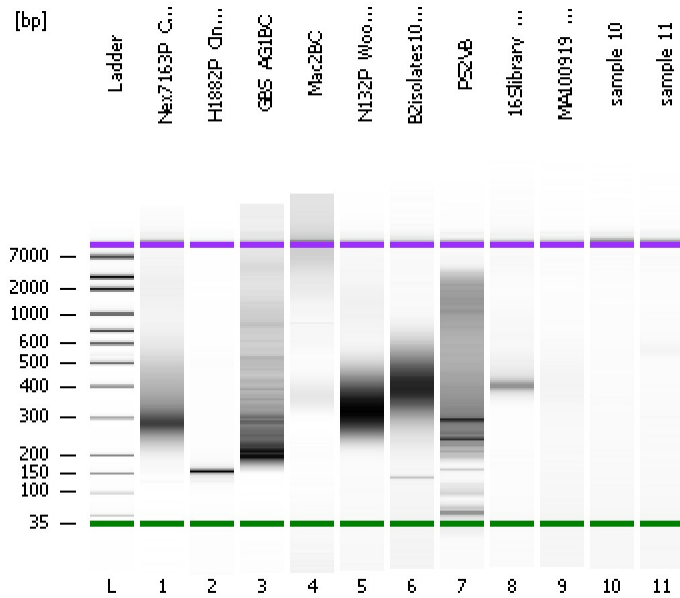


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
Modified: 10/10/2019 2:13:28 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

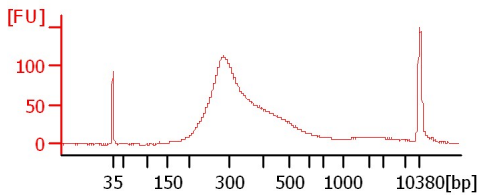
Assay Information:

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

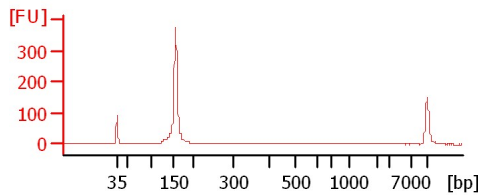
Chip Information:

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

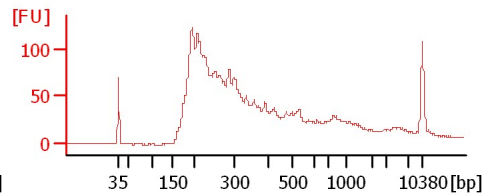
Nex7163P_CincottaBC



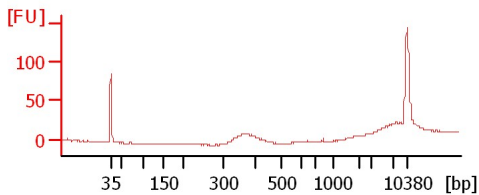
H1882P_CincottaBC



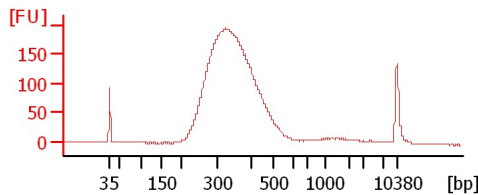
GBS_AG1BC



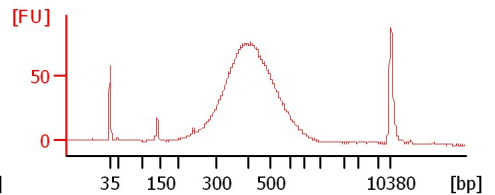
Mac2BC



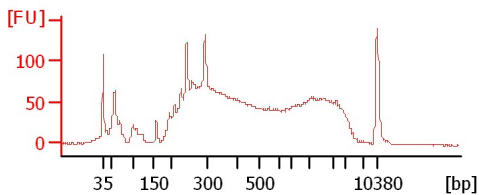
N132P_WoodsBC



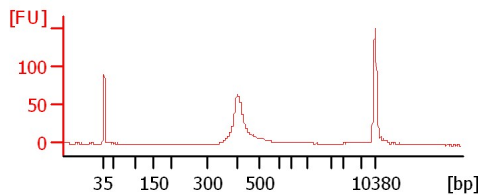
B2isolates10.9.19



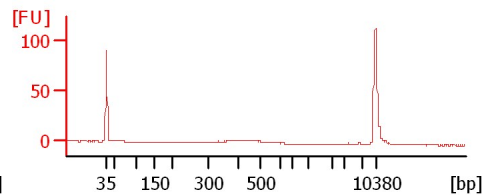
PS2VB



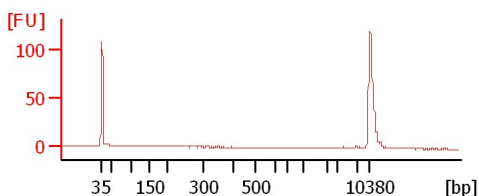
16Slibrary_Moorea_SC_2019



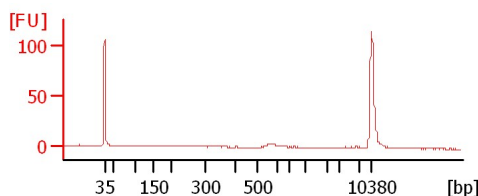
MA100919_UCSB



sample 10



sample 11



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
Modified: 10/10/2019 2:13:28 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Nex7163P_CincottaBC		<input type="checkbox"/>	✓			
H1882P_CincottaBC		<input type="checkbox"/>	✓			
GBS_AG1BC		<input type="checkbox"/>	✓			
Mac2BC		<input type="checkbox"/>	✓			
N132P_WoodsBC		<input type="checkbox"/>	✓			
B2isolates10.9.19		<input type="checkbox"/>	✓			
PS2VB		<input type="checkbox"/>	✓			
16Slibrary_Moorea_SC_2019		<input type="checkbox"/>	✓			
MA100919_UCSB		<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:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
Modified: 10/10/2019 2:13:28 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

