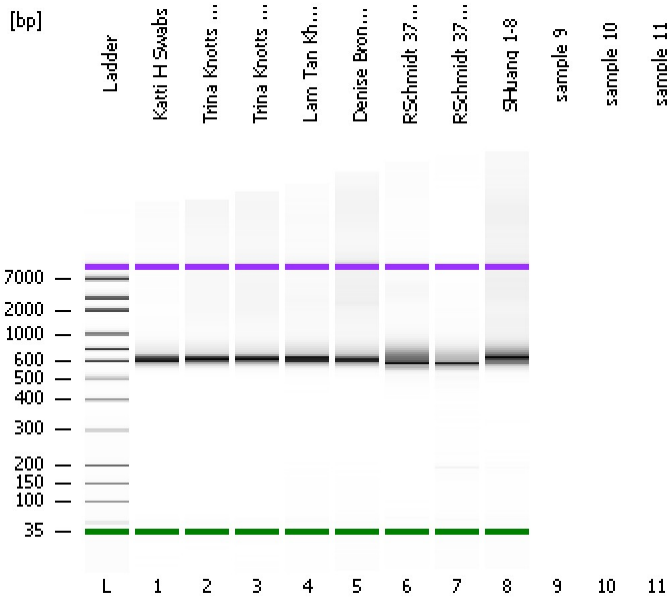


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 PM

Electrophoresis File Run Summary



Instrument Information:

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

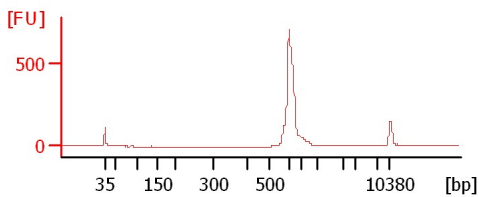
Assay Information:

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

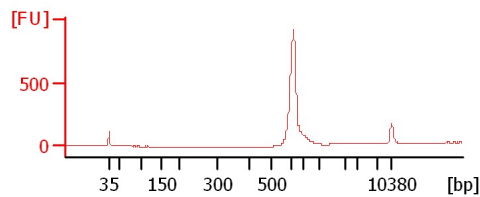
Chip Information:

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

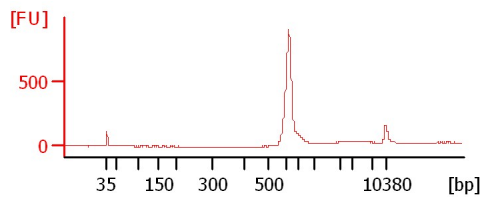
Katti H Swabs



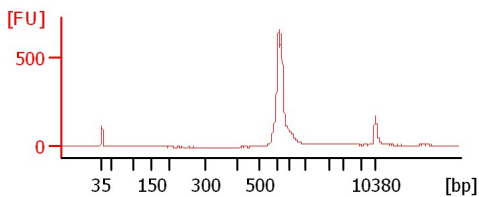
Trina Knotts 1-48



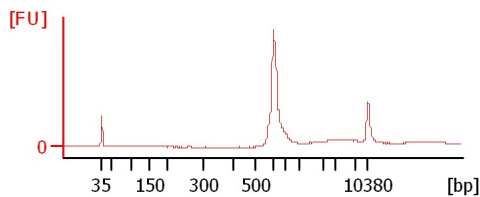
Trina Knotts 49-82



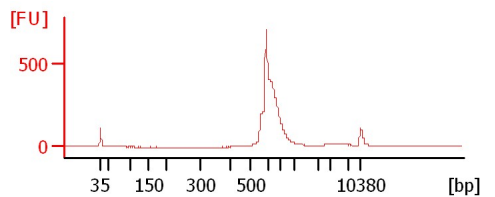
Lam Tan Khuat 1-22



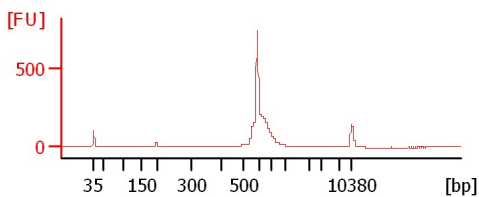
Denise Bronner 1-18 nested



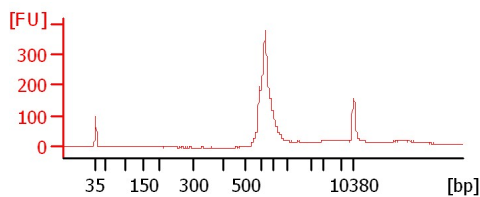
RSchmidt 37-44 V3V4



RSchmidt 37-44 ITS



SHuang 1-8



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Katti H Swabs		<input type="checkbox"/>	✓			
Trina Knotts 1-48		<input type="checkbox"/>	✓			
Trina Knotts 49-82		<input type="checkbox"/>	✓			
Lam Tan Khuat 1-22		<input type="checkbox"/>	✓			
Denise Bronner 1-18 nested		<input type="checkbox"/>	✓			
RSchmidt 37-44 V3V4		<input type="checkbox"/>	✓			
RSchmidt 37-44 ITS		<input type="checkbox"/>	✓			
SHuang 1-8		<input type="checkbox"/>	✓			
sample 9		<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\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 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

