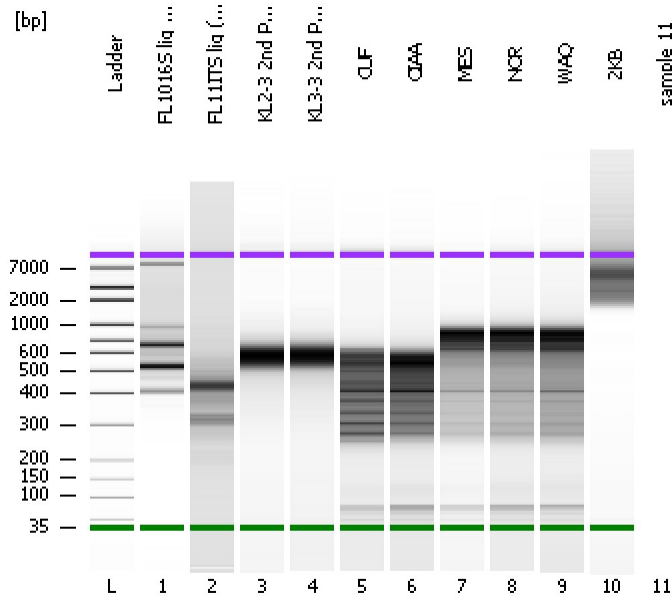


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Electrophoresis File Run Summary



Instrument Information:

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

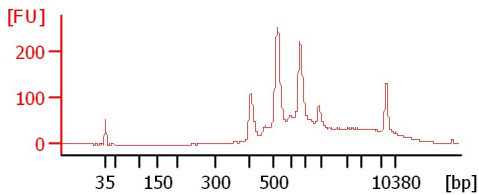
Assay Information:

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

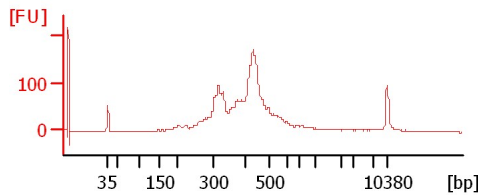
Chip Information:

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

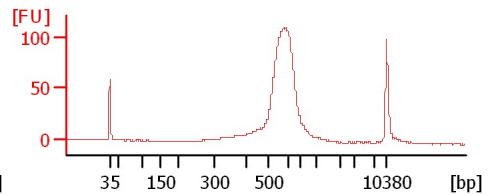
FL1016S lig (1:3)



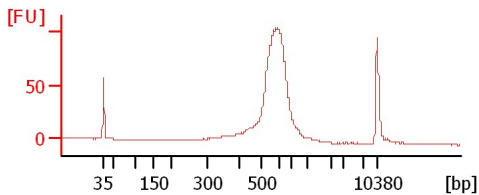
FL111TS lig (1:3)



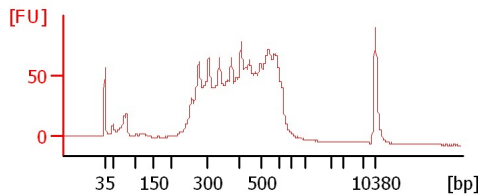
KL2-3 2nd PCR (1:90)



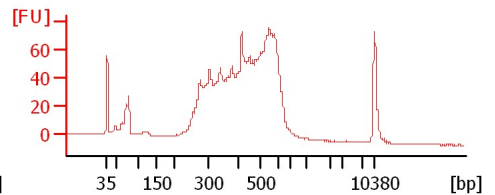
KL3-3 2nd PCR (1:90)



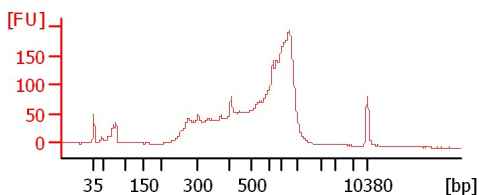
CUF



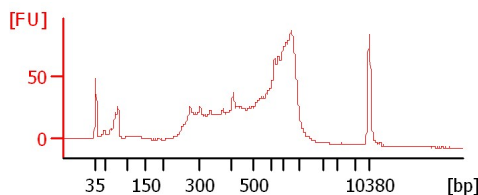
CIAA



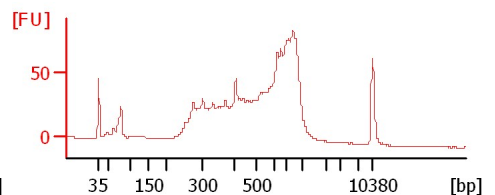
MES



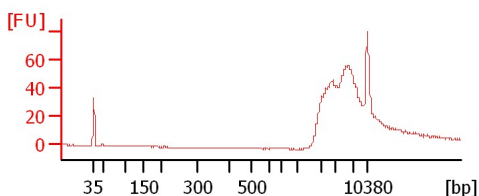
NOR



WAQ



2KB



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
FL1016S lig (1:3)		<input type="checkbox"/>	✓			
FL11ITS lig (1:3)		<input type="checkbox"/>	✓			
KL2-3 2nd PCR (1:90)		<input type="checkbox"/>	✓			
KL3-3 2nd PCR (1:90)		<input type="checkbox"/>	✓			
CUF		<input type="checkbox"/>	✓			
CIAA		<input type="checkbox"/>	✓			
MES		<input type="checkbox"/>	✓			
NOR		<input type="checkbox"/>	✓			
WAQ		<input type="checkbox"/>	✓			
2KB		<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:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

