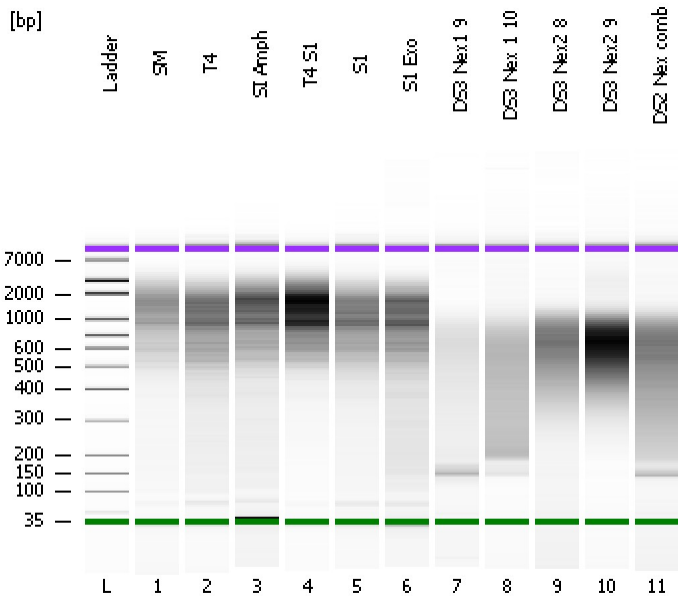


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 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

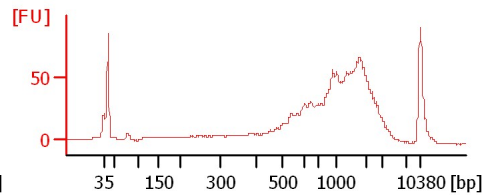
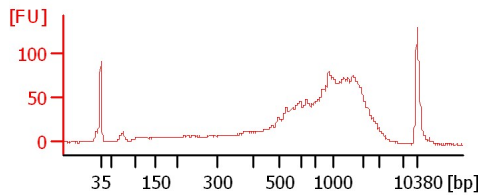
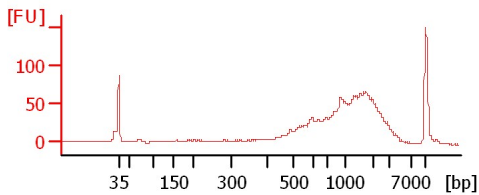
Chip Information:

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

SM

T4

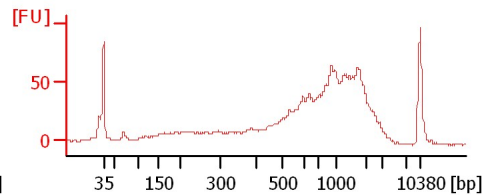
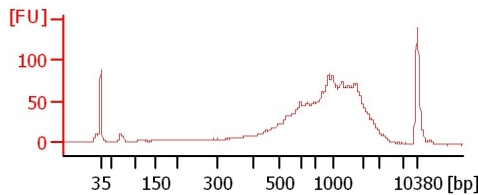
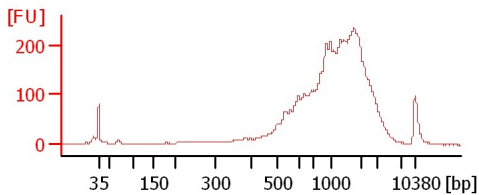
SI Amph



T4 S1

S1

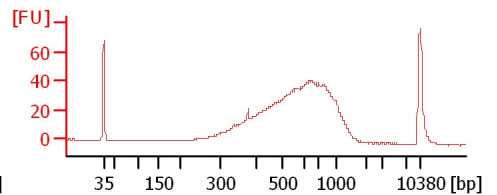
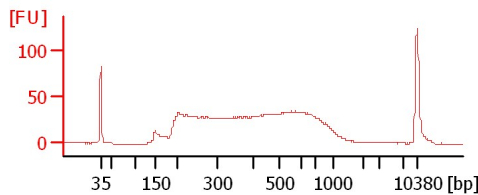
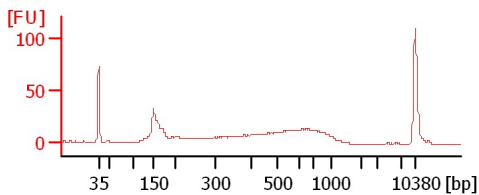
S1 Exo



