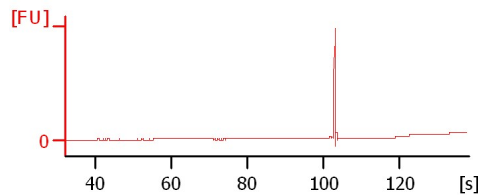
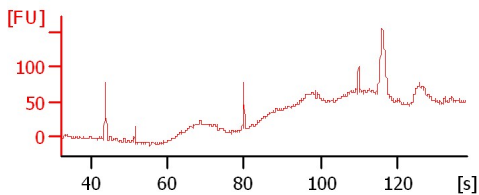
HG 3 E lib 3 (6)



Hi GR lib 1

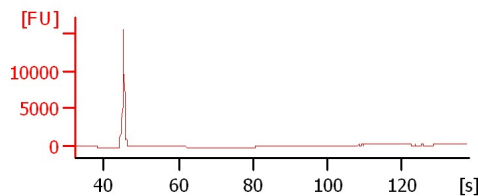
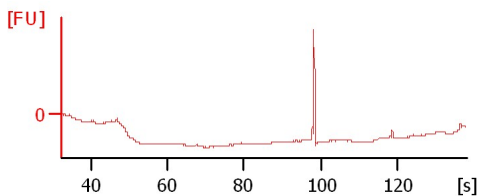
Hi GR lib 2

Botrytis DNA Lib 1



ITSBM14

V4BM14



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
HR 1 E lib 3		<input type="checkbox"/>	✓			
HR 1 F lib 3		<input type="checkbox"/>	✓			
HG 1 E lib 3		<input type="checkbox"/>	✓			
HG 1 F lib 3		<input type="checkbox"/>	✓			
HG 3 E lib 3 (5)	2 tubes with same name	<input type="checkbox"/>	✓			
HG 3 E lib 3 (6)	2 tubes with same name	<input type="checkbox"/>	✓			
Hi GR llib 1		<input type="checkbox"/>	✓			
Hi GR llib 2		<input type="checkbox"/>	✓			
Botrytis DNA Lib 1		<input type="checkbox"/>	✓			
ITSBM14		<input type="checkbox"/>	✓			
V4BM14		<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\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 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

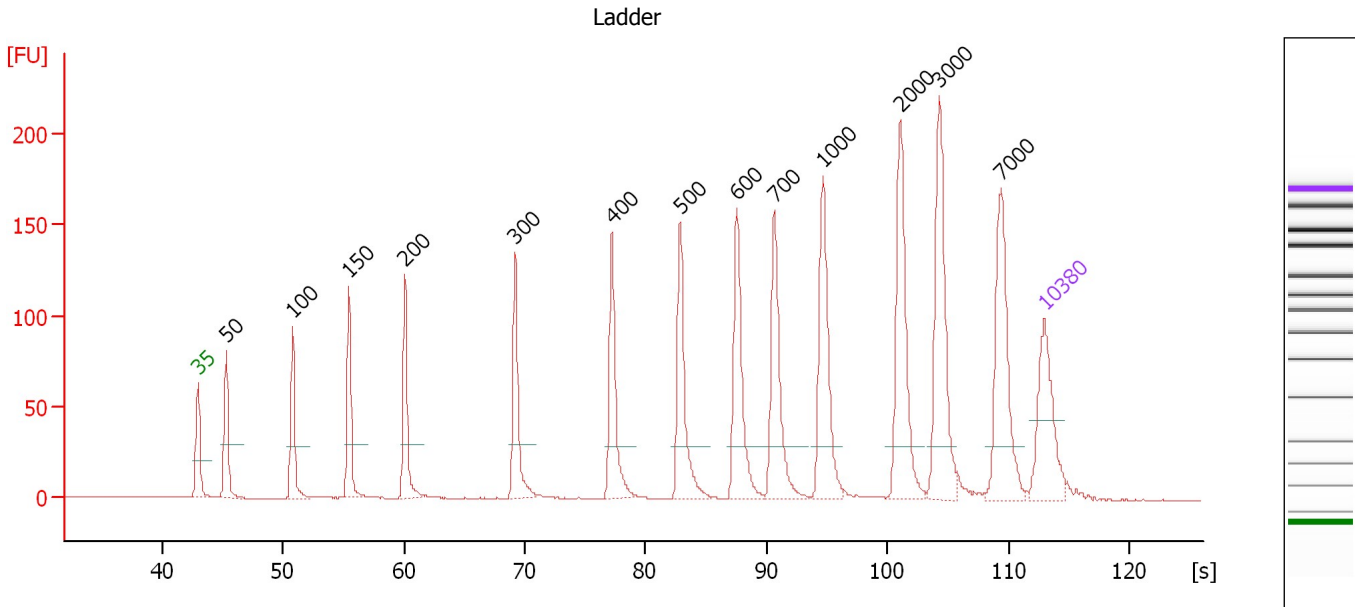
Ladder

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\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