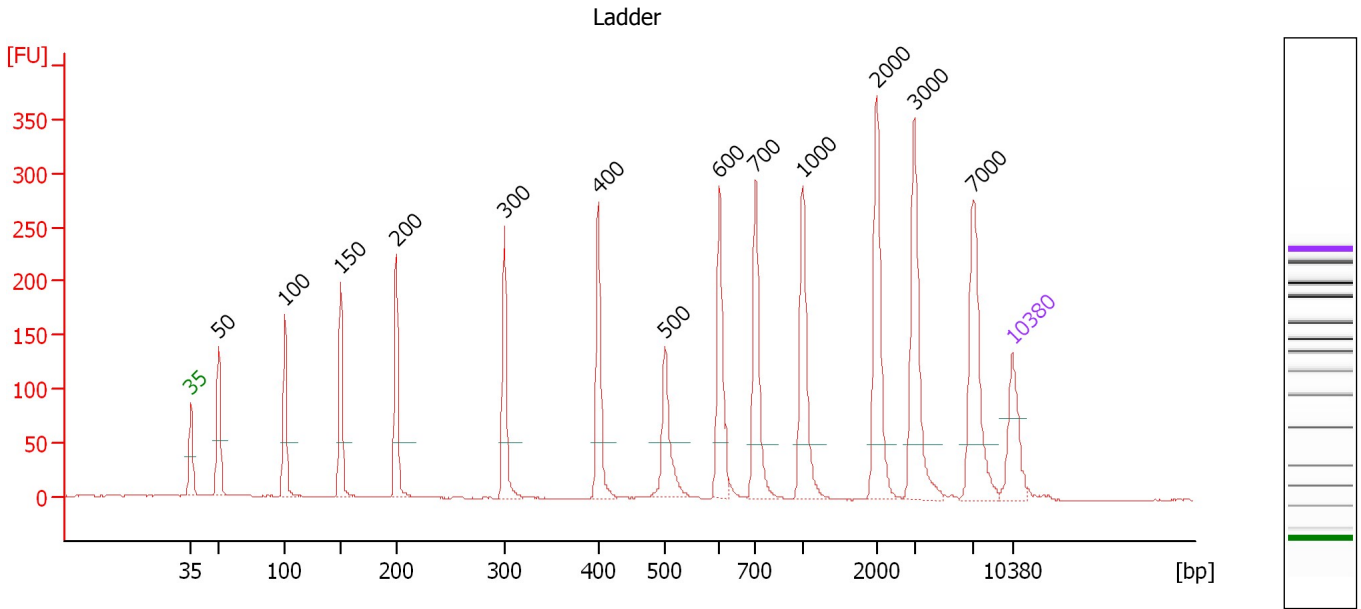
Ladder

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\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

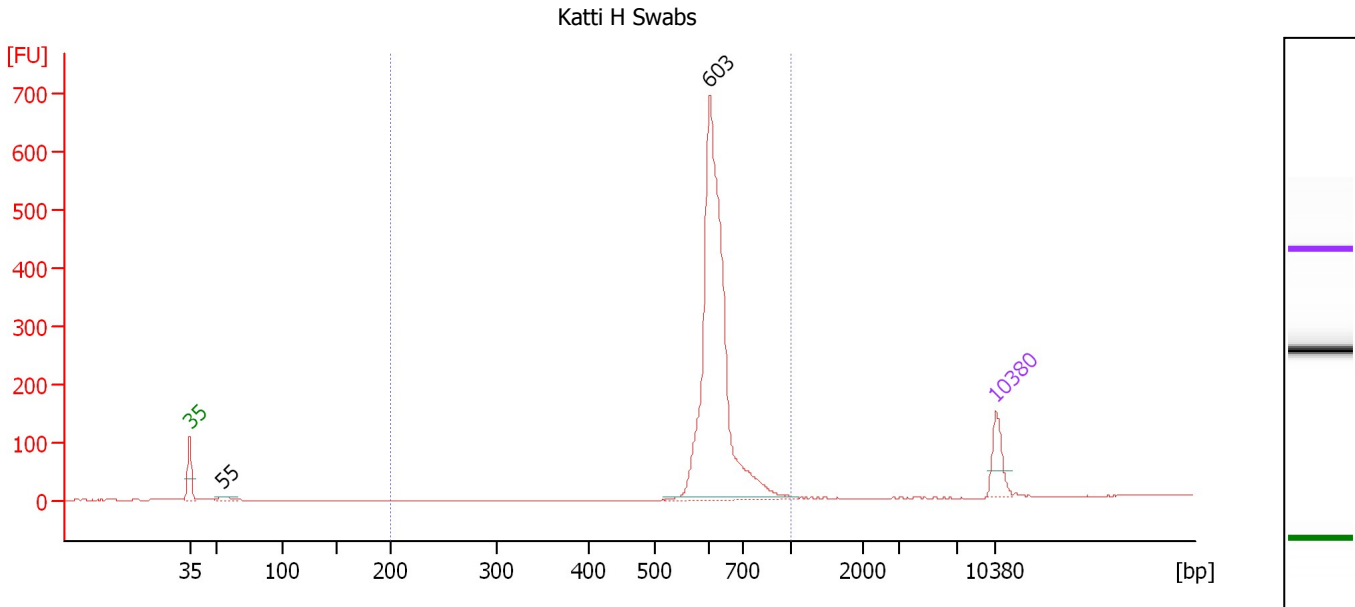
Peak table for Ladder

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	50.99
4	150	150.00	1,515.2	Ladder Peak	55.76
5	200	150.00	1,136.4	Ladder Peak	60.49
6	300	150.00	757.6	Ladder Peak	69.71
7	400	150.00	568.2	Ladder Peak	77.70
8	500	150.00	454.5	Ladder Peak	83.38
9	600	150.00	378.8	Ladder Peak	88.06
10	700	150.00	324.7	Ladder Peak	91.11
11	1,000	150.00	227.3	Ladder Peak	95.15
12	2,000	150.00	113.6	Ladder Peak	101.42
13	3,000	150.00	75.8	Ladder Peak	104.64
14	7,000	150.00	32.5	Ladder Peak	109.64
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Katti H Swabs