DS3 Nex1 9

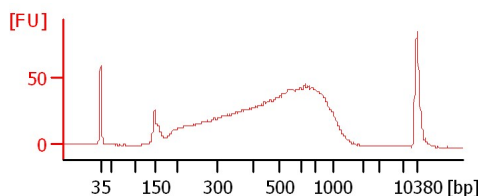
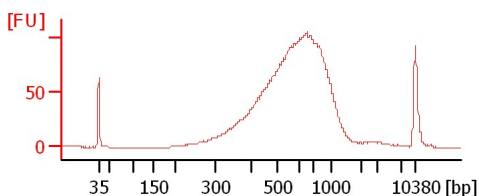
DS3 Nex1 10

DS3 Nex2 8



DS3 Nex2 9

DS2 Nex comb



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
SM		<input type="checkbox"/>	✓			
T4		<input type="checkbox"/>	✓			
SI Amph		<input type="checkbox"/>	✓			
T4 S1		<input type="checkbox"/>	✓			
S1		<input type="checkbox"/>	✓			
S1 Exo		<input type="checkbox"/>	✓			
DS3 Nex1 9		<input type="checkbox"/>	✓			
DS3 Nex 1 10		<input type="checkbox"/>	✓			
DS3 Nex2 8		<input type="checkbox"/>	✓			
DS3 Nex2 9		<input type="checkbox"/>	✓			
DS2 Nex comb		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

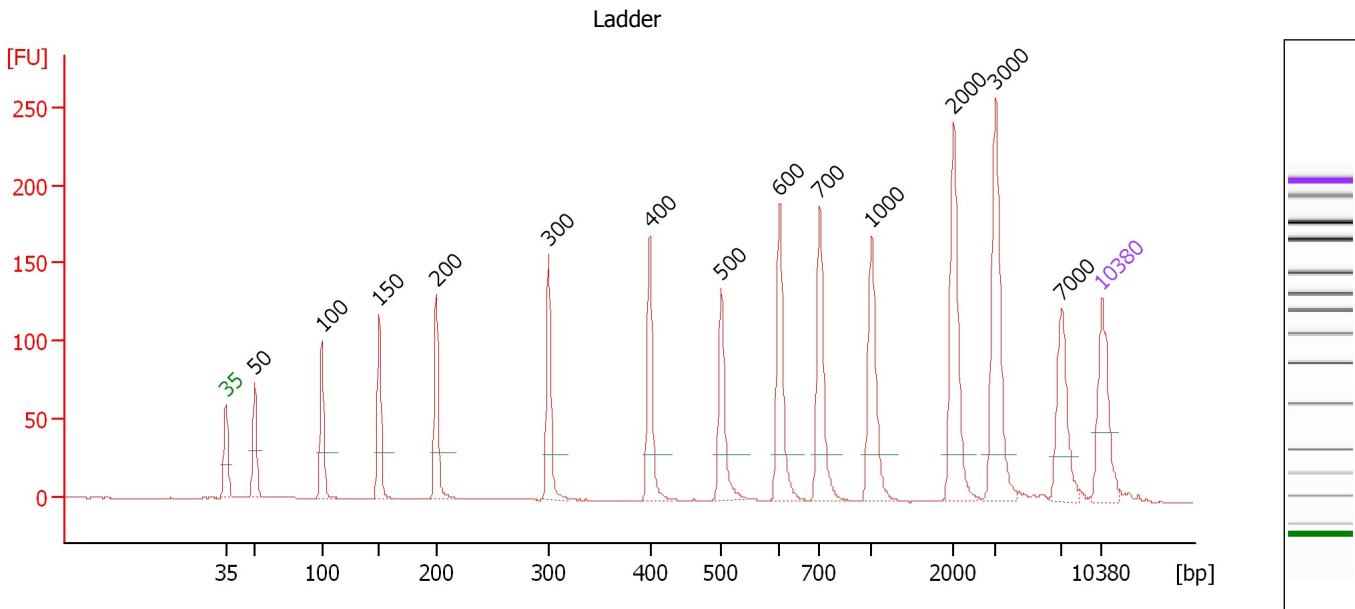
Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
 Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