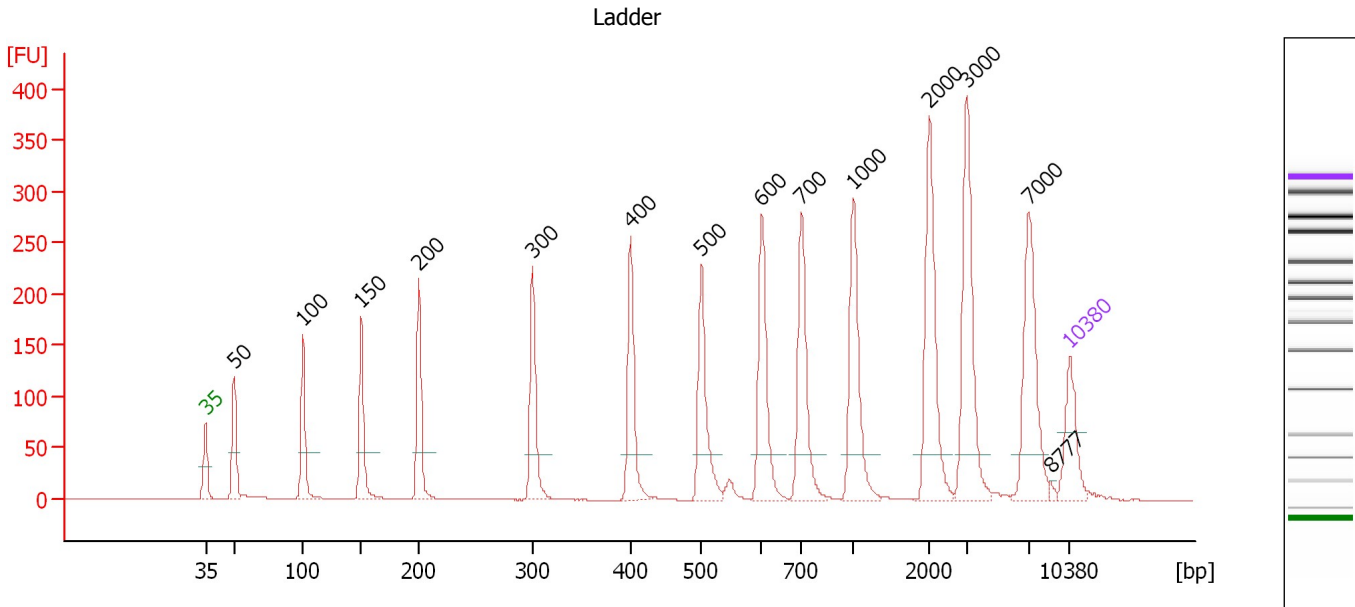
Ladder

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:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

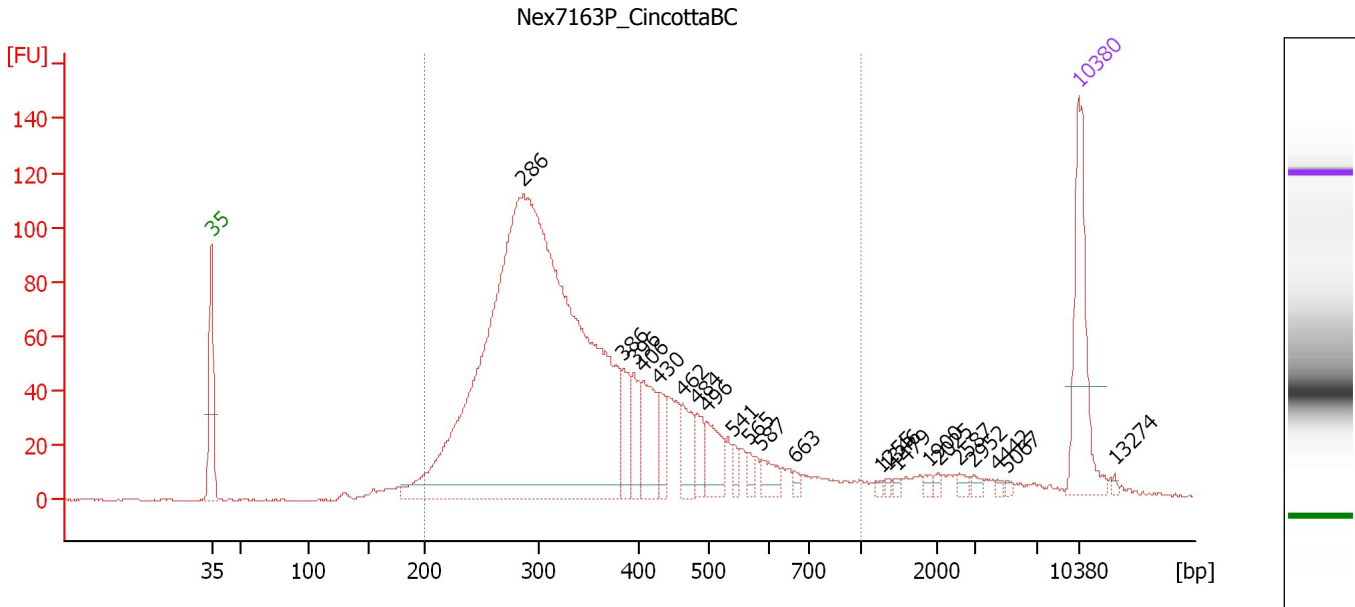
Peak table for Ladder

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	45.29
3	100	150.00	2,272.7	Ladder Peak	50.86
4	150	150.00	1,515.2	Ladder Peak	55.61
5	200	150.00	1,136.4	Ladder Peak	60.27
6	300	150.00	757.6	Ladder Peak	69.47
7	400	150.00	568.2	Ladder Peak	77.42
8	500	150.00	454.5	Ladder Peak	83.16
9	600	150.00	378.8	Ladder Peak	88.04
10	700	150.00	324.7	Ladder Peak	91.24
11	1,000	150.00	227.3	Ladder Peak	95.47
12	2,000	150.00	113.6	Ladder Peak	101.60
13	3,000	150.00	75.8	Ladder Peak	104.67
14	7,000	150.00	32.5	Ladder Peak	109.63
15	8,777	0.00	0.0		111.40
16	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Nex7163P_CincottaBC

Number of peaks found: 22 Corr. Area 1: 2,041.9
 Noise: 0.2

Peak table for sample 1 : Nex7163P_CincottaBC

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	286	1,281.19	6,787.1		68.18
3	386	38.16	149.9		76.28
4	396	33.56	128.4		77.10
5	406	52.92	197.3		77.78
6	430	24.98	88.1		79.12
7	462	33.75	110.7		80.97
8	484	23.29	73.0		82.22
9	496	35.54	108.5		82.95
10	541	9.00	25.2		85.15
11	565	7.83	21.0		86.35
12	587	15.00	38.7		87.43
13	663	4.18	9.5		90.05
14	1,255	2.47	3.0		97.03
15	1,346	2.20	2.5		97.59
16	1,479	2.67	2.7		98.41
17	1,900	3.25	2.6		100.99
18	2,025	2.80	2.1		101.68
19	2,587	3.90	2.3		103.40
20	2,952	3.45	1.8		104.52
21	4,442	1.91	0.7		106.46
22	5,067	1.95	0.6		107.23
23	10,380	75.00	10.9	Upper Marker	113.00
24	13,274	0.00	0.0		115.88

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...