Number of peaks found: 2 Corr. Area 1: 1,597.3
 Noise: 0.3

Peak table for sample 1 : Katti H Swabs

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	55	20.14	555.7		45.92
3	603	1,251.59	3,147.3		88.14
4	10,380	75.00	10.9	Upper Marker	113.00

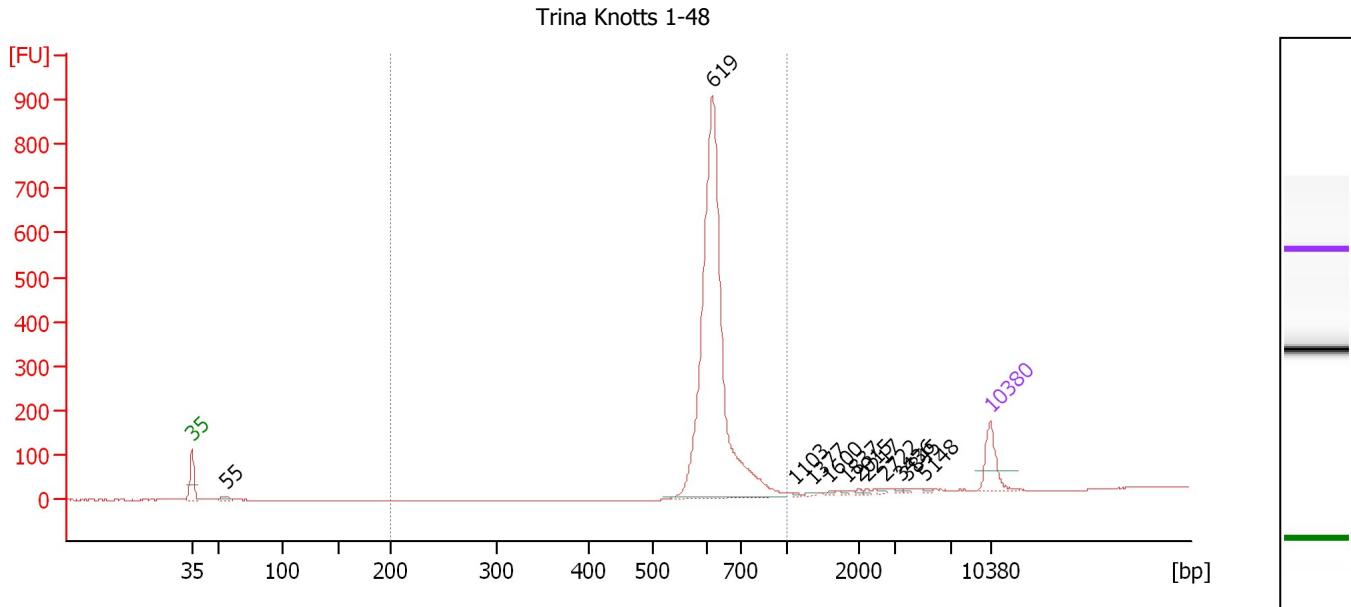
Region table for sample 1 : Katti H Swabs

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	626	1,204.97	1,597.3	2,931.8	99	7.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Trina Knotts 1-48

Number of peaks found: 12 Corr. Area 1: 1,963.1
 Noise: 0.5

Peak table for sample 2 : Trina Knotts 1-48

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	55	16.58	453.0		45.98
3	619	1,387.61	3,398.7		88.63
4	1,103	2.13	2.9		95.79
5	1,377	3.17	3.5		97.51
6	1,600	4.65	4.4		98.91
7	1,837	3.30	2.7		100.40
8	2,015	3.62	2.7		101.47
9	2,217	3.19	2.2		102.12
10	2,722	5.00	2.8		103.74
11	3,436	3.93	1.7		105.19
12	3,845	3.60	1.4		105.70
13	5,148	3.69	1.1		107.33
14	10,380	75.00	10.9	Upper Marker	113.00

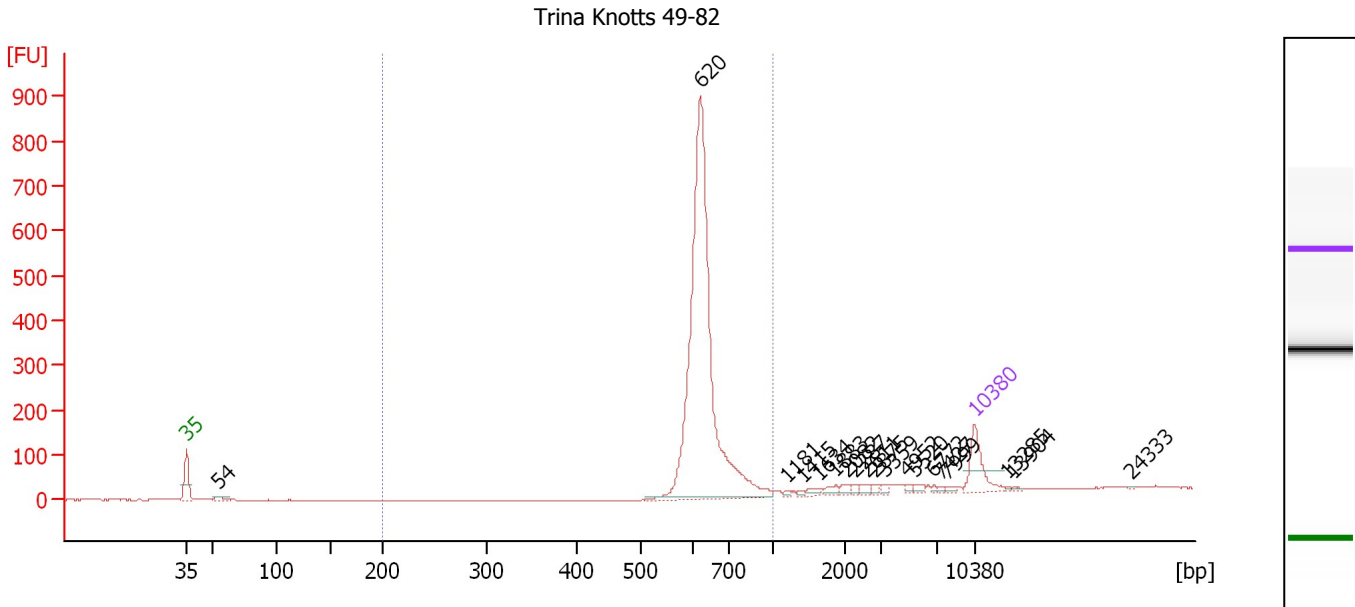
Region table for sample 2 : Trina Knotts 1-48