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

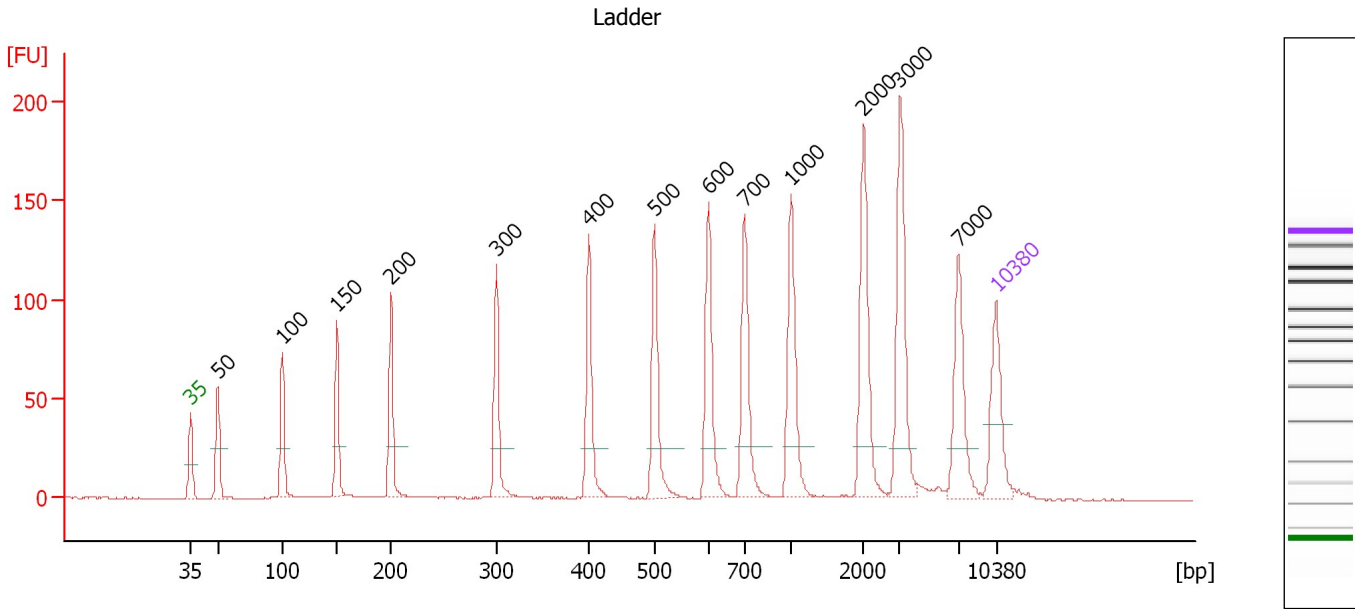
Ladder

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:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

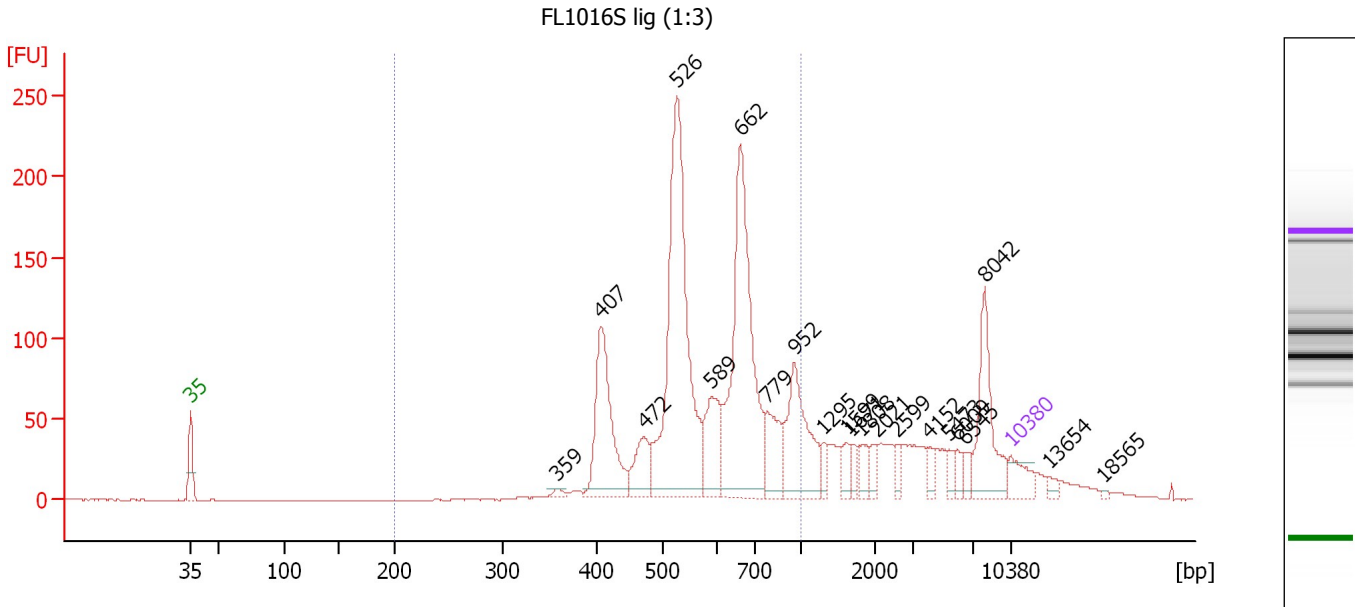
Peak table for Ladder

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	45.36
3	100	150.00	2,272.7	Ladder Peak	50.97
4	150	150.00	1,515.2	Ladder Peak	55.69
5	200	150.00	1,136.4	Ladder Peak	60.37
6	300	150.00	757.6	Ladder Peak	69.55
7	400	150.00	568.2	Ladder Peak	77.61
8	500	150.00	454.5	Ladder Peak	83.30
9	600	150.00	378.8	Ladder Peak	87.98
10	700	150.00	324.7	Ladder Peak	91.13
11	1,000	150.00	227.3	Ladder Peak	95.16
12	2,000	150.00	113.6	Ladder Peak	101.46
13	3,000	150.00	75.8	Ladder Peak	104.61
14	7,000	150.00	32.5	Ladder Peak	109.71
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : FL1016S lig (1:3)

Number of peaks found: 21 Corr. Area 1: 1,691.4
 Noise: 0.2

Peak table for sample 1 : FL1016S lig (1:3)


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	359	18.13	76.5		74.31
3	407	524.98	1,954.8		78.00
4	472	161.58	519.1		81.69
5	526	1,314.79	3,789.2		84.51
6	589	241.29	620.7		87.47
7	662	1,015.48	2,325.5		89.92
8	779	175.02	340.2		92.20
9	952	373.70	594.7		94.52
10	1,295	40.12	46.9		97.02
11	1,599	48.96	46.4		98.94
12	1,671	36.49	33.1		99.39
13	1,808	54.42	45.6		100.26
14	2,021	39.88	29.9		101.53
15	2,599	34.21	19.9		103.35
16	4,152	31.62	11.5		106.08
17	5,473	34.69	9.6		107.77
18	6,009	29.45	7.4		108.45
19	6,545	34.82	8.1		109.13
20	8,042	273.70	51.6		110.72
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,654	0.00	0.0		116.19
23	18,565	0.00	0.0		120.96

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...