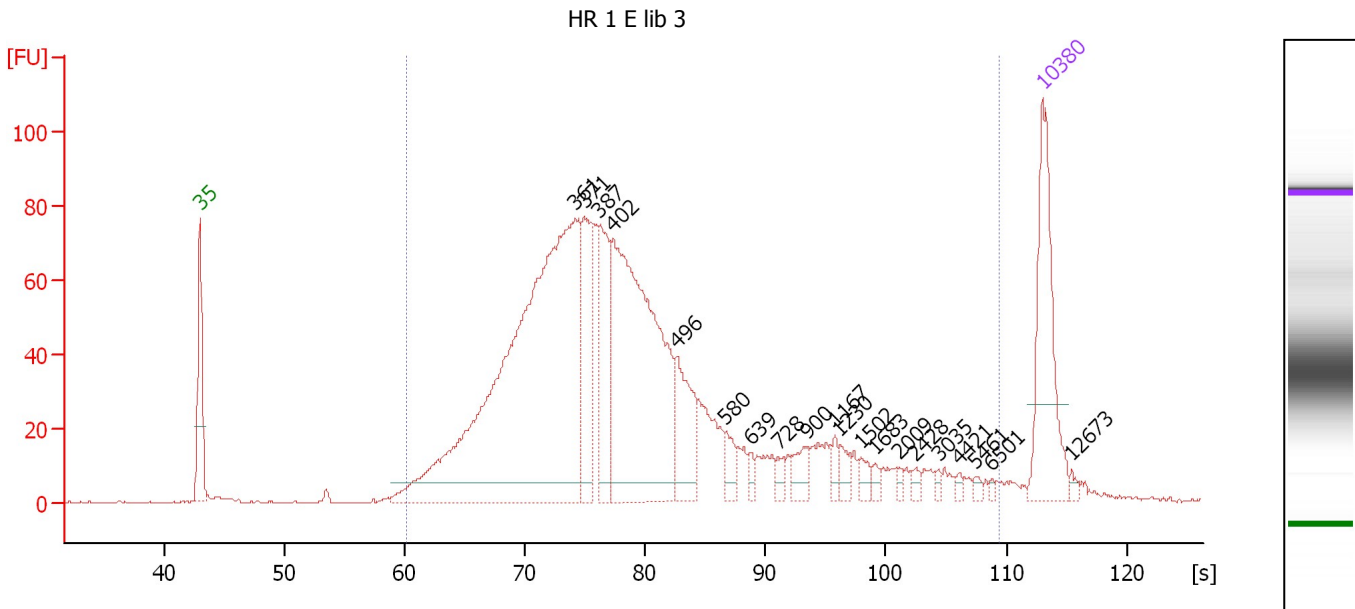
Peak table for Ladder

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\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : HR 1 E lib 3

Number of peaks found: 20 Corr. Area 1: 1,827.3
 Noise: 0.2

Peak table for sample 1 : HR 1 E lib 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	361	585.28	2,457.4	
3	371	79.99	326.3	
4	387	77.44	303.5	
5	402	295.35	1,114.5	
6	496	58.30	178.2	
7	580	13.90	36.3	
8	639	5.94	14.1	
9	728	6.59	13.7	
10	900	14.78	24.9	
11	1,167	7.02	9.1	
12	1,230	8.82	10.9	
13	1,502	7.45	7.5	
14	1,683	4.07	3.7	
15	2,009	2.91	2.2	
16	2,428	4.30	2.7	
17	3,035	2.77	1.4	
18	4,421	3.02	1.0	
19	5,461	2.76	0.8	
20	6,501	1.71	0.4	
21	10,380	75.00	10.9	Upper Marker
22	12,673	0.00	0.0	

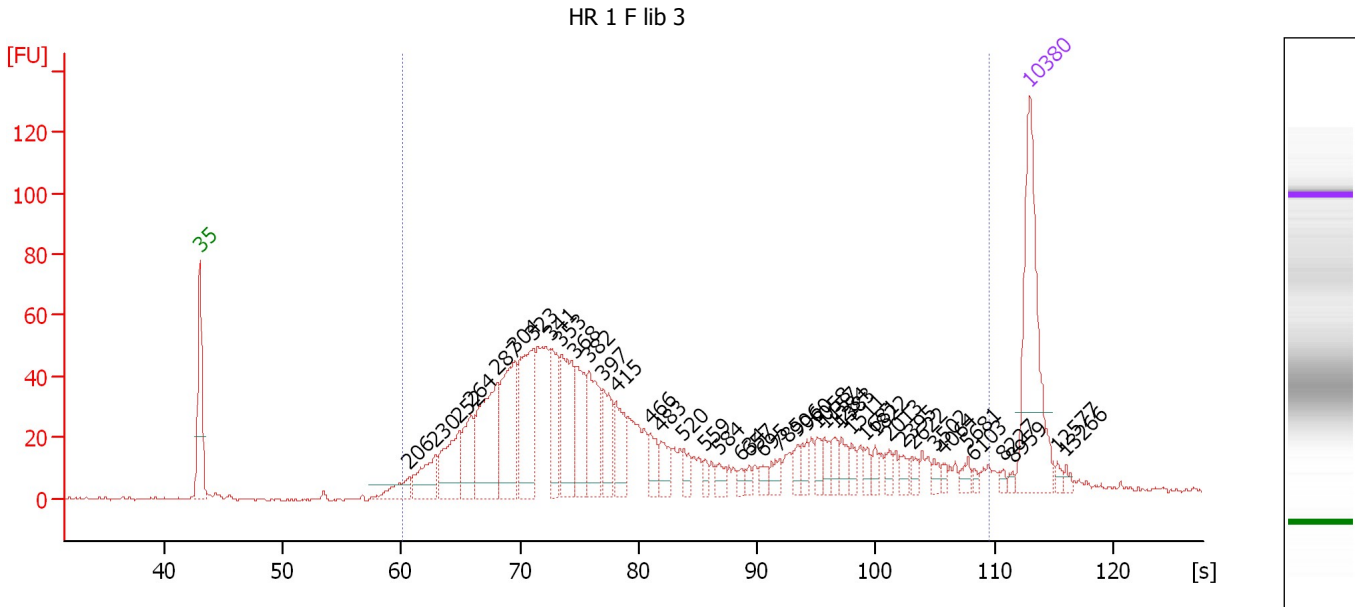
Region table for sample 1 : HR 1 E lib 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	613	6,987	1,827.3	98	100.0	1,397.98	5,499.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : HR 1 F lib 3

Number of peaks found: 42 Corr. Area 1: 1,282.4
 Noise: 0.4

Peak table for sample 2 : HR 1 F lib 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	206	18.17	133.6	
3	230	31.01	204.0	
4	252	44.17	265.5	
5	264	33.84	194.6	
6	287	78.32	413.1	
7	304	70.05	349.0	
8	323	70.79	331.7	
9	341	37.75	167.5	
10	353	54.91	235.5	
11	368	36.40	149.8	
12	382	45.44	180.4	
13	397	27.52	105.0	
14	415	26.19	95.6	
15	466	14.83	48.2	
16	483	14.88	46.6	
17	520	8.72	25.4	
18	559	5.00	13.6	
19	584	6.85	17.8	
20	634	3.86	9.2	
21	657	4.13	9.5	
22	695	5.88	12.8	
23	785	8.08	15.6	
24	906	6.28	10.5	
25	960	7.09	11.2	
26	1,058	6.97	10.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad


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 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 2 : HR 1 F lib 3

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27	1,157	8.10	10.6	
28	1,284	7.07	8.3	
29	1,383	7.76	8.5	
30	1,511	5.06	5.1	
31	1,681	4.86	4.4	
32	1,822	4.33	3.6	
33	2,013	4.03	3.0	
34	2,395	4.67	3.0	
35	2,622	3.38	2.0	
36	3,502	3.57	1.5	
37	4,064	2.67	1.0	
38	5,681	4.81	1.3	
39	6,103	2.06	0.5	
40	8,227	2.23	0.4	
41	8,959	1.80	0.3	
42	10,380	75.00	10.9	Upper Marker
43	12,577	0.00	0.0	
44	13,266	0.00	0.0	