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	627	1,292.28	1,963.1	3,133.1	99	7.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Trina Knotts 49-82

Number of peaks found: 19 Corr. Area 1: 1,954.8
 Noise: 0.5

Peak table for sample 3 : Trina Knotts 49-82

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	54	15.64	436.8		45.84
3	620	1,221.72	2,987.8		88.66
4	1,181	4.96	6.4		96.28
5	1,415	4.39	4.7		97.75
6	1,634	10.26	9.5		99.12
7	1,883	12.84	10.3		100.69
8	2,082	10.66	7.8		101.68
9	2,287	6.95	4.6		102.34
10	2,611	11.81	6.9		103.39
11	2,875	7.67	4.0		104.24
12	3,359	8.47	3.8		105.09
13	4,952	5.59	1.7		107.08
14	5,520	8.64	2.4		107.79
15	6,772	4.23	0.9		109.35
16	7,427	4.14	0.8		110.06
17	7,999	5.19	1.0		110.63
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,285	0.00	0.0		115.89
20	13,904	0.00	0.0		116.50
21	24,333	0.00	0.0		126.88

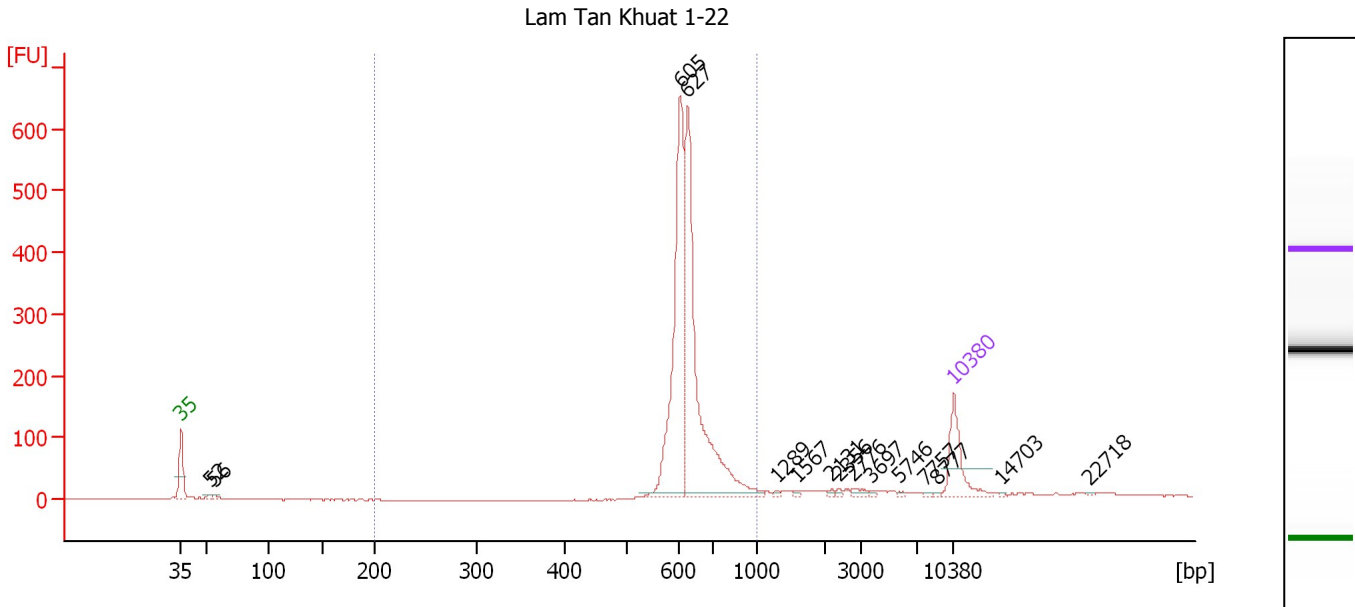
Region table for sample 3 : Trina Knotts 49-82

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	632	1,145.26	1,954.8	2,760.4	90	8.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Lam Tan Khuat 1-22

Number of peaks found: 15 Corr. Area 1: 1,716.7
 Noise: 0.7

Peak table for sample 4 : Lam Tan Khuat 1-22

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	52	4.80	139.4		45.61
3	56	4.74	129.2		46.00
4	605	453.82	1,136.3		88.22
5	627	534.41	1,290.7		88.89
6	1,289	3.09	3.6		96.96
7	1,567	2.84	2.7		98.70
8	2,131	3.26	2.3		101.84
9	2,356	3.78	2.4		102.57
10	2,776	8.18	4.5		103.92
11	3,697	2.75	1.1		105.51
12	5,746	1.92	0.5		108.07
13	7,757	1.94	0.4		110.39
14	8,777	1.48	0.3		111.41
15	10,380	75.00	10.9	Upper Marker	113.00
16	14,703	0.00	0.0		117.30
17	22,718	0.00	0.0		125.27