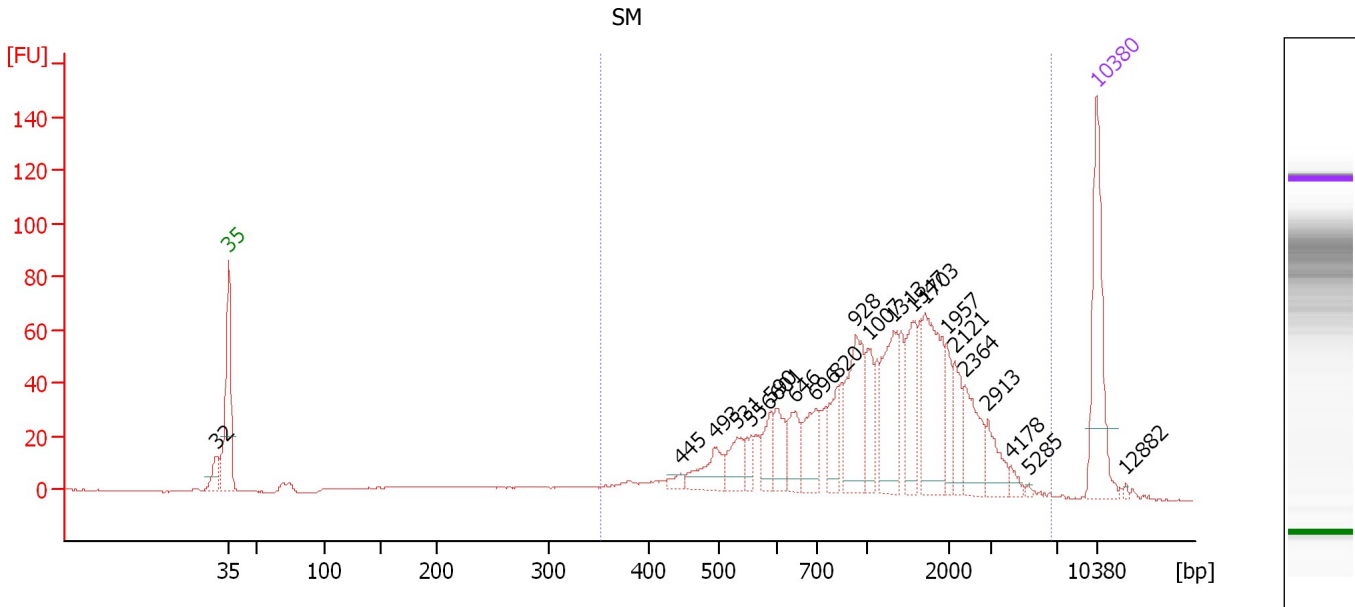
Peak table for Ladder

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	45.30
3	100	150.00	2,272.7	Ladder Peak	50.67
4	150	150.00	1,515.2	Ladder Peak	55.27
5	200	150.00	1,136.4	Ladder Peak	59.79
6	300	150.00	757.6	Ladder Peak	68.78
7	400	150.00	568.2	Ladder Peak	76.87
8	500	150.00	454.5	Ladder Peak	82.58
9	600	150.00	378.8	Ladder Peak	87.22
10	700	150.00	324.7	Ladder Peak	90.46
11	1,000	150.00	227.3	Ladder Peak	94.55
12	2,000	150.00	113.6	Ladder Peak	101.16
13	3,000	150.00	75.8	Ladder Peak	104.52
14	7,000	150.00	32.5	Ladder Peak	109.80
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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 Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : SM