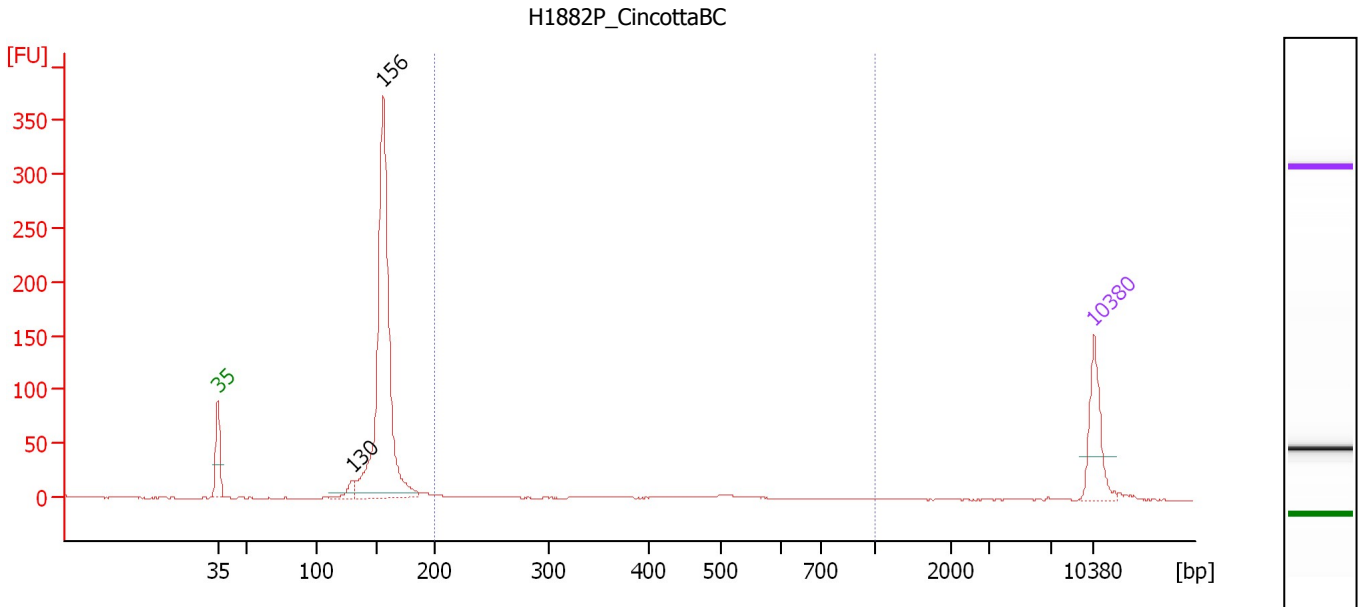
... Region table for sample 1 : Nex7163P CincottaBC

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co lor % of Total	Size distribution in CV [%]
200	1,000	358	1,606.62	2,041.9	7,593.0	 93	33.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : H1882P_CincottaBC

Number of peaks found: 2 Corr. Area 1: 80.5
 Noise: 0.3

Peak table for sample 2 : H1882P_CincottaBC

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	130	23.55	274.8		53.69
3	156	779.41	7,550.1		56.21
4	10,380	75.00	10.9	Upper Marker	113.00

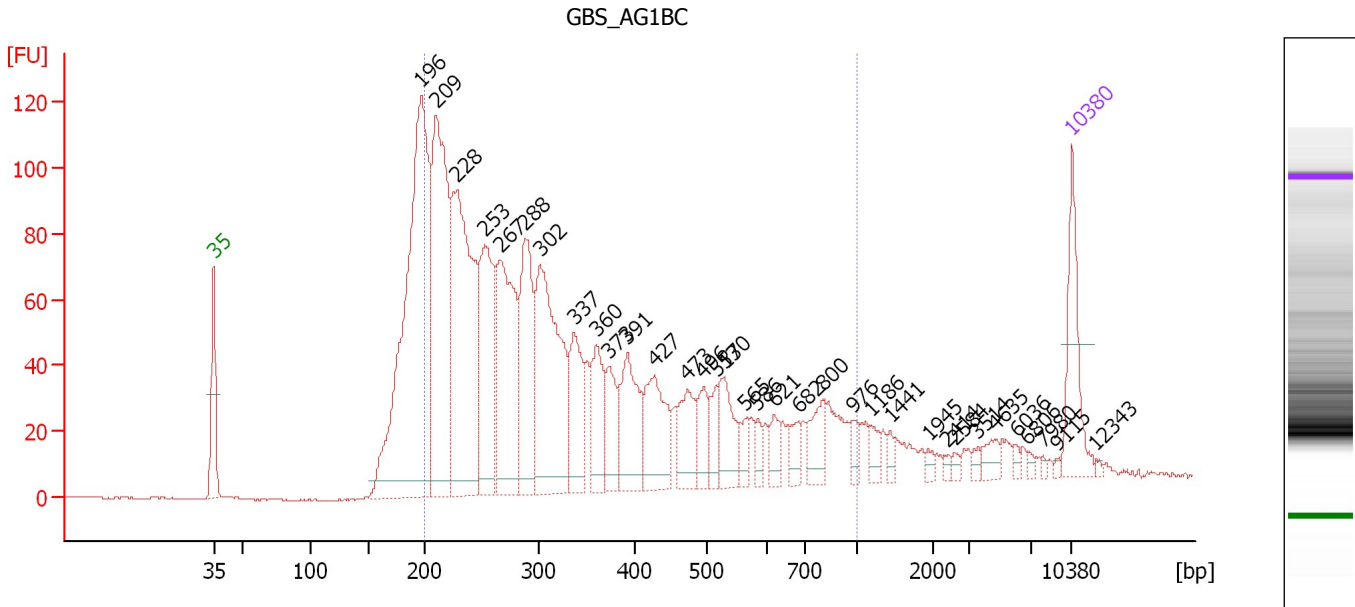
Region table for sample 2 : H1882P_CincottaBC

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	434	64.64	80.5	279.8	9	37.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : GBS_AG1BC