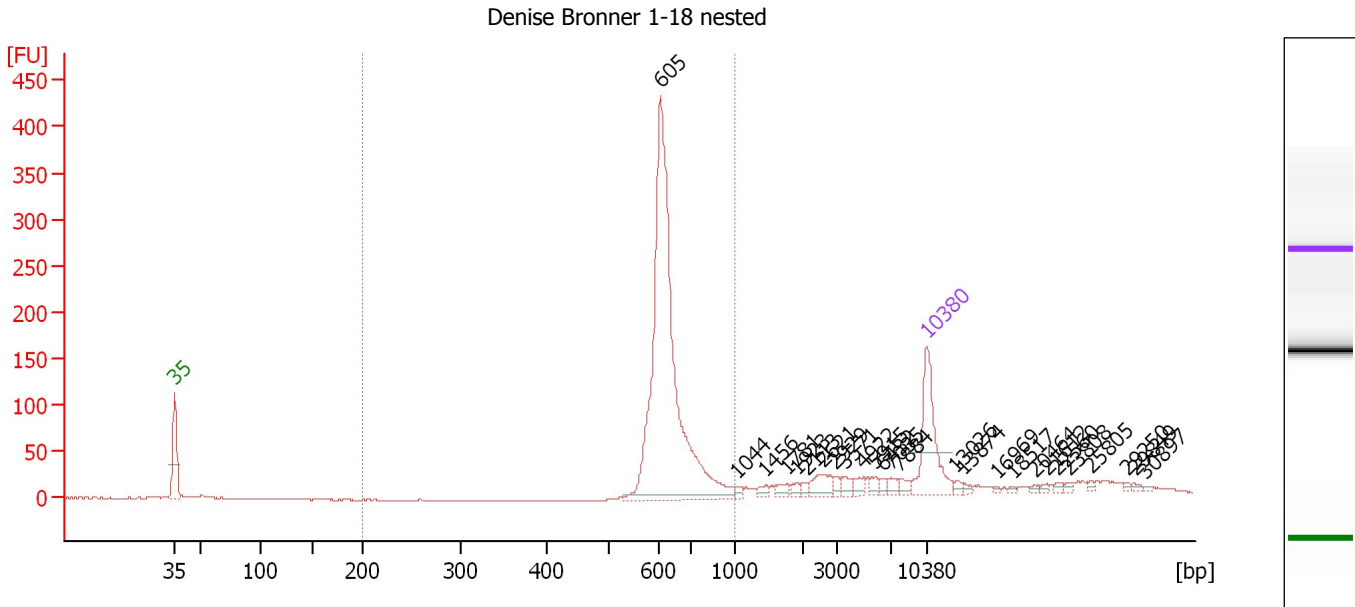
Region table for sample 4 : Lam Tan Khuat 1-22

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	638	988.80	1,716.7	2,369.3	88	10.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Denise Bronner 1-18 nested

Number of peaks found: 26 Corr. Area 1: 1,014.2
 Noise: 0.5

Peak table for sample 5 : Denise Bronner 1-18 nested

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	605	527.77	1,320.8		88.23
3	1,044	4.28	6.2		95.43
4	1,456	6.56	6.8		98.01
5	1,781	6.74	5.7		100.04
6	1,923	5.12	4.0		100.94
7	2,113	4.71	3.4		101.78
8	2,621	18.18	10.5		103.42
9	2,929	5.61	2.9		104.41
10	3,571	8.54	3.6		105.35
11	4,922	8.74	2.7		107.04
12	5,915	6.56	1.7		108.28
13	6,432	4.52	1.1		108.93
14	7,035	7.46	1.6		109.67
15	7,884	6.83	1.3		110.52
16	10,380	75.00	10.9	Upper Marker	113.00
17	13,026	0.00	0.0		115.63
18	13,874	0.00	0.0		116.48
19	16,969	0.00	0.0		119.55
20	18,517	0.00	0.0		121.09
21	20,464	0.00	0.0		123.03
22	21,612	0.00	0.0		124.17
23	22,560	0.00	0.0		125.11
24	23,808	0.00	0.0		126.35
25	25,805	0.00	0.0		128.34


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...**... Peak table for sample 5 : Denise Bronner 1-18 nested**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	29,250	0.00	0.0		131.77
27	29,849	0.00	0.0		132.36
28	30,897	0.00	0.0		133.40

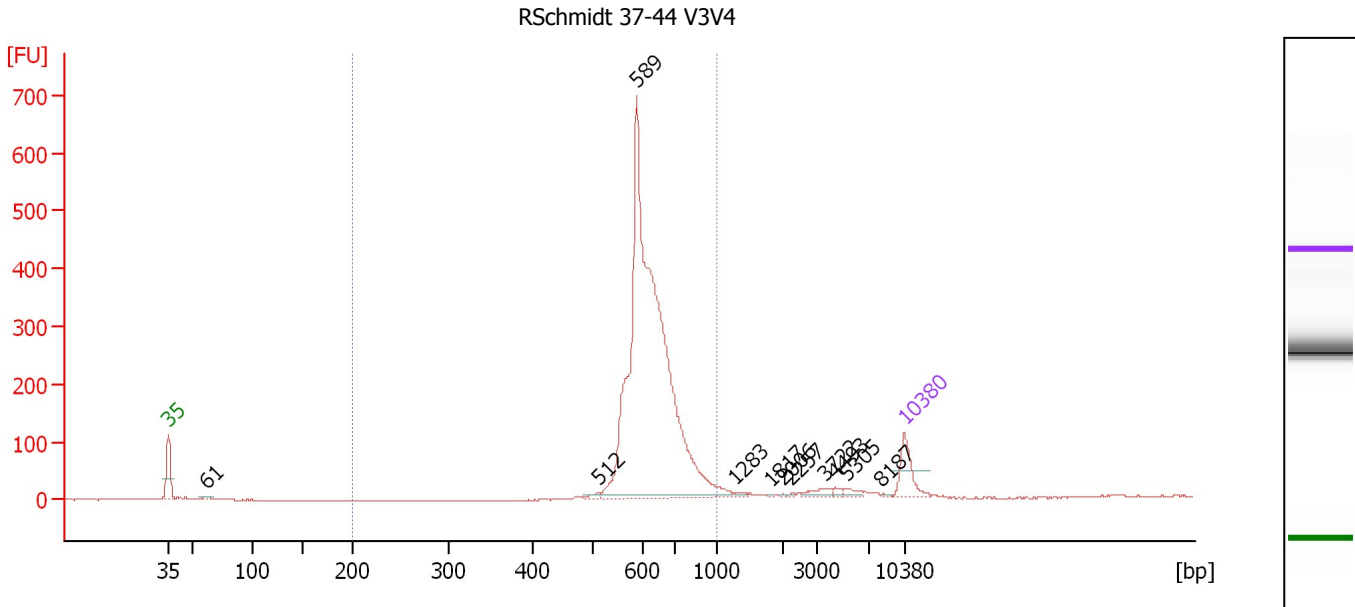
Region table for sample 5 : Denise Bronner 1-18 nested