Number of peaks found: 22 Corr. Area 1: 1,101.2
 Noise: 0.2

Peak table for sample 1 : SM

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.01
2	35	125.00	5,411.3	Lower Marker	43.00
3	445	7.08	24.1		79.46
4	493	32.70	100.5		82.17
5	531	27.87	79.5		84.02
6	556	14.89	40.6		85.18
7	590	22.08	56.7		86.77
8	601	27.02	68.2		87.24
9	646	28.76	67.5		88.71
10	696	38.63	84.1		90.34
11	820	31.22	57.7		92.10
12	928	70.01	114.4		93.57
13	1,007	29.48	44.4		94.60
14	1,313	60.83	70.2		96.62
15	1,547	40.66	39.8		98.17
16	1,703	77.33	68.8		99.20
17	1,957	19.12	14.8		100.87
18	2,121	26.26	18.8		101.56
19	2,364	37.22	23.9		102.38
20	2,913	22.14	11.5		104.23
21	4,178	5.50	2.0		106.08
22	5,285	1.29	0.4		107.54
23	10,380	75.00	10.9	Upper Marker	113.00
24	12,882	0.00	0.0		115.36


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...

... Region table for sample 1 :

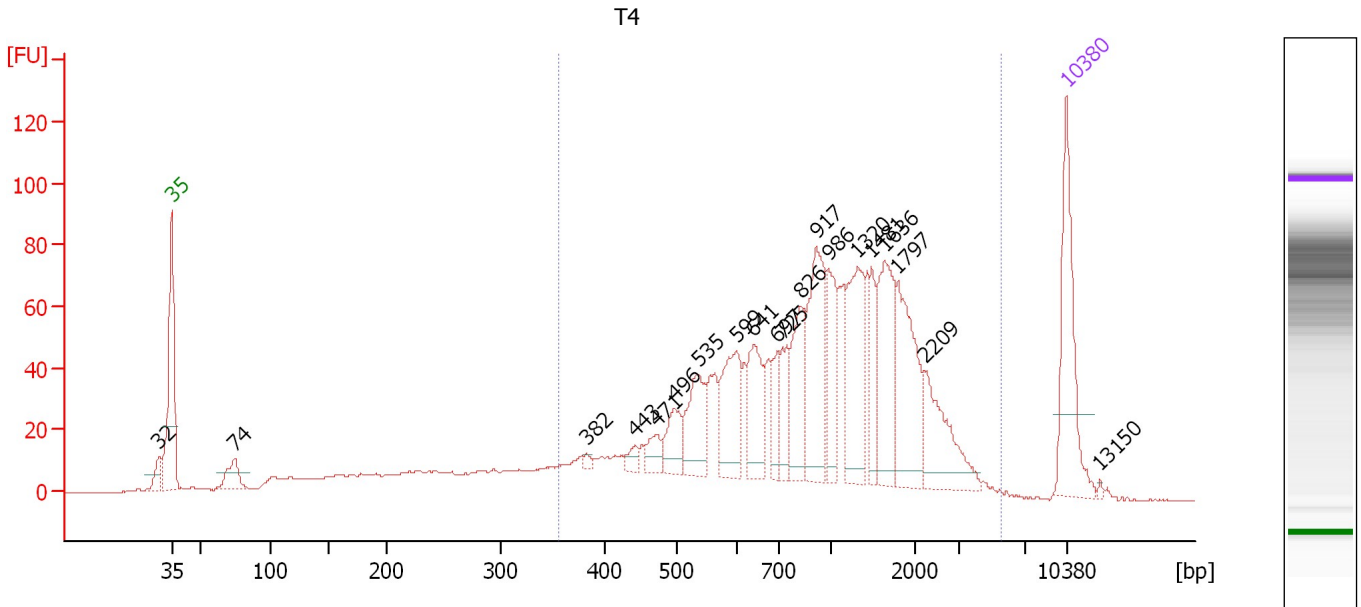
SM

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
352	6,684	1,361	1,101.2	1,329.1	 790.30	90	62.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
 Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : T4