Number of peaks found: 34 Corr. Area 1: 2,186.5
 Noise: 0.2

Peak table for sample 3 : GBS_AG1BC

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	196	754.74	5,830.3		59.91
3	209	453.45	3,282.3		61.13
4	228	451.26	2,995.8		62.87
5	253	236.59	1,417.9		65.13
6	267	272.64	1,550.0		66.39
7	288	194.59	1,022.8		68.39
8	302	315.79	1,582.9		69.65
9	337	109.32	491.9		72.39
10	360	89.07	375.1		74.22
11	373	61.94	251.7		75.26
12	391	118.09	457.1		76.74
13	427	108.79	386.2		78.96
14	473	67.44	216.0		81.61
15	496	42.45	129.5		82.96
16	517	30.31	88.8		84.00
17	530	65.53	187.4		84.61
18	565	20.62	55.3		86.35
19	586	14.89	38.5		87.35
20	621	24.15	59.0		88.70
21	682	25.84	57.4		90.65
22	800	44.59	84.4		92.65
23	976	14.76	22.9		95.13
24	1,186	17.02	21.7		96.61
25	1,441	8.52	9.0		98.17

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : GBS_AG1BC

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,945	6.20	4.8		101.26
27	2,414	4.61	2.9		102.87
28	2,584	5.07	3.0		103.39
29	3,514	5.79	2.5		105.30
30	4,635	15.49	5.1		106.70
31	6,036	4.82	1.2		108.43
32	6,806	3.72	0.8		109.39
33	7,980	2.93	0.6		110.61
34	9,115	2.54	0.4		111.74
35	10,380	75.00	10.9	Upper Marker	113.00
36	12,343	0.00	0.0		114.96

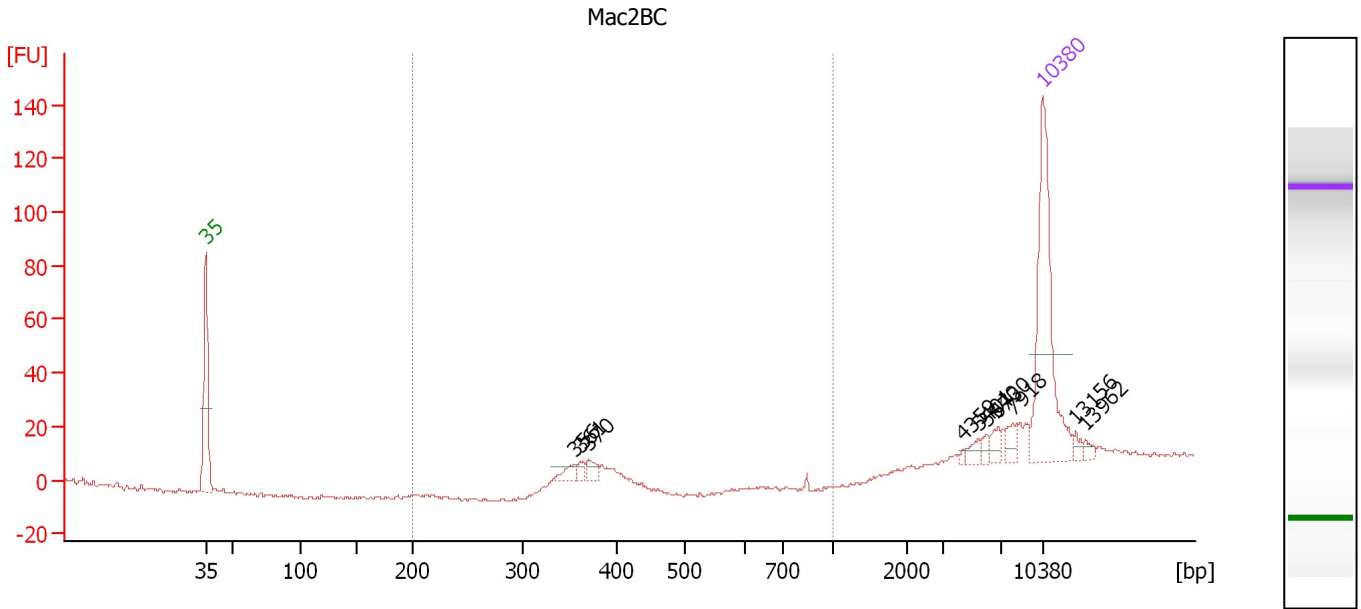
Region table for sample 3 : GBS_AG1BC

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	365	2,980.46	2,186.5	15,269.4	79	46.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Mac2BC

Number of peaks found: 10 Corr. Area 1: 10.1
 Noise: 0.5

Peak table for sample 4 : Mac2BC

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	356	7.58	32.3		73.92
3	361	4.32	18.1		74.32
4	370	6.99	28.6		75.03
5	4,359	1.41	0.5		106.35
6	5,401	4.42	1.2		107.65
7	5,940	3.07	0.8		108.32
8	6,730	5.44	1.2		109.30
9	7,918	5.72	1.1		110.55
10	10,380	75.00	10.9	Upper Marker	113.00
11	13,156	0.00	0.0		115.77
12	13,962	0.00	0.0		116.57