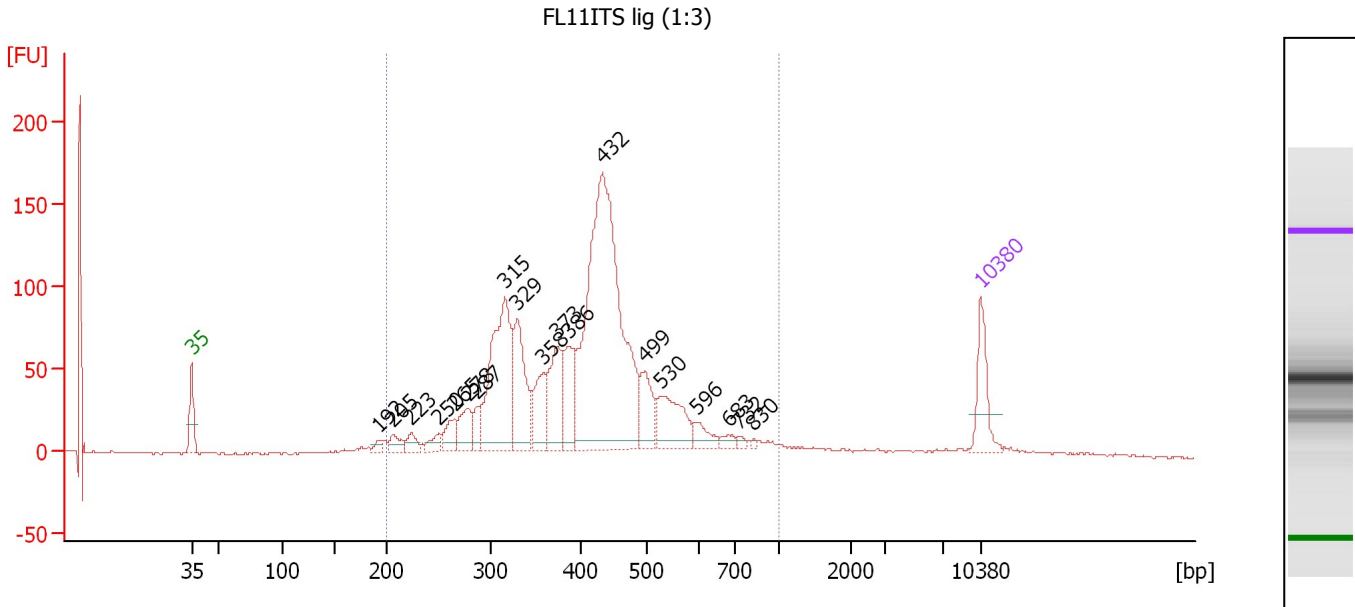
... Region table for sample 1 : FL1016S lig (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	601	1,691.4	10,238.0	 3,783.64	70	24.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : FL11ITS lig (1:3)

Number of peaks found: 19 Corr. Area 1: 1,939.2
 Noise: 0.5

Peak table for sample 2 : FL11ITS lig (1:3)

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	192	14.94	118.0		59.61
3	205	32.19	237.8		60.84
4	223	28.46	193.6		62.45
5	250	28.53	172.9		64.96
6	265	50.83	290.6		66.33
7	278	73.10	398.6		67.52
8	287	35.58	188.1		68.32
9	315	392.78	1,887.2		70.78
10	329	190.04	875.6		71.87
11	358	99.00	419.4		74.19
12	373	143.57	582.5		75.47
13	386	107.85	422.9		76.51
14	432	958.54	3,360.0		79.44
15	499	91.70	278.6		83.23
16	530	130.05	372.0		84.70
17	596	35.51	90.4		87.77
18	683	16.12	35.7		90.61
19	732	8.29	17.2		91.56
20	830	3.69	6.7		92.89
21	10,380	75.00	10.9	Upper Marker	113.00

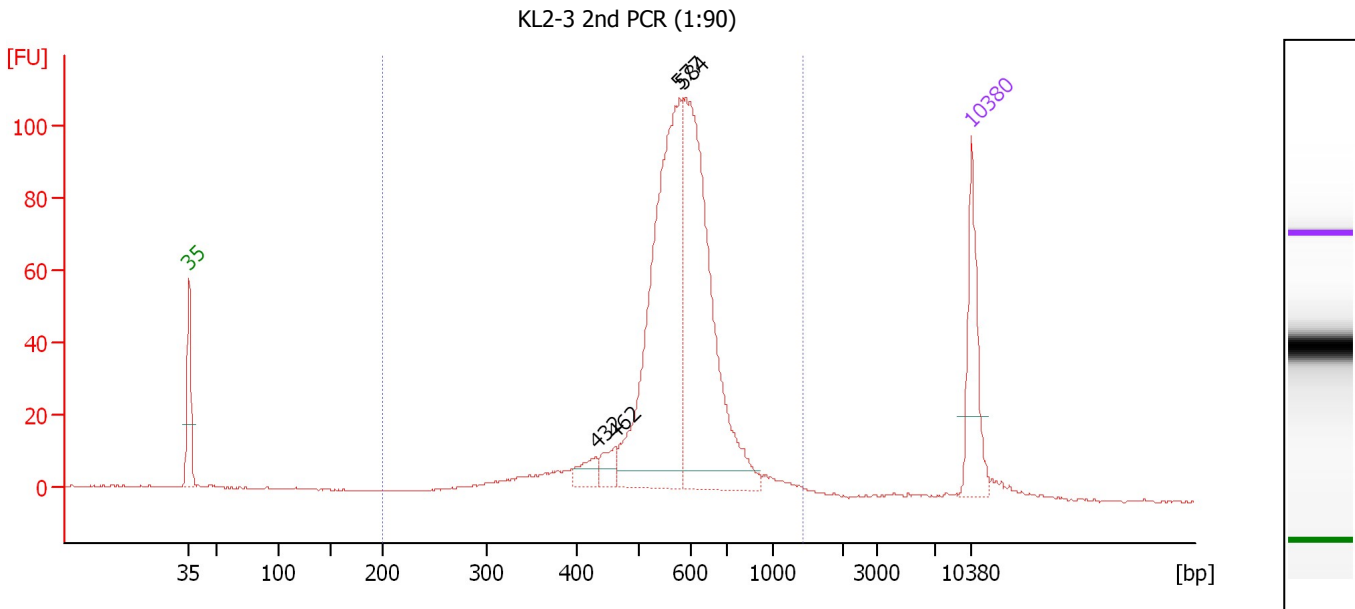
Region table for sample 2 : FL11ITS lig (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	416	1,939.2	10,377.5	2,602.25	89	26.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : KL2-3 2nd PCR (1:90)

Number of peaks found: 4 Corr. Area 1: 919.0
 Noise: 0.2

Peak table for sample 3 : KL2-3 2nd PCR (1:90)

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	432	23.60	82.9		79.41
3	462	24.22	79.5		81.13
4	577	448.38	1,177.8		86.90
5	584	421.61	1,093.9		87.23
6	10,380	75.00	10.9	Upper Marker	113.00