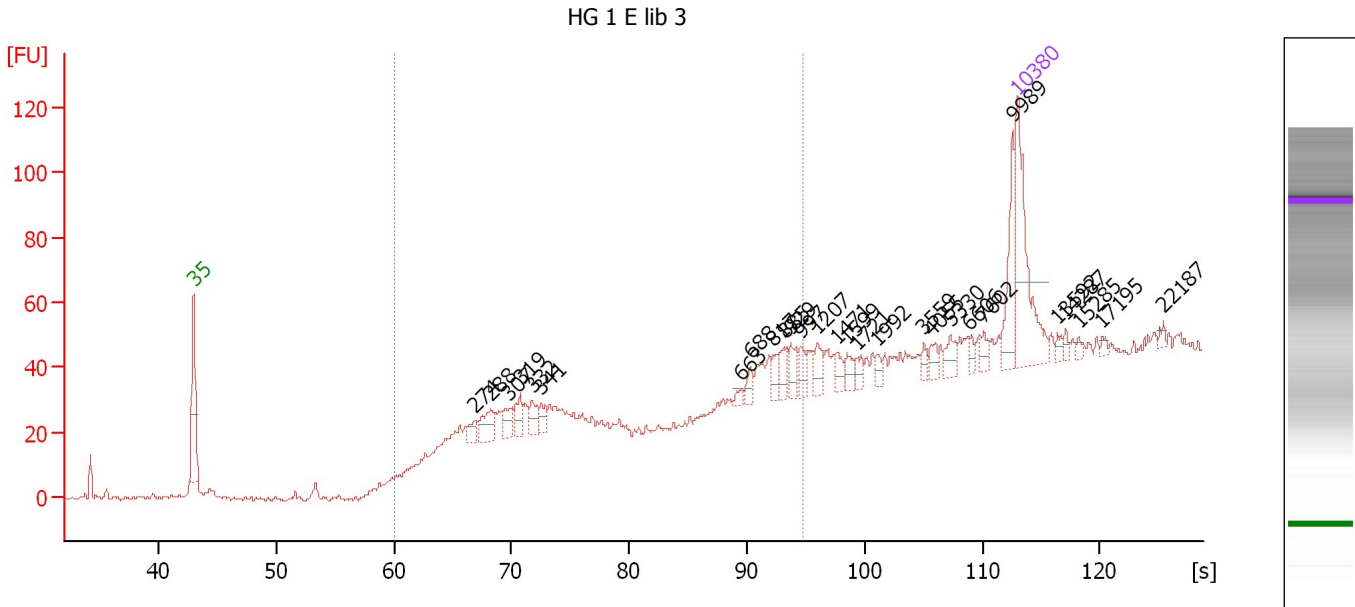
Region table for sample 2 : HR 1 F lib 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Color
200	782	7,136	1,282.4	97	100.0	880.67	3,539.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : HG 1 E lib 3

Number of peaks found: 28 Corr. Area 1: 237.2
 Noise: 0.8

Peak table for sample 3 : HG 1 E lib 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	274	10.46	57.9	
3	288	26.73	140.4	
4	307	16.81	83.0	
5	319	17.32	82.3	
6	332	13.45	61.4	
7	341	10.17	45.2	
8	663	5.25	12.0	
9	688	9.52	21.0	
10	817	10.56	19.6	
11	885	14.37	24.6	
12	929	14.05	22.9	
13	997	10.74	16.3	
14	1,207	12.80	16.1	
15	1,471	9.42	9.7	
16	1,599	8.83	8.4	
17	1,721	6.69	5.9	
18	1,992	5.20	4.0	
19	3,559	5.55	2.4	
20	4,055	7.21	2.7	
21	5,330	10.85	3.1	
22	6,606	5.53	1.3	
23	7,602	7.79	1.6	
24	9,989	37.50	5.7	
25	10,380	75.00	10.9	Upper Marker
26	13,592	0.00	0.0	


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...**... Peak table for sample 3 : HG 1 E lib 3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	14,287	0.00	0.0	
28	15,285	0.00	0.0	
29	17,195	0.00	0.0	
30	22,187	0.00	0.0	

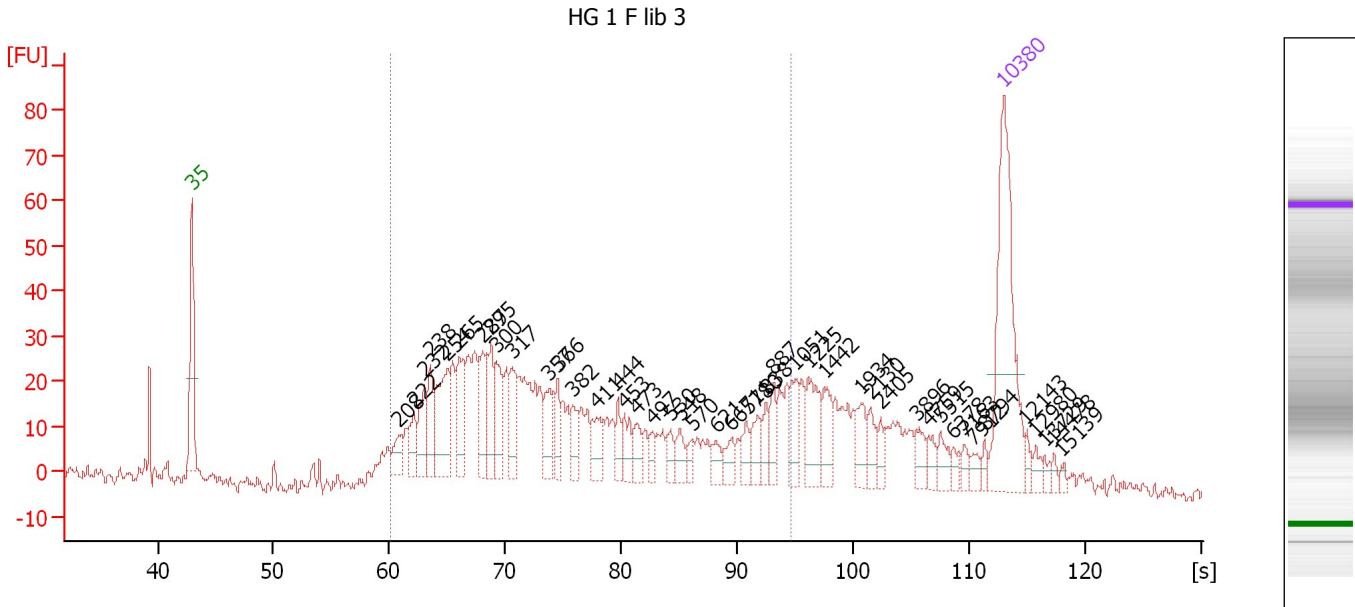
Region table for sample 3 : HG 1 E lib 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	540	1,000	237.2	43	47.6	305.49	1,177.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HG 1 F lib 3