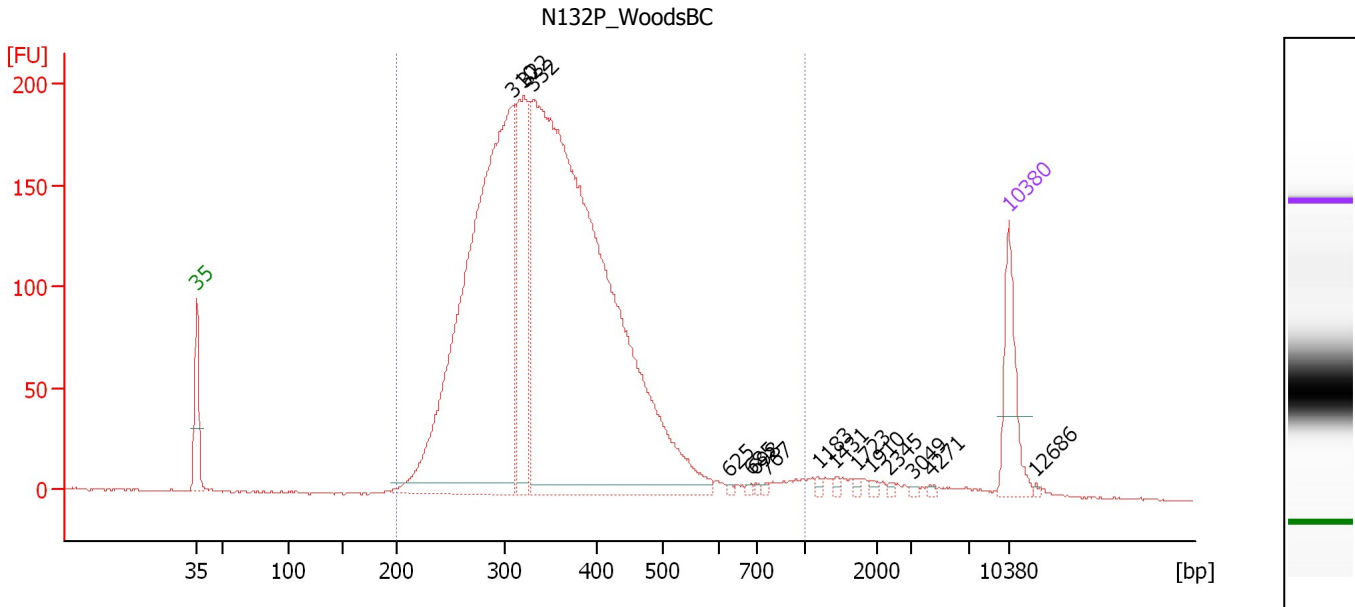
Region table for sample 4 : Mac2BC

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	368	7.02	10.1	28.9	11	2.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : N132P WoodsBC

Number of peaks found: 15 Corr. Area 1: 3,798.7
 Noise: 0.4

Peak table for sample 5 : N132P WoodsBC

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	310	1,153.22	5,628.3		70.30
3	322	241.09	1,134.4		71.22
4	332	1,747.62	7,965.4		72.05
5	625	3.15	7.6		88.82
6	685	2.82	6.2		90.75
7	698	2.98	6.5		91.17
8	767	3.20	6.3		92.18
9	1,183	3.64	4.7		96.59
10	1,431	3.50	3.7		98.11
11	1,723	3.60	3.2		99.90
12	1,910	4.31	3.4		101.05
13	2,345	2.86	1.8		102.66
14	3,049	2.26	1.1		104.73
15	4,271	2.50	0.9		106.24
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,686	0.00	0.0		115.30

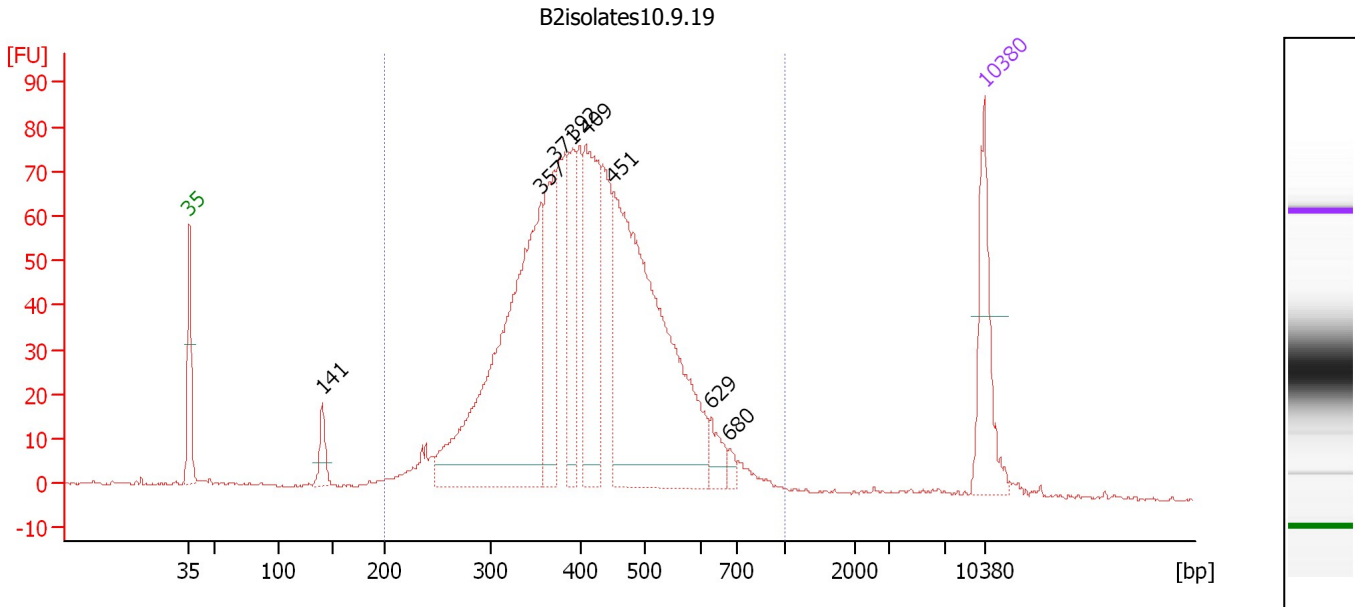
Region table for sample 5 : N132P WoodsBC

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	355	3,165.28	3,798.7	14,430.6	97	25.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : B2isolates10.9.19

Number of peaks found: 8 Corr. Area 1: 1,522.6
 Noise: 0.2

Peak table for sample 6 : B2isolates10.9.19

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	141	33.11	356.2		54.74
3	357	481.48	2,044.5		73.99
4	371	133.18	543.9		75.11
5	392	90.25	349.1		76.76
6	409	189.75	703.1		77.93
7	451	507.60	1,703.7		80.37
8	629	24.56	59.2		88.96
9	680	8.74	19.5		90.61
10	10,380	75.00	10.9	Upper Marker	113.00