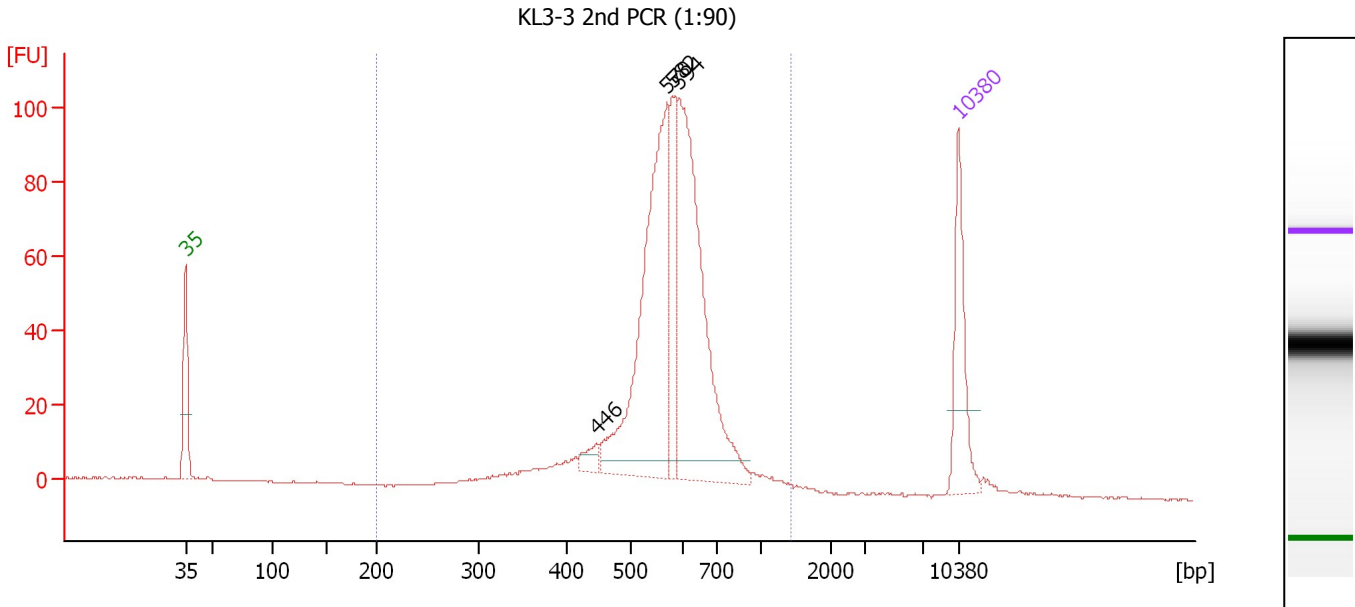
Region table for sample 3 : KL2-3 2nd PCR (1:90)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,440	580	919.0	2,958.3	1,070.56	96	20.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : KL3-3 2nd PCR (1:90)

Number of peaks found: 4 Corr. Area 1: 875.3
 Noise: 0.2

Peak table for sample 4 : KL3-3 2nd PCR (1:90)

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	446	18.08	61.5		80.22
3	570	357.66	950.2		86.59
4	582	96.67	251.8		87.12
5	594	349.62	891.8		87.70
6	10,380	75.00	10.9	Upper Marker	113.00

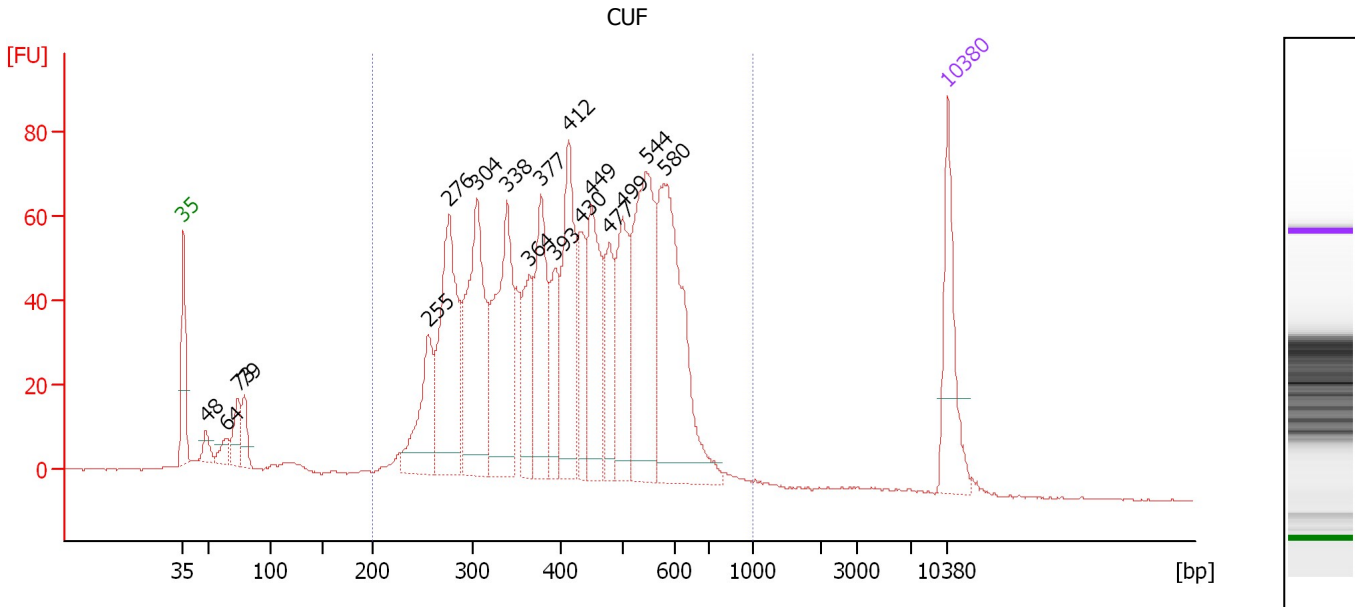
Region table for sample 4 : KL3-3 2nd PCR (1:90)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,411	582	875.3	2,752.3	1,002.73	97	19.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : CUF

Number of peaks found: 18 Corr. Area 1: 1,818.8
 Noise: 0.2

Peak table for sample 5 : CUF

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	48	19.84	625.9		45.05
3	64	20.93	497.3		46.91
4	73	44.35	916.2		47.98
5	79	34.45	660.7		48.61
6	255	107.22	636.0		65.45
7	276	223.48	1,226.1		67.36
8	304	227.14	1,130.7		69.90
9	338	199.02	891.4		72.63
10	364	81.11	337.9		74.68
11	377	128.80	517.6		75.75
12	393	65.53	252.8		77.02
13	412	157.32	578.5		78.29
14	430	71.31	251.3		79.32
15	449	125.13	422.4		80.39
16	477	71.69	227.6		82.00
17	499	116.24	352.6		83.27
18	544	208.13	579.5		85.37
19	580	249.36	651.8		87.03
20	10,380	75.00	10.9	Upper Marker	113.00