Number of peaks found: 45 Corr. Area 1: 784.9
 Noise: 1.6

Peak table for sample 4 : HG 1 F lib 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	208	13.37	97.6	
3	222	15.16	103.4	
4	232	26.82	175.3	
5	238	21.53	137.1	
6	254	47.54	284.0	
7	265	34.23	195.9	
8	287	26.61	140.5	
9	295	26.18	134.6	
10	300	24.87	125.8	
11	317	23.21	110.9	
12	357	21.60	91.7	
13	366	14.73	61.0	
14	382	12.43	49.3	
15	411	16.59	61.1	
16	444	10.02	34.2	
17	453	13.27	44.4	
18	473	10.95	35.1	
19	497	6.93	21.1	
20	530	7.55	21.6	
21	548	10.21	28.2	
22	570	5.37	14.3	
23	621	7.39	18.0	
24	665	8.76	20.0	
25	711	9.65	20.6	
26	780	11.50	22.3	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad


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 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : HG 1 F lib 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	838	8.48	15.3	
28	887	11.70	20.0	
29	1,051	14.94	21.5	
30	1,225	23.71	29.3	
31	1,442	14.58	15.3	
32	1,934	11.26	8.8	
33	2,130	8.78	6.2	
34	2,405	5.30	3.3	
35	3,896	7.26	2.8	
36	4,759	6.05	1.9	
37	5,515	7.60	2.1	
38	6,378	3.94	0.9	
39	7,163	4.55	1.0	
40	7,912	4.59	0.9	
41	8,794	3.20	0.6	
42	10,380	75.00	10.9	Upper Marker
43	12,143	0.00	0.0	
44	12,980	0.00	0.0	
45	13,729	0.00	0.0	
46	14,478	0.00	0.0	
47	15,139	0.00	0.0	

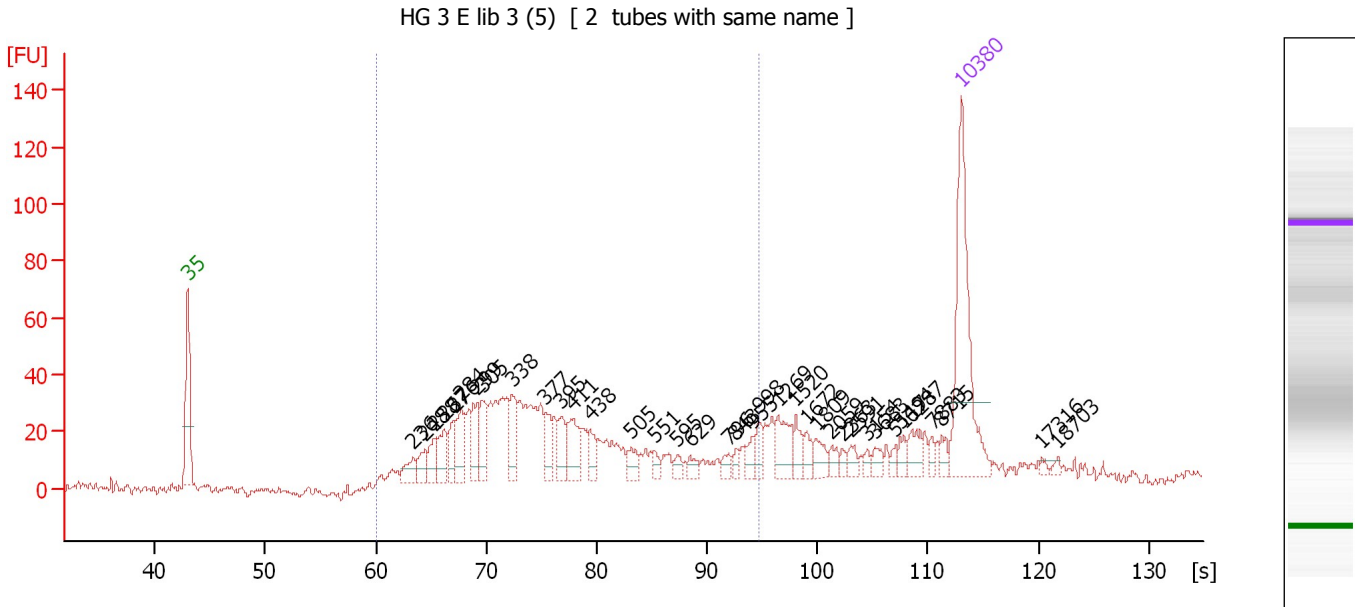
Region table for sample 4 : HG 1 F lib 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	425	1,000	784.9	71	47.2	730.87	3,290.7	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : HG 3 E lib 3 (5)

Number of peaks found: 37 Corr. Area 1: 709.0
 Noise: 1.2

Peak table for sample 5 : HG 3 E lib 3 (5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	236	12.90	82.7	
3	248	12.24	74.8	
4	258	14.51	85.2	
5	267	19.48	110.8	
6	276	15.51	85.1	
7	284	24.79	132.5	
8	299	24.04	121.6	
9	305	23.19	115.0	
10	338	22.62	101.3	
11	377	16.25	65.4	
12	395	16.44	63.1	
13	411	24.85	91.7	
14	438	9.08	31.4	
15	505	9.58	28.8	
16	551	5.15	14.2	
17	595	4.77	12.1	
18	629	5.91	14.2	
19	796	5.62	10.7	
20	843	4.21	7.6	
21	955	7.02	11.1	
22	998	7.81	11.9	
23	1,269	19.01	22.7	
24	1,520	9.07	9.0	
25	1,672	6.22	5.6	
26	1,809	8.93	7.5	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad


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 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 5 : HG 3 E lib 3 (5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	2,059	5.15	3.8	
28	2,363	2.96	1.9	
29	2,591	5.03	2.9	
30	3,154	3.50	1.7	
31	3,683	5.03	2.1	
32	5,117	3.33	1.0	
33	5,494	5.56	1.5	
34	6,287	10.50	2.5	
35	7,883	3.82	0.7	
36	8,715	4.92	0.9	
37	10,380	75.00	10.9	Upper Marker
38	17,316	0.00	0.0	
39	18,703	0.00	0.0	