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	634	501.68	1,014.2	1,208.3	 73	10.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : RSchmidt 37-44 V3V4

Number of peaks found: 11 Corr. Area 1: 2,367.0
 Noise: 0.6

Peak table for sample 6 : RSchmidt 37-44 V3V4

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	61	12.25	304.2		46.60
3	512	10.21	30.2		83.93
4	589	1,991.43	5,124.8		87.54
5	1,283	7.58	9.0		96.92
6	1,817	2.09	1.7		100.27
7	2,006	2.35	1.8		101.43
8	2,257	2.90	1.9		102.25
9	3,722	22.70	9.2		105.54
10	4,493	10.10	3.4		106.51
11	5,305	13.47	3.8		107.52
12	8,187	2.58	0.5		110.82
13	10,380	75.00	10.9	Upper Marker	113.00

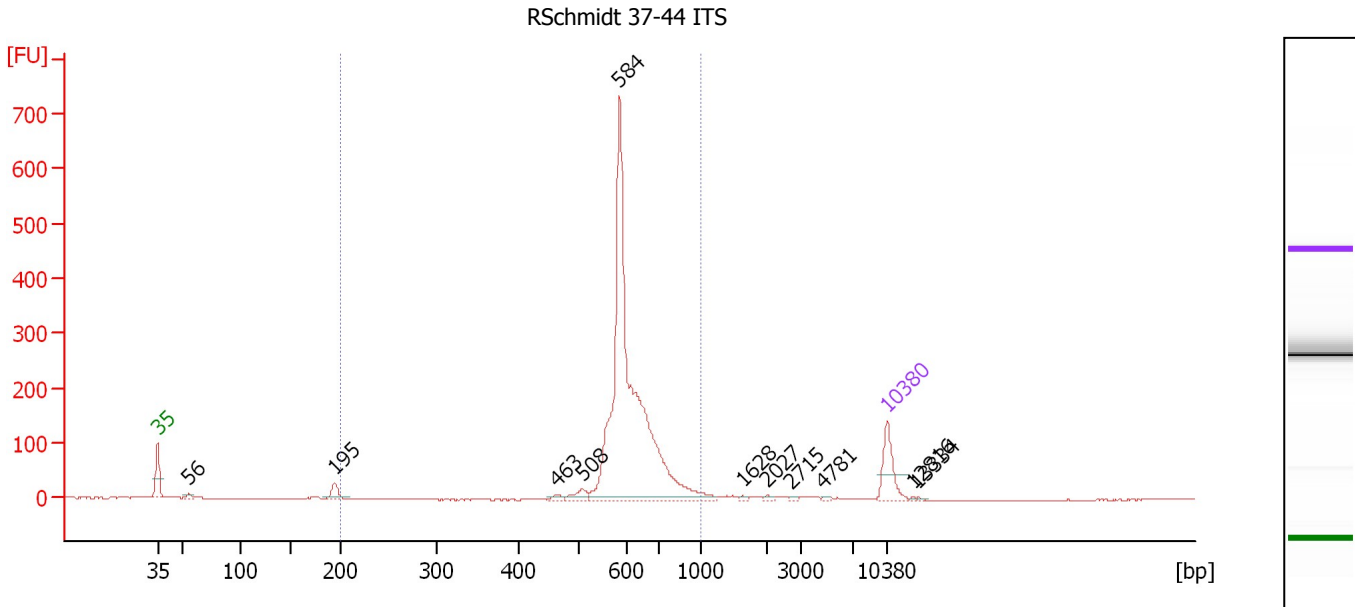
Region table for sample 6 : RSchmidt 37-44 V3V4

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	638	1,972.94	2,367.0	4,744.5	92	12.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : RSchmidt 37-44 ITS

Number of peaks found: 11 Corr. Area 1: 1,608.2
 Noise: 0.6

Peak table for sample 7 : RSchmidt 37-44 ITS

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	56	10.38	283.1		45.99
3	195	35.10	272.8		60.01
4	463	9.67	31.6		81.30
5	508	26.11	77.8		83.77
6	584	1,156.24	2,998.0		87.33
7	1,628	3.55	3.3		99.08
8	2,027	5.28	3.9		101.51
9	2,715	3.64	2.0		103.72
10	4,781	2.49	0.8		106.87
11	10,380	75.00	10.9	Upper Marker	113.00
12	12,816	0.00	0.0		115.42
13	13,334	0.00	0.0		115.94