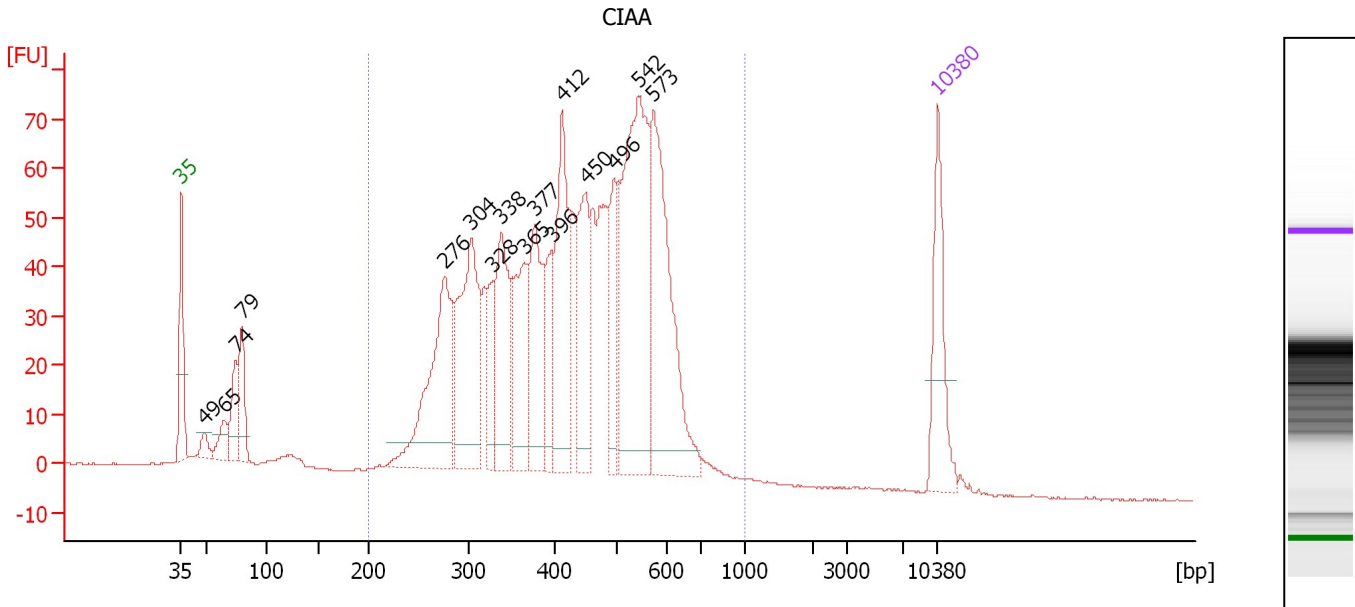
Region table for sample 5 : CUF

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	428	1,818.8	8,676.1	2,182.99	92	27.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : CIAA

Number of peaks found: 16 Corr. Area 1: 1,590.4
 Noise: 0.2

Peak table for sample 6 : CIAA

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	49	18.36	570.4		45.17
3	65	33.10	776.8		47.00
4	74	56.91	1,168.3		48.03
5	79	72.45	1,387.9		48.62
6	276	229.56	1,259.0		67.37
7	304	217.86	1,087.2		69.84
8	328	52.32	241.6		71.81
9	338	141.07	631.5		72.65
10	365	111.23	461.2		74.82
11	377	123.91	497.9		75.76
12	396	63.77	244.0		77.28
13	412	161.78	595.4		78.27
14	450	118.18	398.1		80.44
15	496	77.92	238.2		83.06
16	542	321.38	898.2		85.28
17	573	247.88	655.8		86.71
18	10,380	75.00	10.9	Upper Marker	113.00

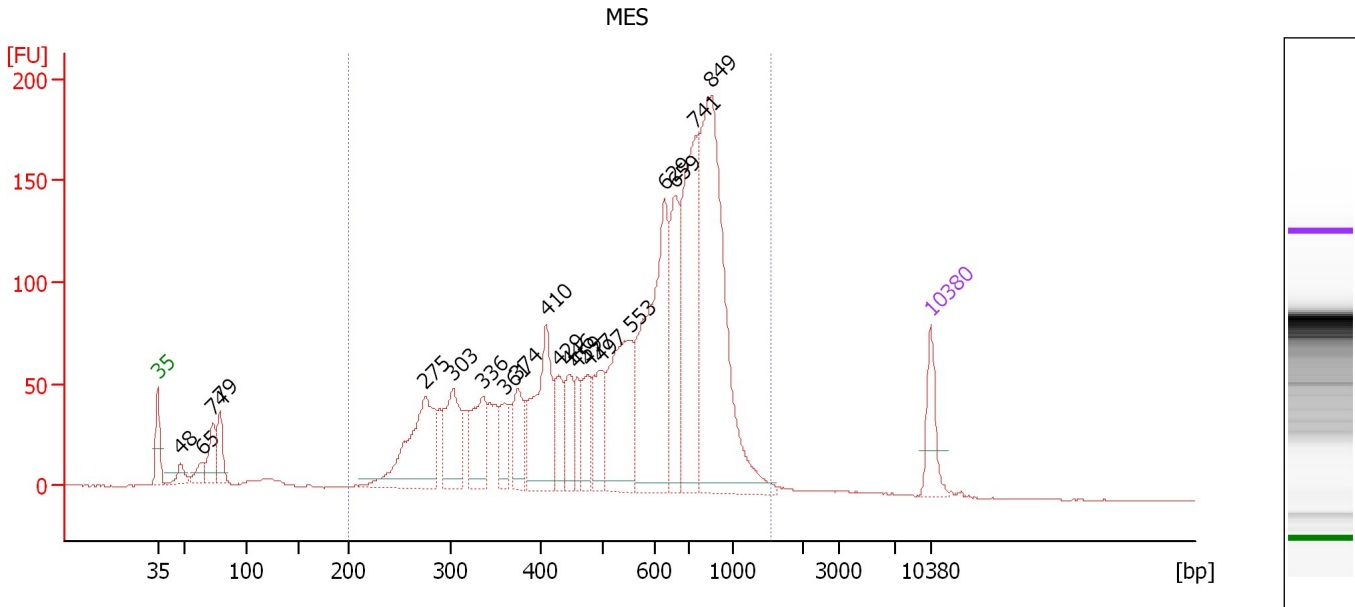
Region table for sample 6 : CIAA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	438	1,590.4	8,834.2	2,297.31	90	26.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : MES