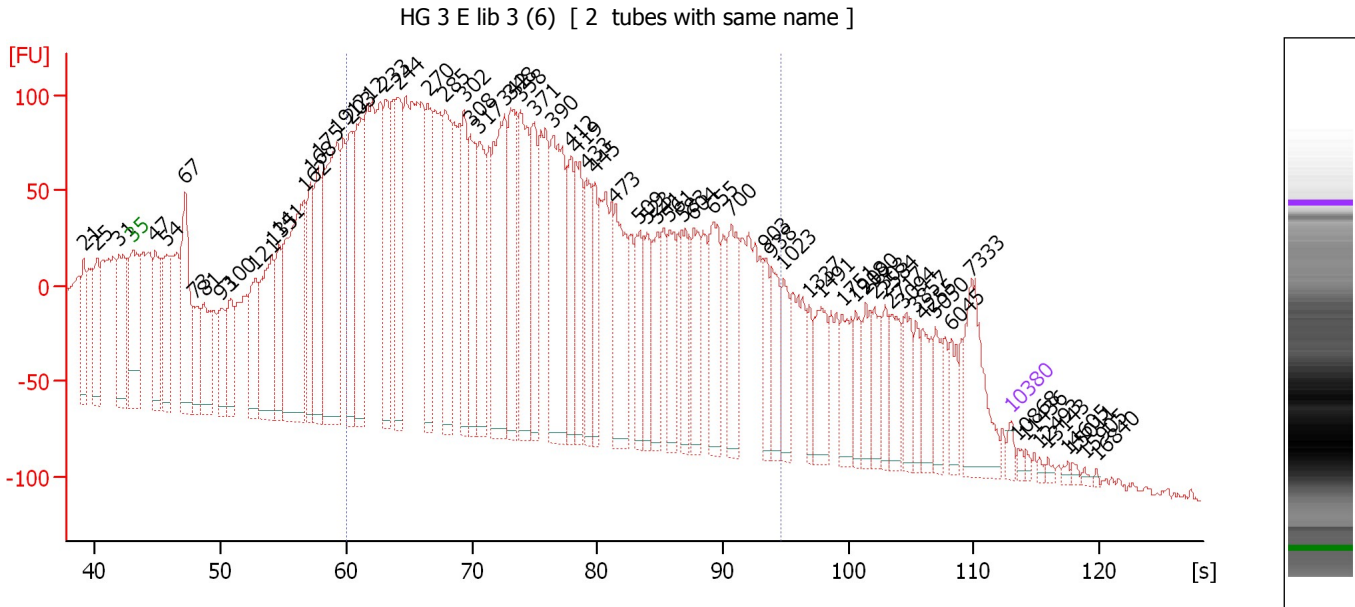
Region table for sample 5 : HG 3 E lib 3 (5)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	418	1,000	709.0	73	41.0	527.13	2,255.2	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : HG 3 E lib 3 (6)

Number of peaks found: 69 Corr. Area 1: 5,649.8
 Noise: 2.5

Peak table for sample 6 : HG 3 E lib 3 (6)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	21	0.00	0.0	
2	25	0.00	0.0	
3	31	0.00	0.0	
4	35	125.00	5,411.3	Lower Marker
5	47	1,290.16	41,903.5	
6	54	1,070.41	30,119.5	
7	67	2,086.91	47,173.0	
8	73	756.02	15,724.8	
9	81	1,099.66	20,671.9	
10	93	759.23	12,368.3	
11	100	894.27	13,604.1	
12	121	1,219.25	15,319.4	
13	135	1,873.11	21,056.4	
14	141	897.52	9,631.3	
15	162	3,113.60	29,053.7	
16	168	1,197.40	10,774.4	
17	175	1,332.12	11,543.0	
18	191	3,155.41	25,026.2	
19	203	1,228.77	9,167.5	
20	212	1,820.56	13,014.2	
21	233	1,358.42	8,835.1	
22	244	1,336.32	8,308.4	
23	270	1,134.68	6,371.4	
24	285	1,476.90	7,845.1	
25	302	1,075.23	5,398.0	
26	308	1,220.33	5,999.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad


Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 6 : HG 3 E lib 3 (6)

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27	317	1,169.60	5,594.5	
28	342	2,041.87	9,035.4	
29	348	1,288.31	5,613.3	
30	358	1,567.62	6,636.5	
31	371	967.48	3,948.5	
32	390	1,922.01	7,457.8	
33	412	749.69	2,755.4	
34	419	891.14	3,222.0	
35	433	740.21	2,587.4	
36	445	617.99	2,105.0	
37	473	1,133.58	3,632.5	
38	509	636.40	1,893.5	
39	523	475.35	1,377.1	
40	541	638.78	1,788.0	
41	561	599.49	1,617.8	
42	583	548.39	1,424.1	
43	604	672.00	1,685.8	
44	655	826.66	1,912.8	
45	700	822.31	1,779.0	
46	903	344.90	578.6	
47	938	492.23	795.0	
48	1,023	378.49	560.5	
49	1,337	234.52	265.8	
50	1,491	474.46	482.3	
51	1,751	373.82	323.5	
52	1,918	286.31	226.2	
53	2,090	278.44	201.9	
54	2,303	307.51	202.3	
55	2,584	244.30	143.3	
56	2,717	361.40	201.5	
57	3,094	322.51	157.9	
58	3,857	188.85	74.2	
59	4,255	305.99	109.0	
60	5,050	263.49	79.1	
61	6,045	243.32	61.0	
62	7,333	795.79	164.4	
63	10,380	75.00	10.9	Upper Marker
64	10,868	0.00	0.0	
65	11,436	0.00	0.0	
66	12,493	0.00	0.0	
67	13,143	0.00	0.0	
68	14,605	0.00	0.0	
69	15,011	0.00	0.0	
70	15,905	0.00	0.0	
71	16,840	0.00	0.0	