Number of peaks found: 20 Corr. Area 1: 1,445.5
 Noise: 0.2

Peak table for sample 2 : T4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.02
2	35	125.00	5,411.3	Lower Marker	43.00
3	74	28.54	580.7		47.93
4	382	4.23	16.8		75.40
5	443	9.19	31.5		79.31
6	471	16.03	51.6		80.91
7	496	30.82	94.2		82.33
8	535	55.01	155.9		84.19
9	599	63.26	160.1		87.17
10	641	51.09	120.8		88.54
11	697	24.01	52.2		90.36
12	725	32.12	67.1		90.81
13	826	57.23	105.0		92.18
14	917	91.57	151.2		93.42
15	986	48.14	74.0		94.36
16	1,320	76.95	88.3		96.67
17	1,481	28.63	29.3		97.73
18	1,636	74.43	68.9		98.75
19	1,797	79.75	67.2		99.82
20	2,209	56.19	38.5		101.86
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,150	0.00	0.0		115.62

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...

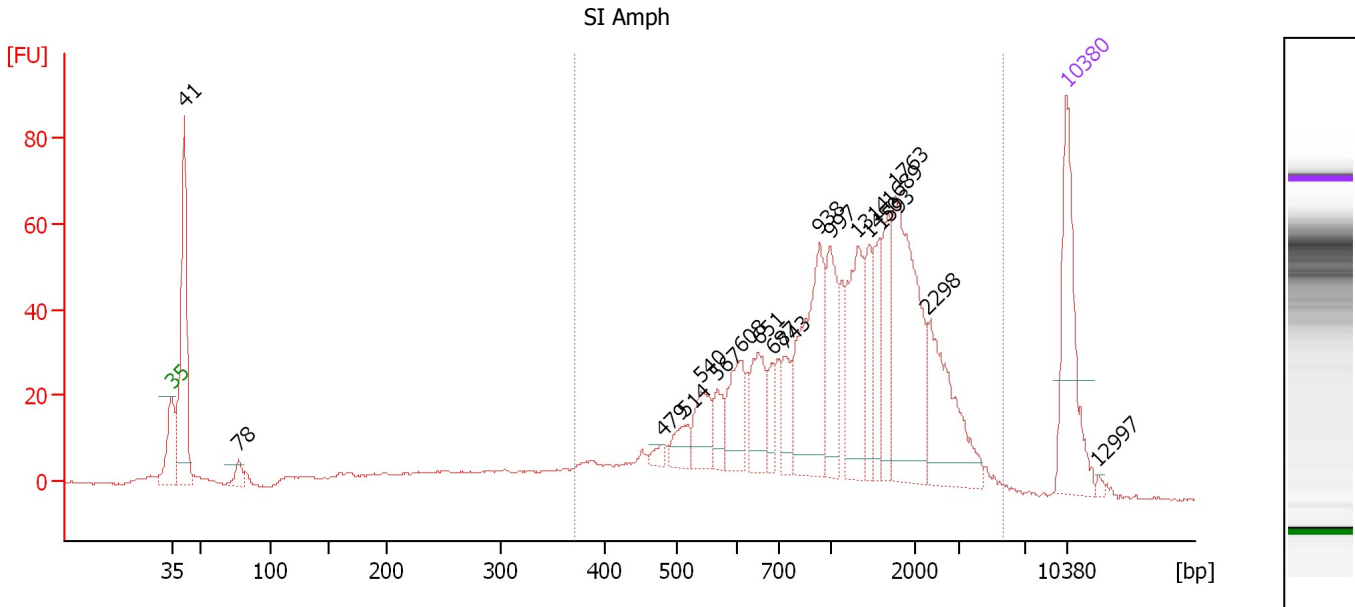
... Region table for sample 2 : T4

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
356	5,514	1,157	1,445.5	2,166.4	1,123.78	82	61.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
 Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : SI Amph