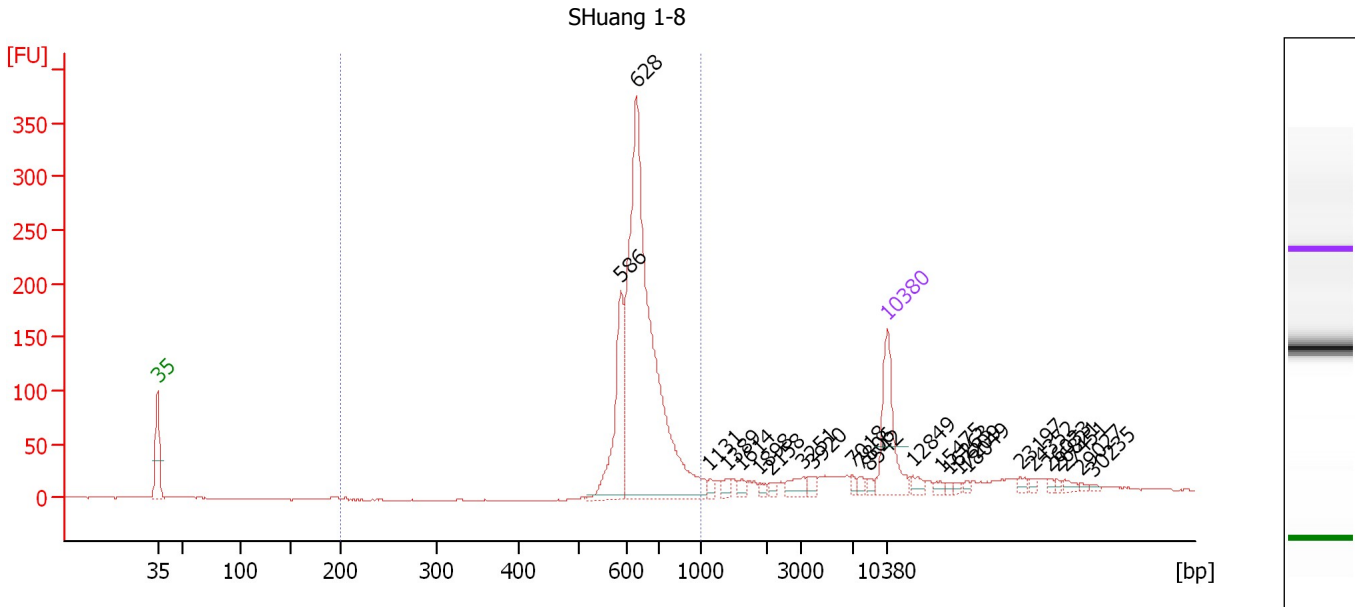
Region table for sample 7 : RSchmidt 37-44 ITS

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	620	1,136.57	1,608.1	2,819.8	94	12.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : SHuang 1-8

Number of peaks found: 24 Corr. Area 1: 1,216.6
 Noise: 0.4

Peak table for sample 8 : SHuang 1-8

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	586	131.86	341.0		87.40
3	628	608.83	1,468.6		88.92
4	1,131	7.01	9.4		95.97
5	1,389	7.96	8.7		97.59
6	1,614	6.83	6.4		99.00
7	1,898	3.92	3.1		100.78
8	2,158	5.15	3.6		101.93
9	3,251	15.15	7.1		104.96
10	3,920	7.95	3.1		105.79
11	7,018	5.18	1.1		109.66
12	7,806	5.53	1.1		110.44
13	8,542	5.60	1.0		111.17
14	10,380	75.00	10.9	Upper Marker	113.00
15	12,849	0.00	0.0		115.46
16	15,475	0.00	0.0		118.07
17	16,263	0.00	0.0		118.85
18	17,209	0.00	0.0		119.79
19	18,049	0.00	0.0		120.63
20	23,197	0.00	0.0		125.75
21	24,352	0.00	0.0		126.90
22	26,033	0.00	0.0		128.57
23	26,821	0.00	0.0		129.35
24	27,451	0.00	0.0		129.98
25	29,027	0.00	0.0		131.54

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad


Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Electropherogram Summary Continued ...