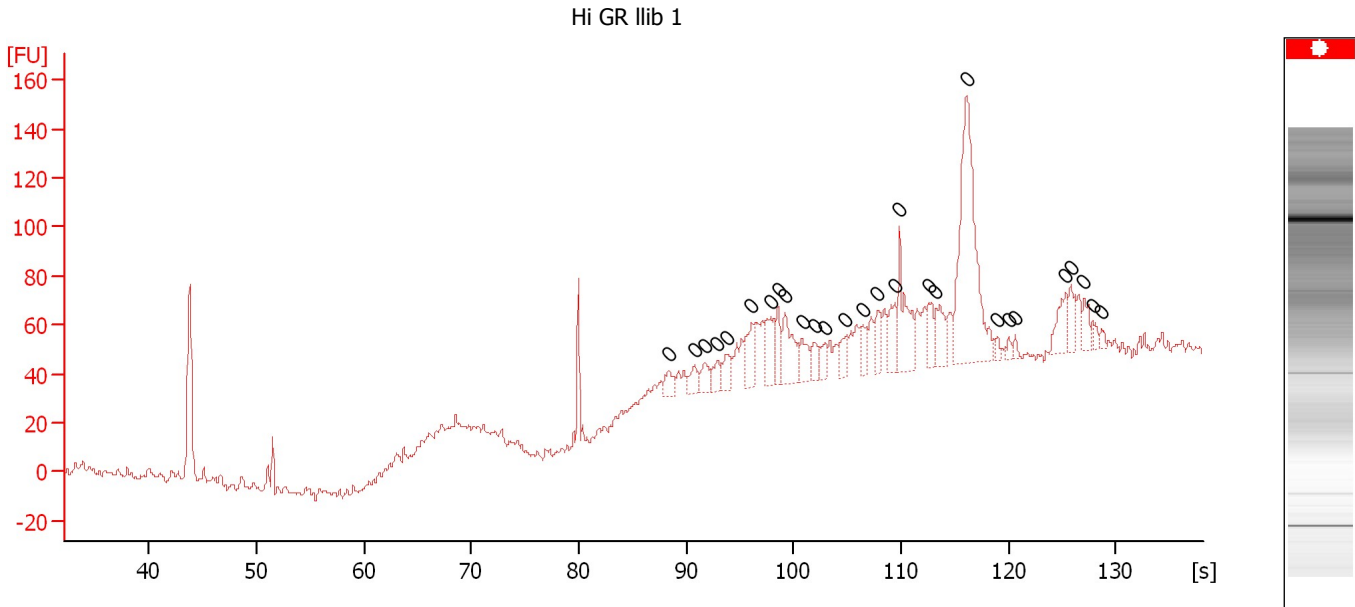
Region table for sample 6 : HG 3 E lib 3 (6)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Color
200	424	1,000	5,649.8	69	43.5	34,886.91	157,461.6	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Hi GR llib 1

Number of peaks found: 0 Noise: 2.0

Peak table for sample 7 : Hi GR llib 1

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	0	0.00	0.0	
2	0	0.00	0.0	
3	0	0.00	0.0	
4	0	0.00	0.0	
5	0	0.00	0.0	
6	0	0.00	0.0	
7	0	0.00	0.0	
8	0	0.00	0.0	
9	0	0.00	0.0	
10	0	0.00	0.0	
11	0	0.00	0.0	
12	0	0.00	0.0	
13	0	0.00	0.0	
14	0	0.00	0.0	
15	0	0.00	0.0	
16	0	0.00	0.0	
17	0	0.00	0.0	
18	0	0.00	0.0	
19	0	0.00	0.0	
20	0	0.00	0.0	
21	0	0.00	0.0	
22	0	0.00	0.0	
23	0	0.00	0.0	
24	0	0.00	0.0	
25	0	0.00	0.0	
26	0	0.00	0.0	
27	0	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

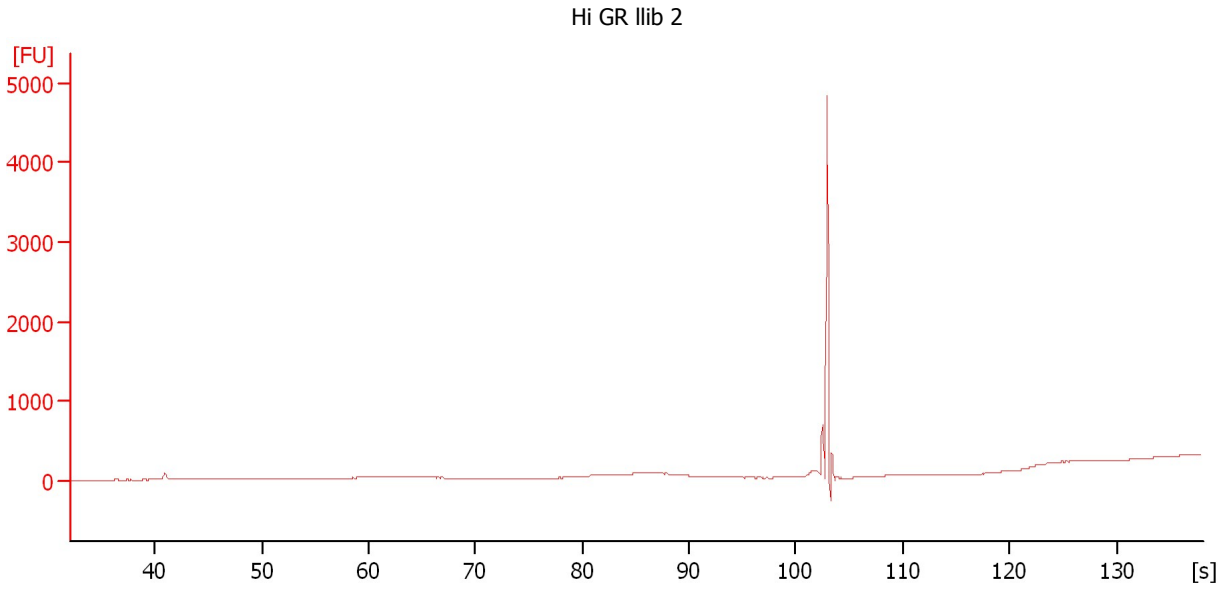
... Peak table for sample 7 : Hi GR Ilib 1

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations
28	0	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