Number of peaks found: 20 Corr. Area 1: 2,801.6
 Noise: 0.1

Peak table for sample 7 : MES

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	48	41.83	1,319.8		45.05
3	65	47.23	1,107.1		47.00
4	74	98.57	2,030.4		48.00
5	79	94.73	1,808.5		48.65
6	275	338.42	1,867.1		67.22
7	303	192.14	961.5		69.77
8	336	164.25	740.0		72.47
9	361	71.63	301.1		74.42
10	374	105.83	428.6		75.52
11	410	270.50	999.7		78.18
12	429	95.47	336.9		79.28
13	446	94.71	321.8		80.23
14	459	66.85	220.6		80.98
15	477	92.90	295.3		81.98
16	497	121.90	371.7		83.13
17	553	302.57	829.1		85.78
18	629	503.61	1,214.0		88.88
19	659	212.51	488.8		89.83
20	741	414.98	848.5		91.68
21	849	765.48	1,366.1		93.14
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...

... Region table for sample 7 :

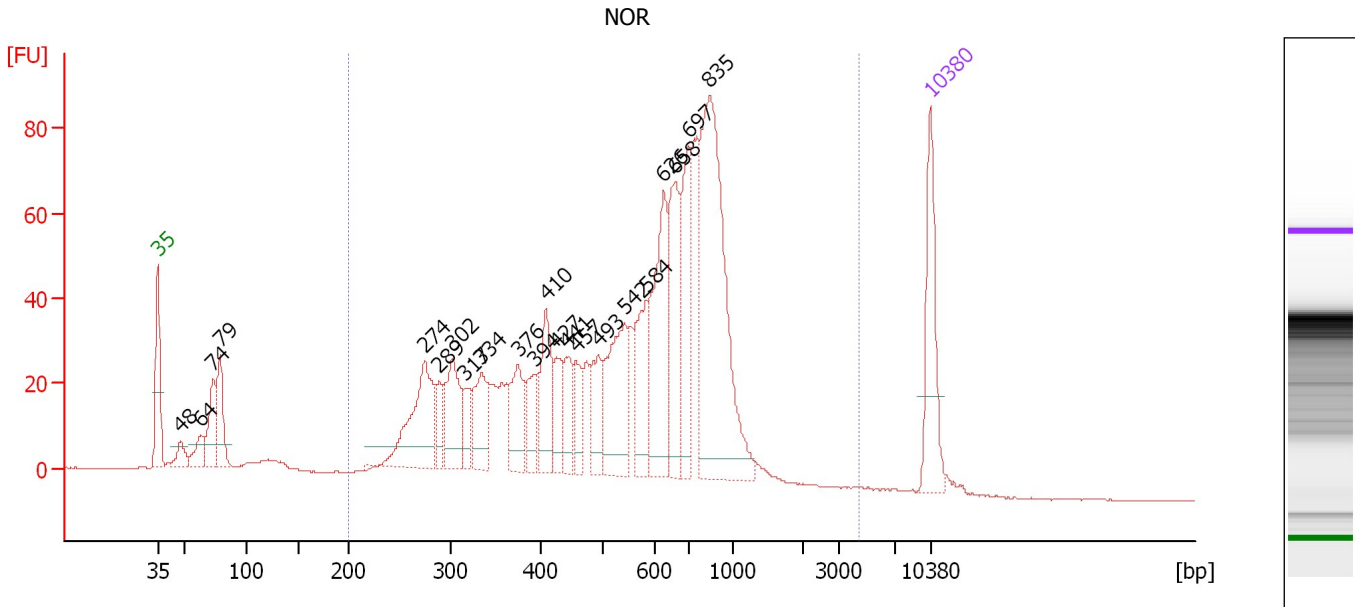
MES

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,545	591	2,801.6	13,218.5	4,154.30	92	36.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : NOR

Number of peaks found: 22 Corr. Area 1: 1,413.0
 Noise: 0.2

Peak table for sample 8 : NOR

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	48	24.39	767.6		45.07
3	64	30.03	710.3		46.94
4	74	63.24	1,295.4		48.05
5	79	68.84	1,314.0		48.66
6	274	138.15	764.5		67.14
7	289	30.45	159.8		68.51
8	302	84.35	423.9		69.67
9	317	29.06	139.1		70.88
10	334	61.46	278.8		72.29
11	376	58.33	235.0		75.68
12	394	41.21	158.6		77.09
13	410	74.16	274.3		78.15
14	427	37.98	134.7		79.16
15	441	44.63	153.2		79.97
16	457	35.69	118.2		80.88
17	493	47.50	146.0		82.90
18	542	109.97	307.4		85.27
19	584	71.31	185.0		87.24
20	626	137.05	331.6		88.81
21	658	108.03	248.7		89.82
22	697	102.42	222.7		91.03
23	835	321.85	583.9		92.95
24	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...

... Region table for sample 8 :