... Peak table for sample 8 : SHuang 1-8

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	30,235	0.00	0.0		132.75

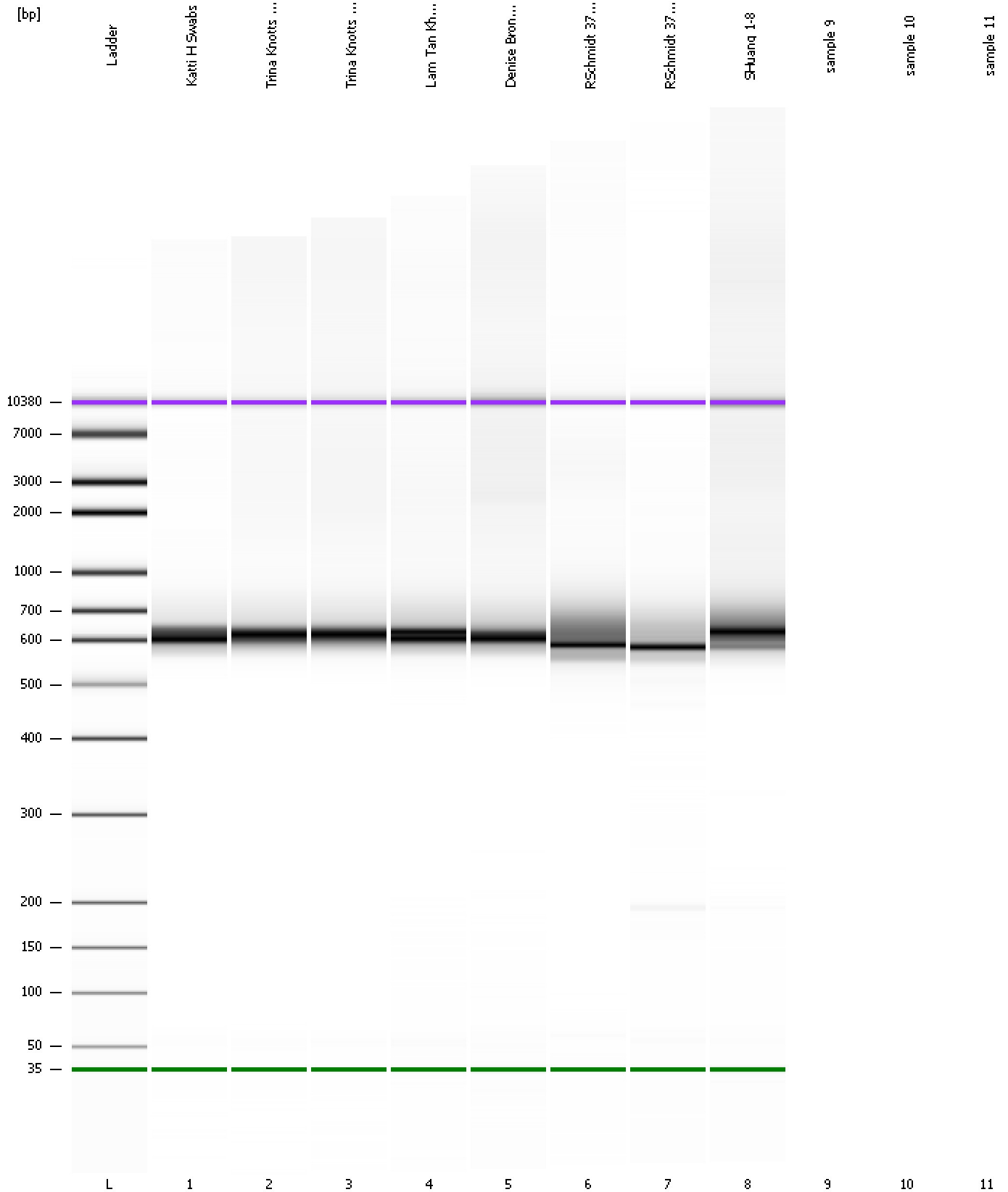
Region table for sample 8 : SHuang 1-8

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	653	695.27	1,216.6	1,631.8	 77	11.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 PM

Gel Image

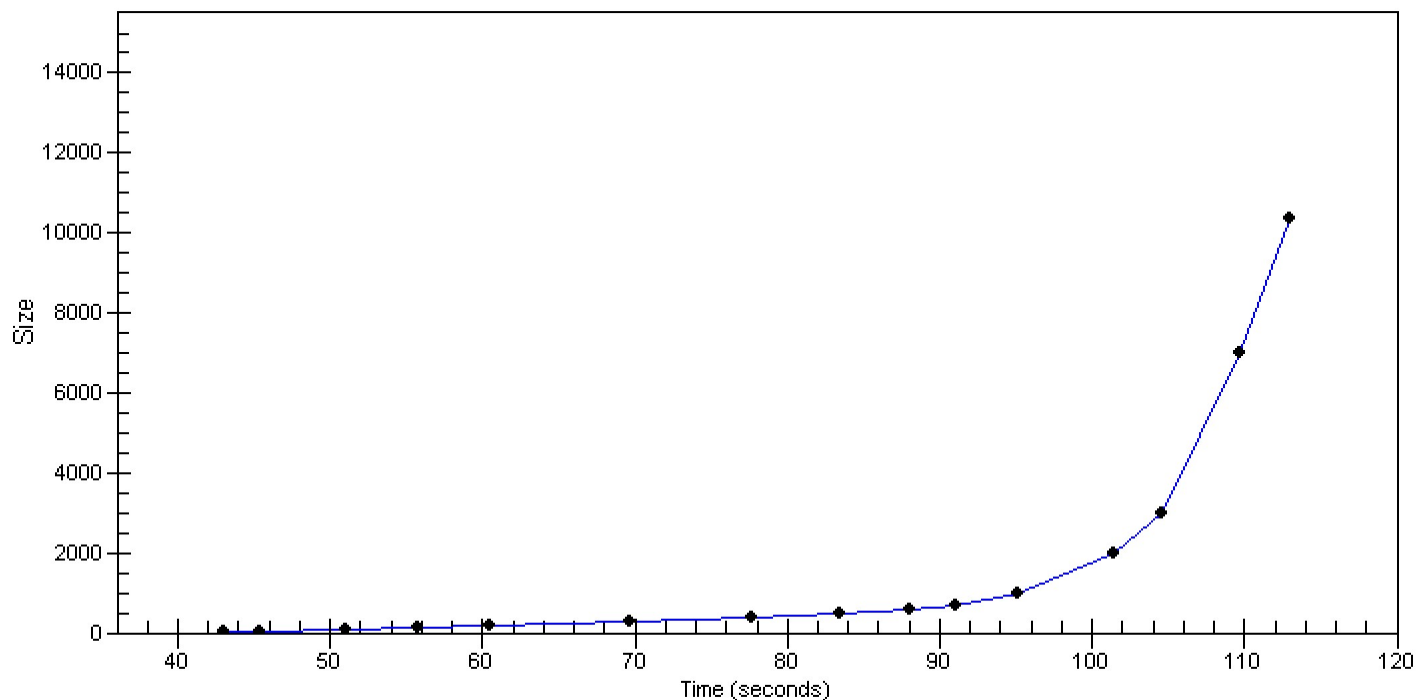


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
Modified: 3/20/2019 2:20:29 PM

Invalid Samples

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-20\2019-03-20_001.xad

Created: 3/20/2019 1:47:42 PM
 Modified: 3/20/2019 2:20:29 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 9)		Instrument	Run		3/20/2019 2:20:24 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\2019-03-20\2019-03-20_001.xad)		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/20/2019 1:47:48 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1