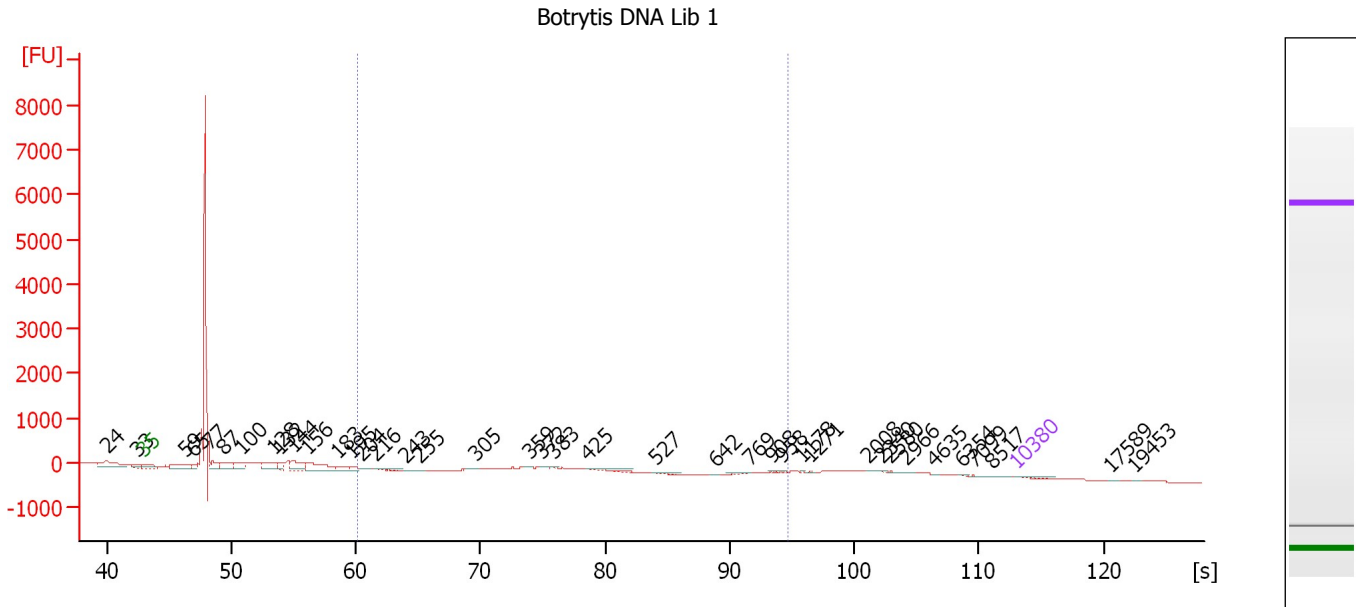
Overall Results for sample 8 : Hi GR Ilib 2

Noise: 3.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
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Electropherogram Summary Continued ...



Overall Results for sample 9 : Botrytis DNA Lib 1

Number of peaks found: 39 Corr. Area 1: 1,219.7
 Noise: 5.4

Peak table for sample 9 : Botrytis DNA Lib 1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	24	0.00	0.0	
2	33	0.00	0.0	
3	35	125.00	5,411.3	Lower Marker
4	59	8,591.30	219,946.8	
5	65	3,559.41	83,490.6	
6	77	7,487.33	147,398.9	
7	87	9,424.54	164,106.2	
8	100	7,487.81	113,712.5	
9	128	10,014.44	118,512.9	
10	132	4,581.81	52,536.2	
11	144	9,467.38	99,608.5	
12	156	13,210.50	127,983.0	
13	183	3,868.56	32,007.7	
14	195	2,513.22	19,516.8	
15	204	94.87	704.9	
16	216	163.44	1,144.0	
17	243	59.81	372.9	
18	255	58.37	346.7	
19	305	58.94	292.7	
20	359	50.01	211.2	
21	372	69.27	282.4	
22	383	69.37	274.6	
23	425	395.35	1,407.9	
24	527	144.38	415.3	
25	642	46.28	109.2	
26	769	129.73	255.7	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad


Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 9 : Botrytis DNA Lib 1

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27	908	61.90	103.3	
28	958	51.18	80.9	
29	1,178	23.76	30.6	
30	1,271	29.72	35.4	
31	2,008	29.64	22.4	
32	2,340	41.40	26.8	
33	2,580	55.32	32.5	
34	2,966	24.31	12.4	
35	4,635	20.31	6.6	
36	6,354	54.81	13.1	
37	7,099	39.97	8.5	
38	8,517	49.18	8.7	
39	10,380	75.00	10.9	Upper Marker
40	17,589	0.00	0.0	
41	19,453	0.00	0.0	

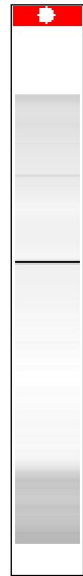
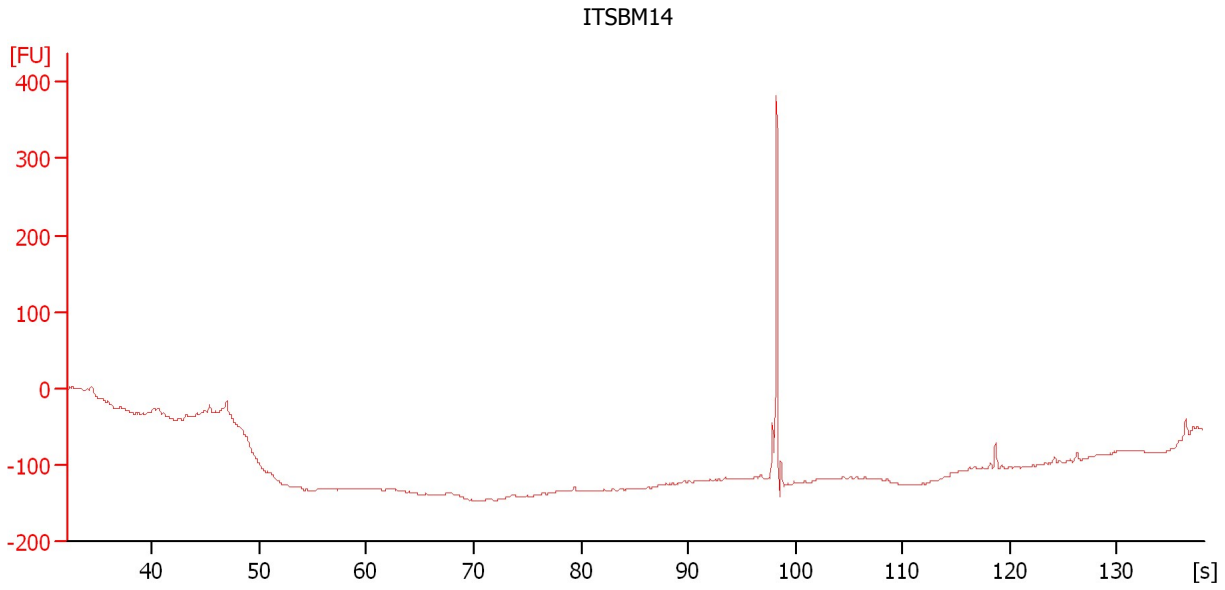
Region table for sample 9 : Botrytis DNA Lib 1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Color
200	515	1,000	1,219.7	14	42.3	17,658.58	61,987.9	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

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Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : ITSBM14