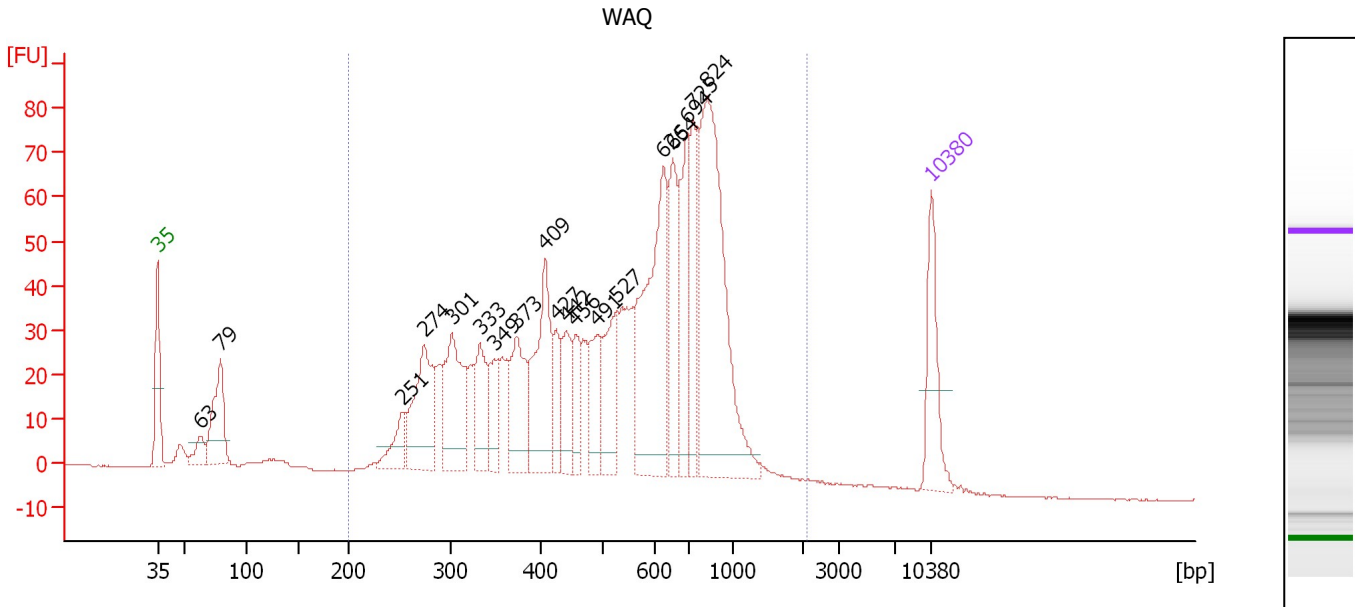
NOR

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	4,408	595	1,413.0	6,250.0	1,932.72	90	44.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : WAQ

Number of peaks found: 19 Corr. Area 1: 1,478.7
 Noise: 0.1

Peak table for sample 9 : WAQ

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	63	32.58	782.9		46.83
3	79	129.19	2,464.1		48.66
4	251	47.45	285.9		65.09
5	274	157.72	873.1		67.13
6	301	160.45	807.4		69.63
7	333	82.15	374.1		72.18
8	349	56.59	246.0		73.46
9	373	107.23	435.3		75.45
10	409	163.69	606.8		78.10
11	427	48.83	173.4		79.12
12	442	64.61	221.6		79.99
13	456	53.00	176.0		80.81
14	491	68.38	211.0		82.80
15	527	94.84	272.5		84.58
16	626	271.16	655.9		88.82
17	654	112.17	259.9		89.68
18	694	131.22	286.3		90.96
19	725	92.46	193.2		91.47
20	824	383.19	704.8		92.80
21	10,380	75.00	10.9	Upper Marker	113.00

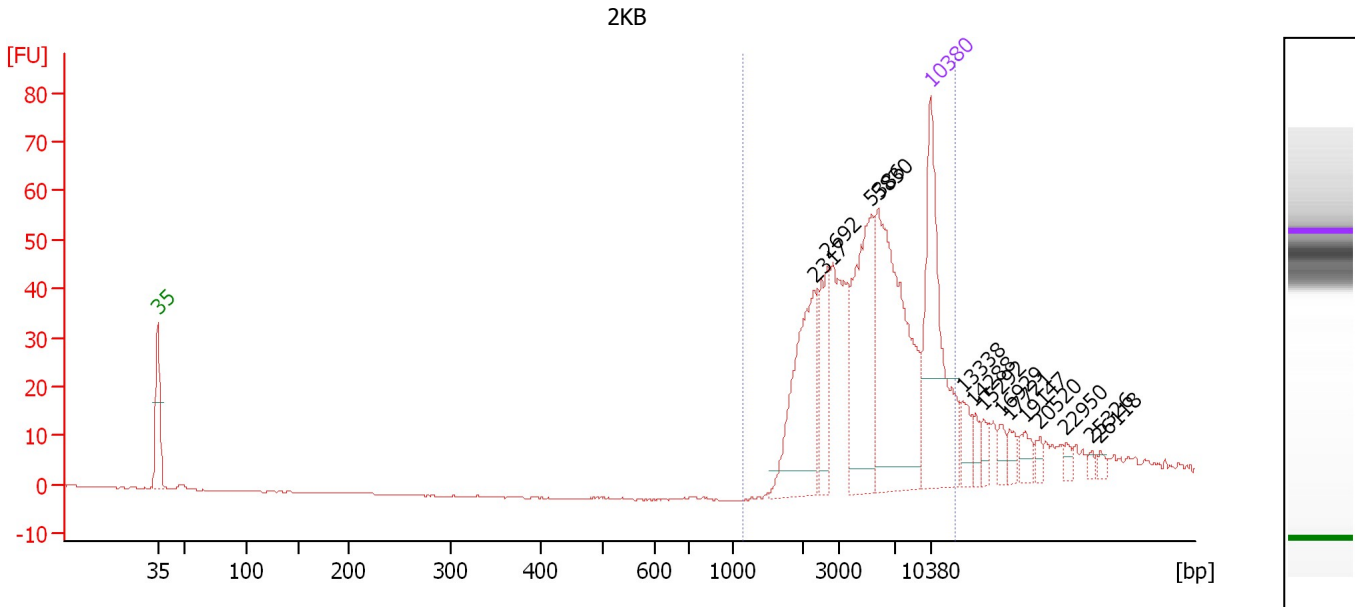
Region table for sample 9 : WAQ

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	2,076	574	1,478.7	8,413.0	2,542.99	93	40.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 2KB

Number of peaks found: 14 Corr. Area 1: 443.4
 Noise: 0.1

Peak table for sample 10 : 2KB

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	2,317	68.52	44.8		102.46
3	2,692	29.74	16.7		103.65
4	5,386	73.09	20.6		107.65
5	5,830	114.70	29.8		108.22
6	10,380	75.00	10.9	Upper Marker	113.00
7	13,338	0.00	0.0		115.88
8	14,288	0.00	0.0		116.80
9	15,292	0.00	0.0		117.78
10	16,929	0.00	0.0		119.37
11	17,721	0.00	0.0		120.14
12	19,147	0.00	0.0		121.53
13	20,520	0.00	0.0		122.87
14	22,950	0.00	0.0		125.23
15	25,326	0.00	0.0		127.54
16	26,118	0.00	0.0		128.32