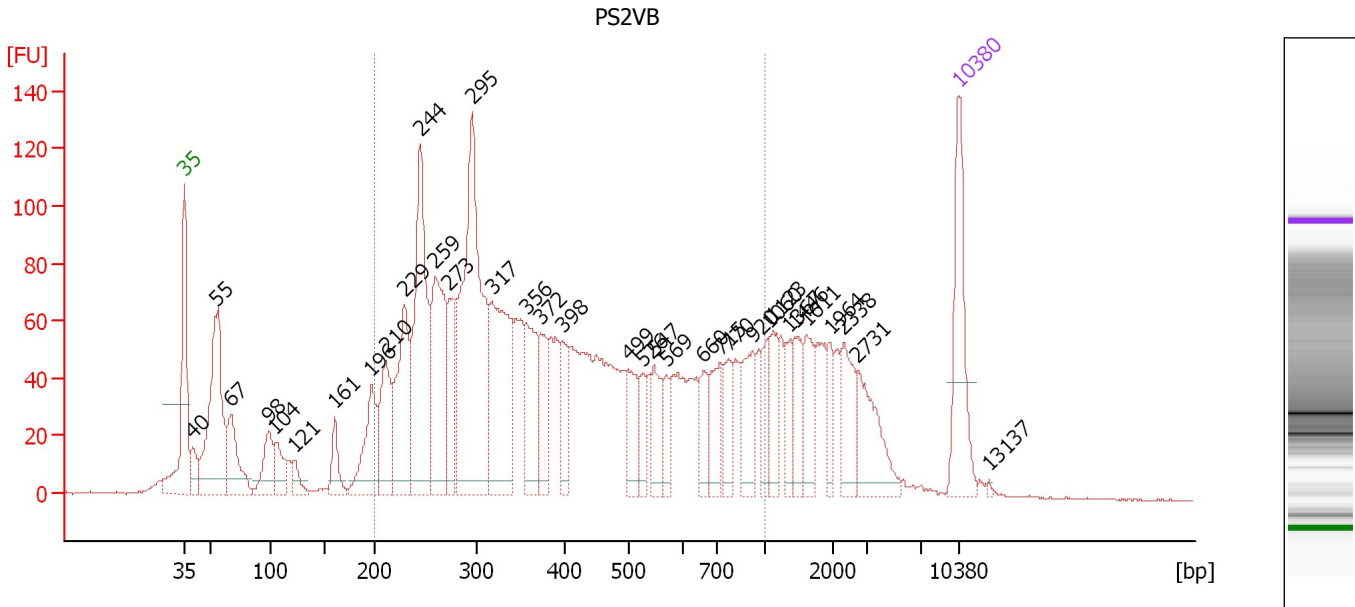
Region table for sample 6 : B2isolates10.9.19

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	423	1,845.86	1,522.6	7,148.6	96	24.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : PS2VB

Number of peaks found: 35 Corr. Area 1: 2,635.8
 Noise: 0.3

Peak table for sample 7 : PS2VB

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	40	31.37	1,186.7		43.77
3	55	218.09	5,962.5		45.89
4	67	75.47	1,704.0		47.19
5	98	52.61	814.8		50.62
6	104	35.03	512.7		51.20
7	121	20.23	252.7		52.88
8	161	33.13	312.6		56.60
9	196	83.08	641.5		59.92
10	210	82.51	594.4		61.22
11	229	131.17	869.1		62.91
12	244	219.44	1,363.4		64.31
13	259	153.21	896.1		65.71
14	273	63.96	355.4		66.96
15	295	333.95	1,713.8		69.03
16	317	173.84	831.1		70.82
17	356	79.53	338.7		73.90
18	372	50.10	204.0		75.20
19	398	42.36	161.4		77.23
20	499	47.30	143.6		83.11
21	526	26.06	75.1		84.41
22	547	41.85	115.8		85.47
23	569	22.31	59.4		86.53
24	660	30.62	70.3		89.96
25	715	42.23	89.5		91.45

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...

... Peak table for sample 7 : PS2VB

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	770	30.67	60.4		92.22
27	920	47.89	78.9		94.34
28	1,060	28.40	40.6		95.84
29	1,123	31.43	42.4		96.22
30	1,367	23.66	26.2		97.72
31	1,446	33.77	35.4		98.20
32	1,611	34.34	32.3		99.21
33	1,964	19.07	14.7		101.38
34	2,338	41.69	27.0		102.63
35	2,731	56.18	31.2		103.84
36	10,380	75.00	10.9	Upper Marker	113.00
37	13,137	0.00	0.0		115.75

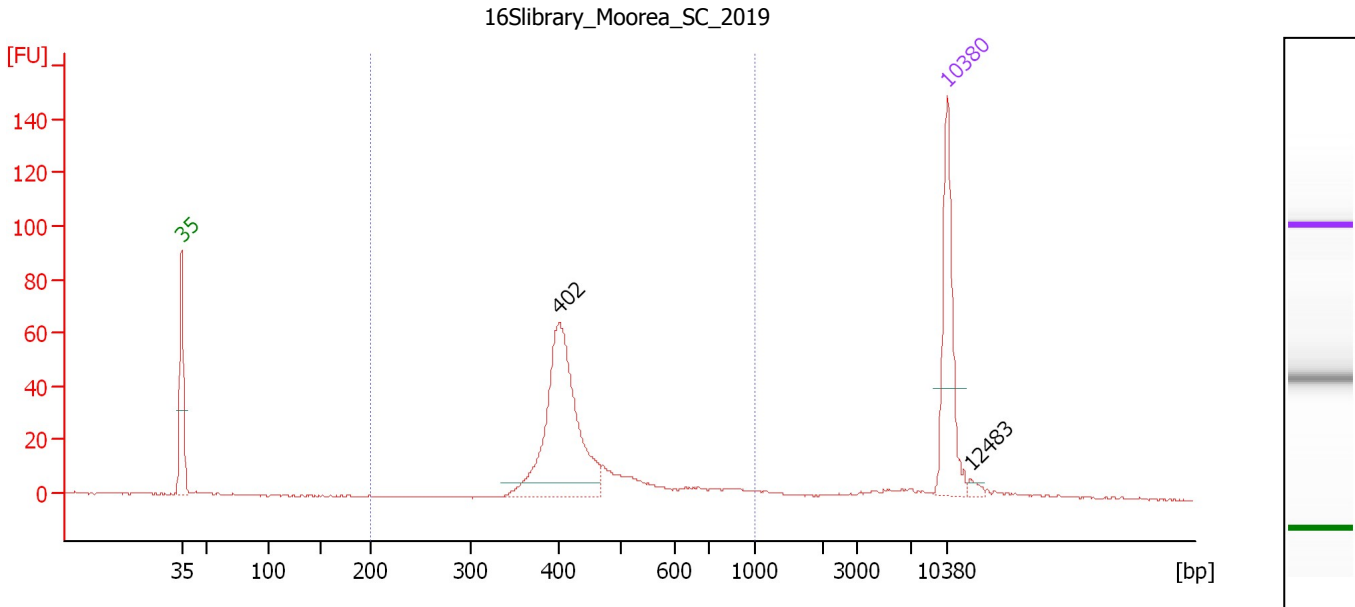
Region table for sample 7 : PS2VB

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	428	2,208.10	2,635.8	9,853.5	70	44.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : 16Slibrary_Moorea_SC_2019

Number of peaks found: 2 Corr. Area 1: 347.0
 Noise: 0.3

Peak table for sample 8 : 16Slibrary_Moorea_SC_2019

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	402	212.57	801.8		77.51
3	10,380	75.00	10.9	Upper Marker	113.00
4	12,483	0.00	0.0		115.10