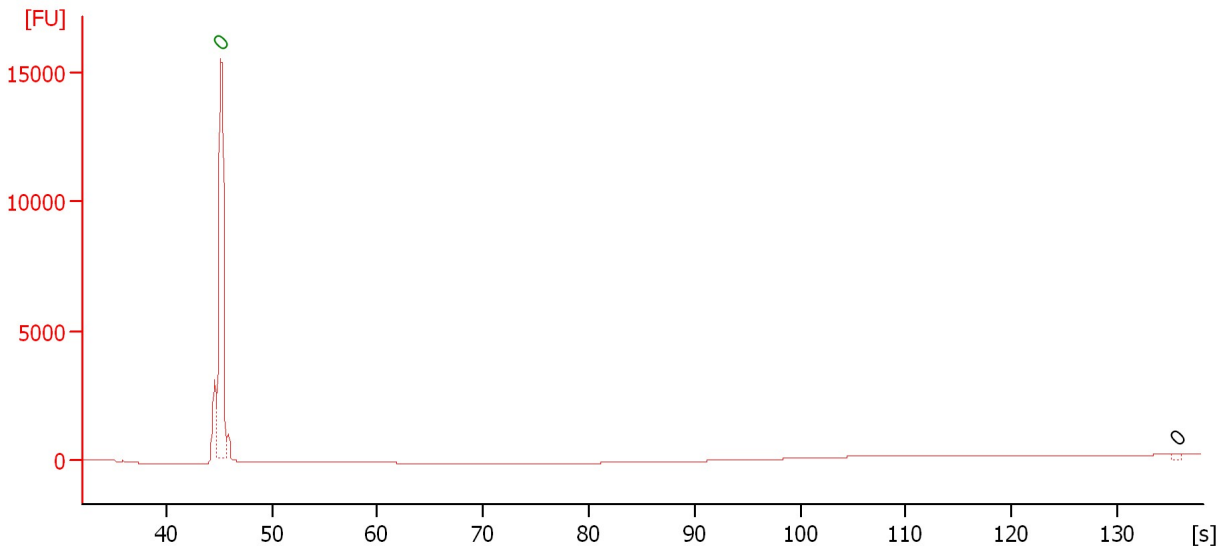
Noise: 1.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Electropherogram Summary Continued ...

V4BM14



Overall Results for sample 11 : V4BM14

Number of peaks found: 1 Noise: 1.1

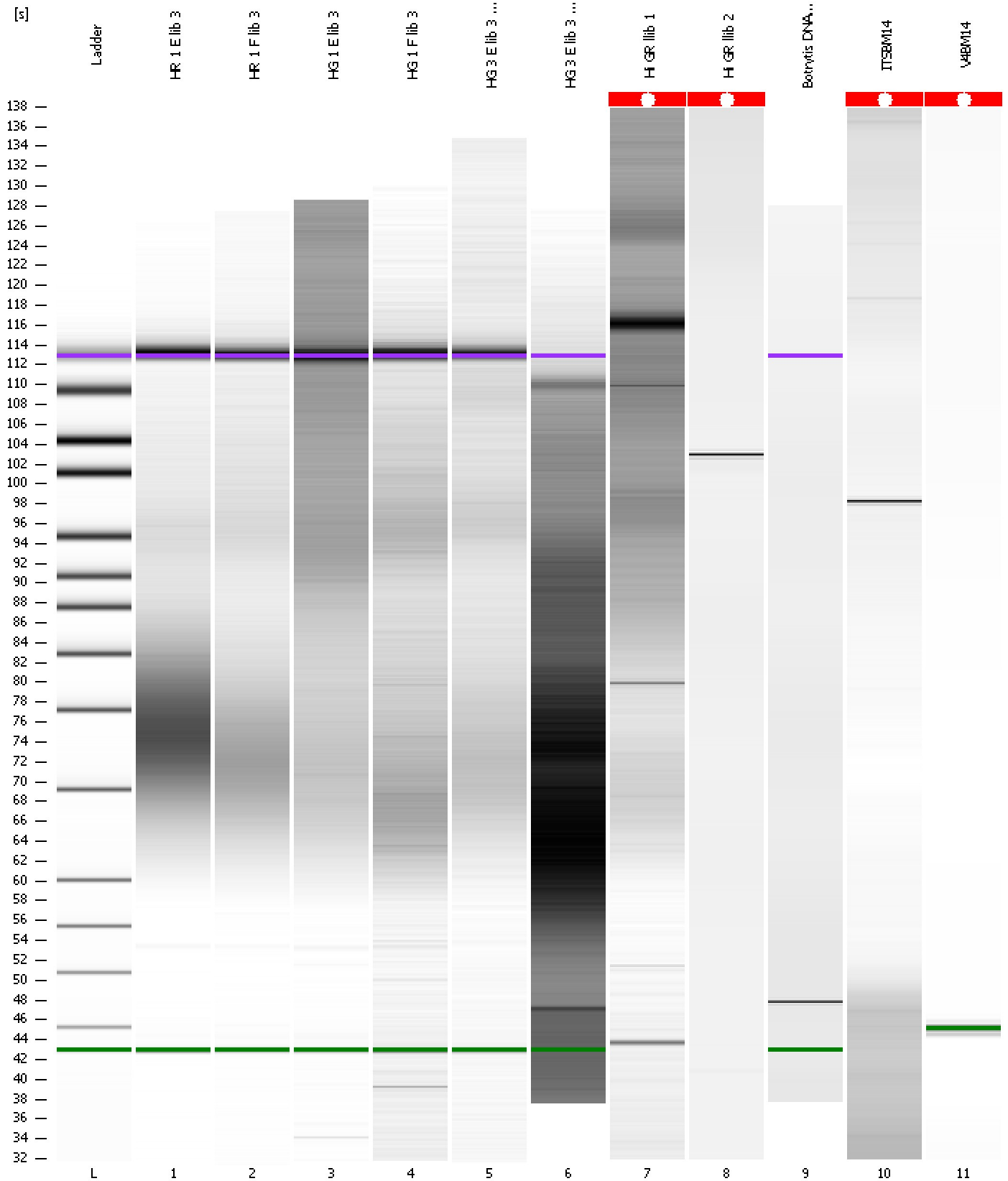
Peak table for sample 11 : V4BM14

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	0	0.00	0.0	Lower Marker
2	0	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
Modified: 11/25/2014 1:37:30 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad

Created: 11/25/2014 12:56:09 PM
 Modified: 11/25/2014 1:37:30 PM

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		11/25/2014 1:37:27 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Instrument error occurred on port 1, Optical signal too high (1605h)	559	Instrument	Run	Sample 11	11/25/2014 1:35:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-11-25\2014-11-25_005.xad)		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/25/2014 12:56:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1