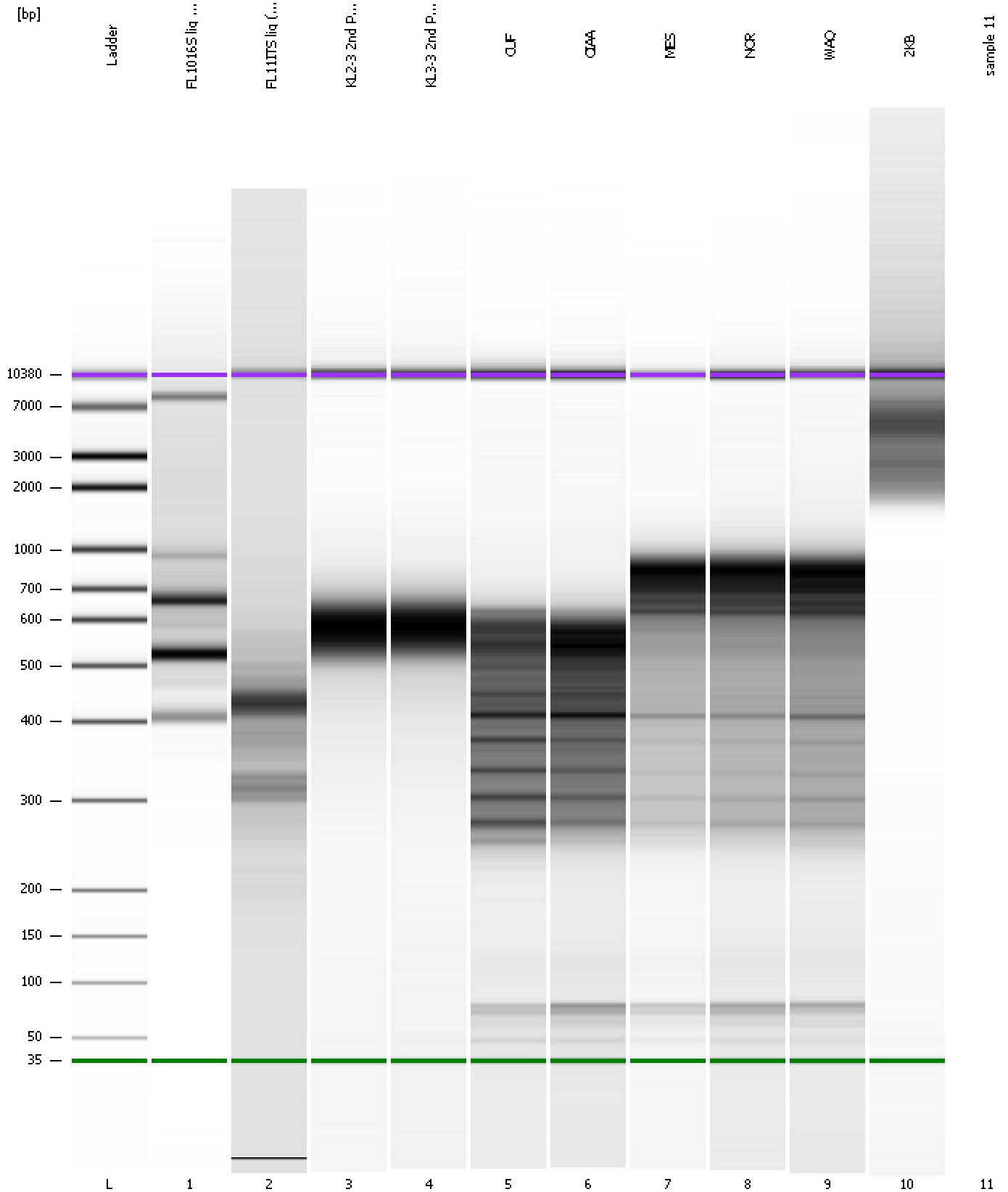
Region table for sample 10 : 2KB

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
1,118	12,659	5,840	443.4	124.0	307.39	81	53.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Gel Image

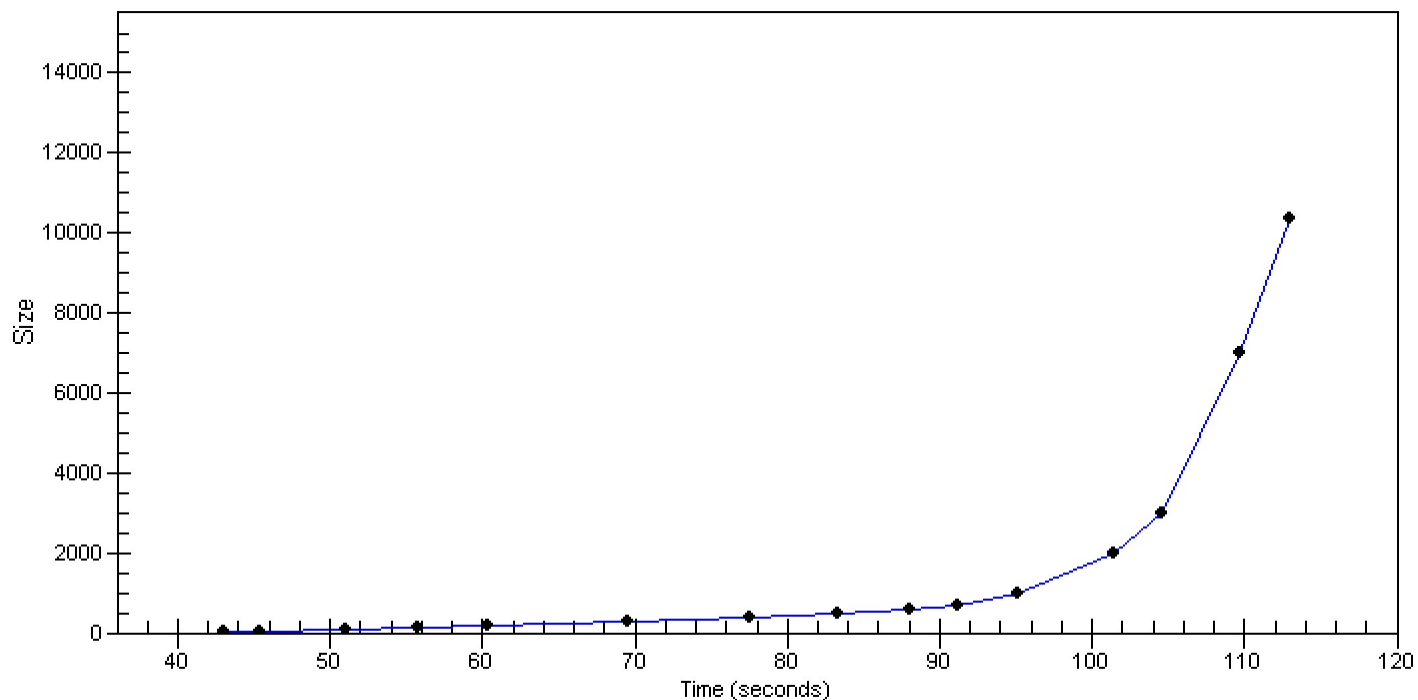


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
Modified: 6/28/2016 10:13:29 AM

Invalid Samples

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad

Created: 6/28/2016 9:31:15 AM
 Modified: 6/28/2016 10:13:29 AM

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		6/28/2016 10:09:42 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-06-28\2016-06-28_001.xad)		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		6/28/2016 9:31:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1