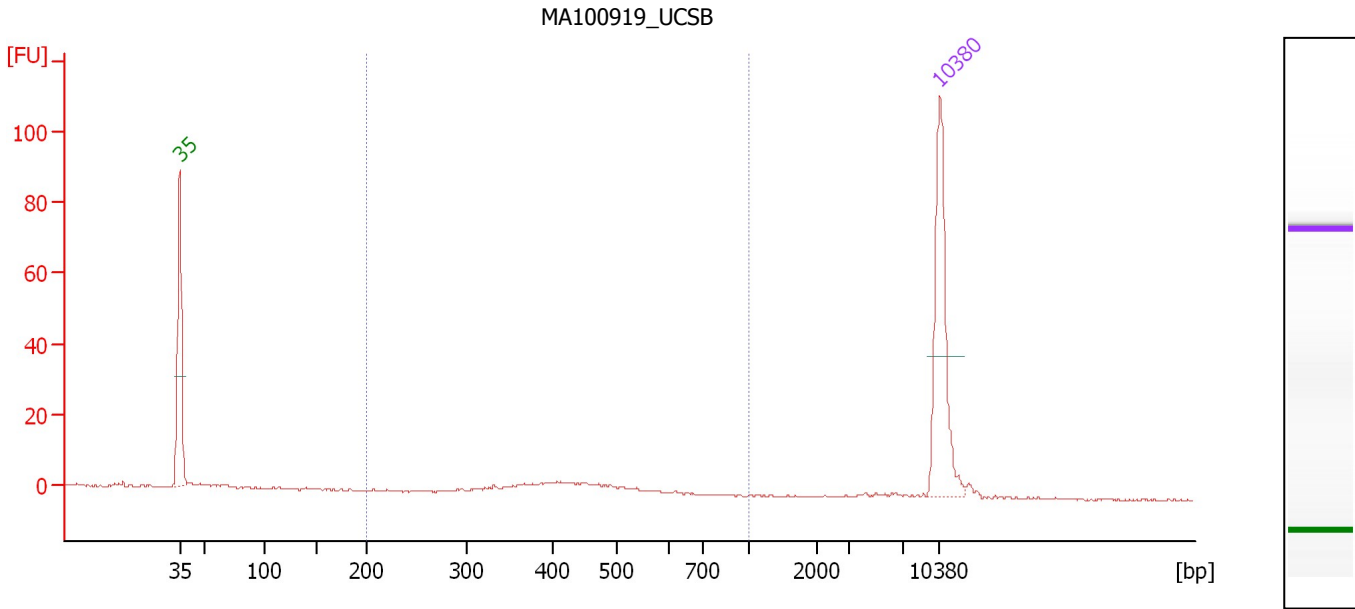
Region table for sample 8 : 16Slibrary_Moorea_SC_2019

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	453	264.06	347.0	924.8	86	23.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : MA100919 UCSB

Number of peaks found: 0 Corr. Area 1: 34.8
 Noise: 0.2

Peak table for sample 9 : MA100919 UCSB

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	10,380	75.00	10.9	Upper Marker	113.00

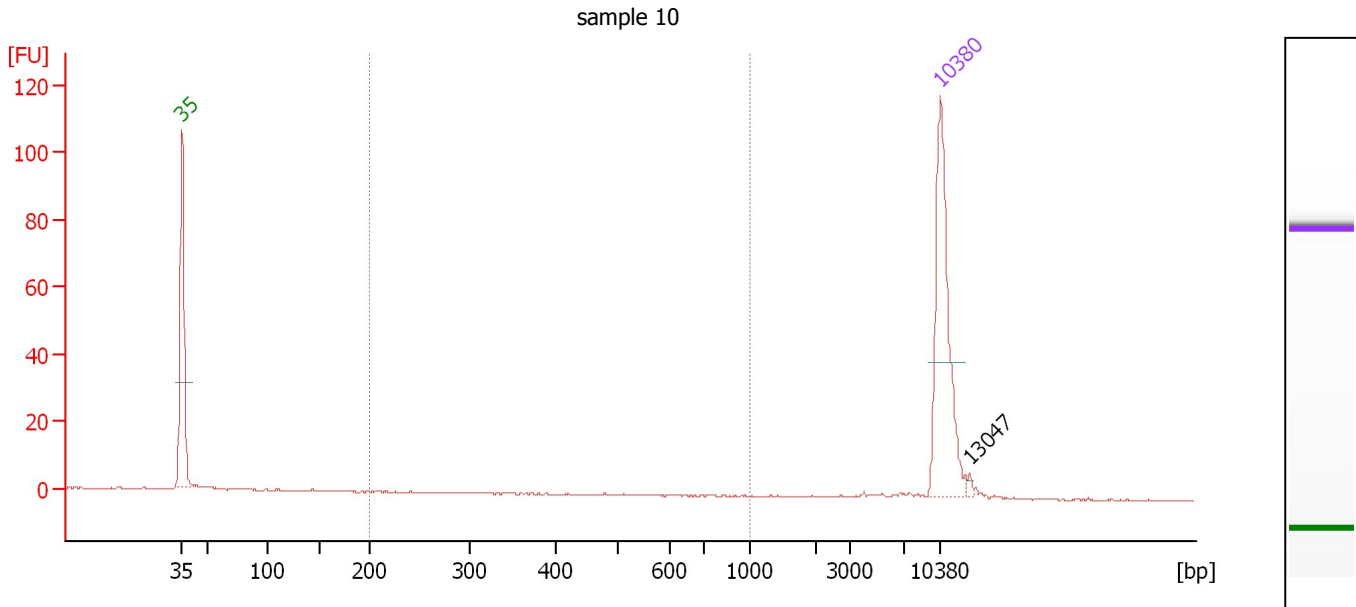
Region table for sample 9 : MA100919 UCSB

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	428	32.07	34.8	116.7	73	15.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 1 Corr. Area 1: 0.0
 Noise: 0.1

Peak table for sample 10 : sample 10

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	10,380	75.00	10.9	Upper Marker	113.00
3	13,047	0.00	0.0		115.66

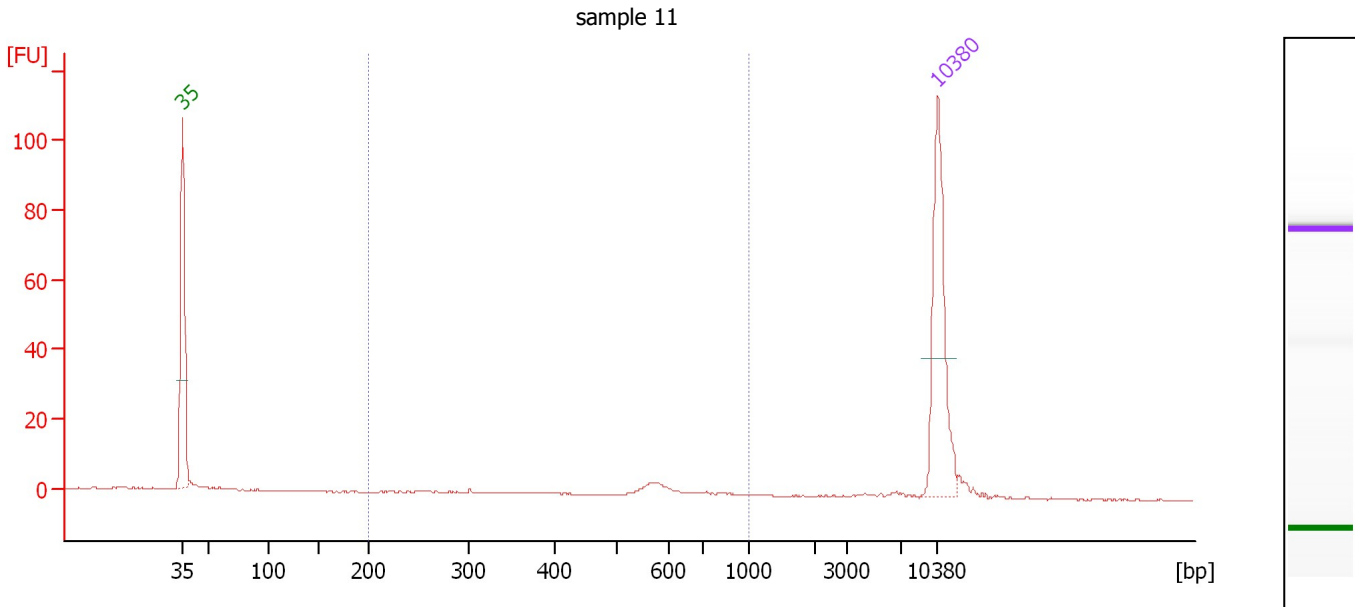
Region table for sample 10 : sample 10

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	559	0.03	0.0	0.1	0	35.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 0 Corr. Area 1: 18.8
 Noise: 0.2

Peak table for sample 11 : sample 11

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	10,380	75.00	10.9	Upper Marker	113.00

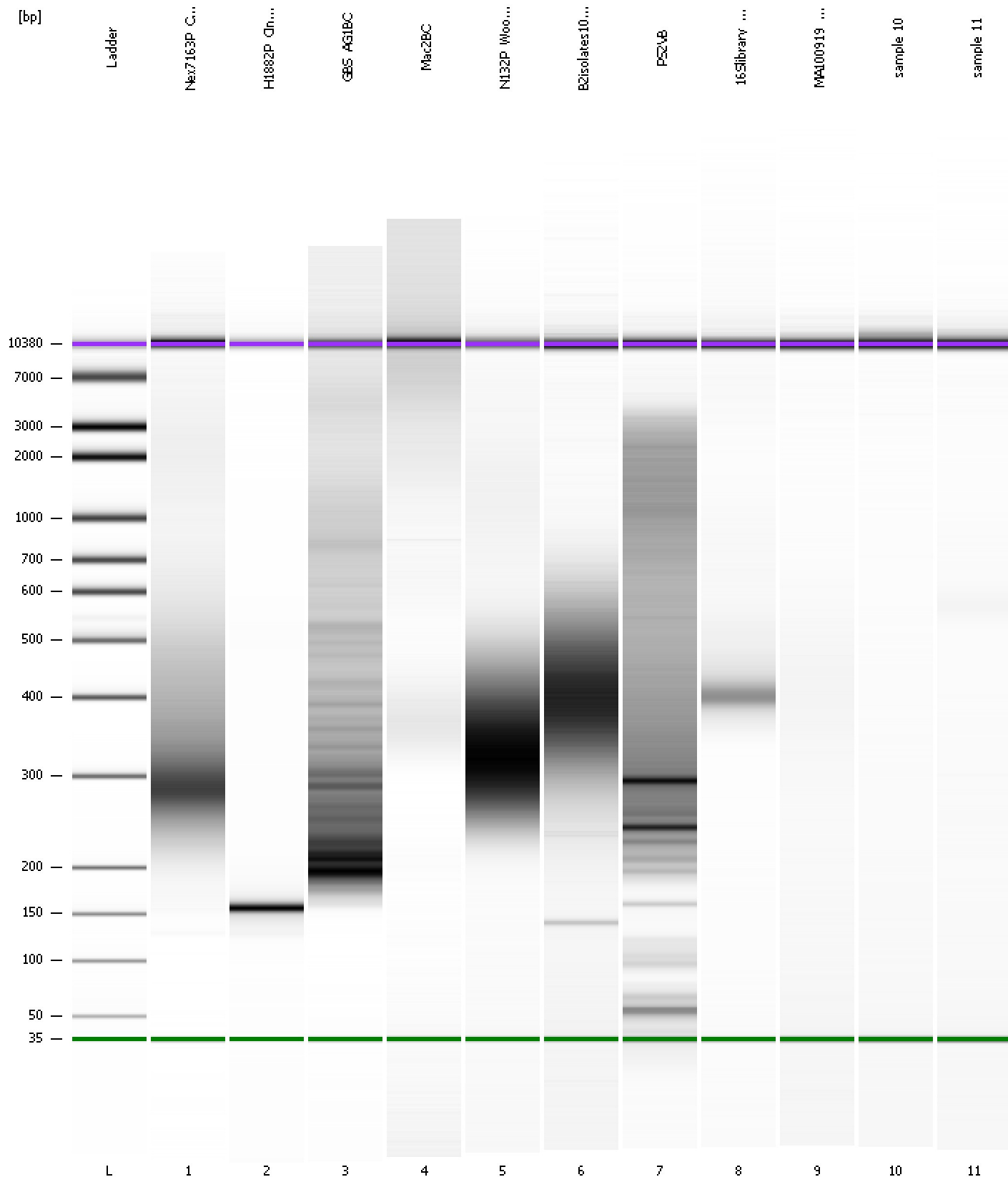
Region table for sample 11 : sample 11

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	585	15.06	18.8	44.9	39	24.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
Modified: 10/10/2019 2:13:28 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad

Created: 10/10/2019 1:32:12 PM
 Modified: 10/10/2019 2:13:28 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		10/10/2019 2:13:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2019-10-10\2019-10-10_002.xad)		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/10/2019 1:32:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1