Number of peaks found: 19 Corr. Area 1: 1,050.5
 Noise: 0.2

Peak table for sample 3 : SI Amph

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	41	184.05	6,727.1		43.99
3	78	13.65	266.5		48.26
4	479	6.86	21.7		81.37
5	514	18.86	55.6		83.22
6	540	32.58	91.4		84.43
7	567	19.87	53.1		85.69
8	608	41.92	104.4		87.49
9	651	38.89	90.5		88.89
10	687	16.78	37.0		90.06
11	743	25.91	52.9		91.05
12	938	105.58	170.6		93.70
13	997	52.06	79.1		94.51
14	1,314	69.26	79.9		96.62
15	1,450	27.81	29.1		97.52
16	1,593	32.20	30.6		98.47
17	1,689	39.11	35.1		99.10
18	1,763	124.22	106.7		99.59
19	2,298	72.19	47.6		102.16
20	10,380	75.00	10.9	Upper Marker	113.00
21	12,997	0.00	0.0		115.47

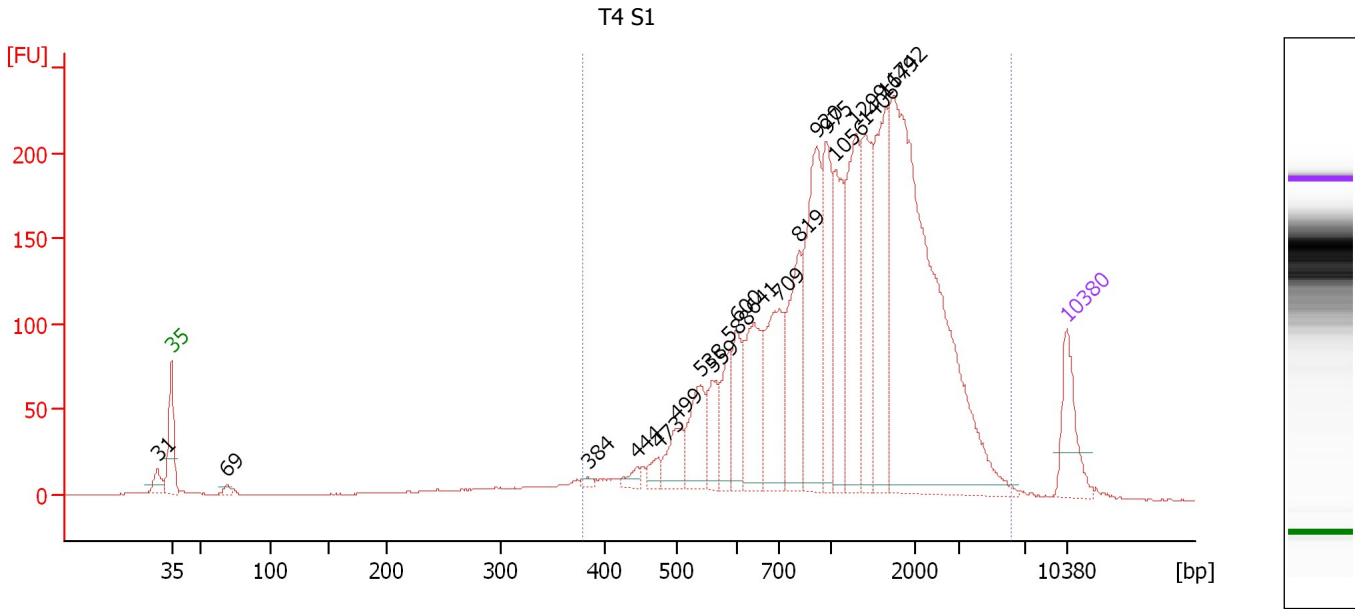
Region table for sample 3 : SI Amph

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
371	5,651	1,340	1,050.5	1,603.2	950.01	79	60.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
 Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : T4 S1

Number of peaks found: 20 Corr. Area 1: 3,562.1
 Noise: 0.2

Peak table for sample 4 : T4 S1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.91
2	35	125.00	5,411.3	Lower Marker	43.00
3	69	10.26	225.3		47.34
4	384	8.00	31.6		75.57
5	444	15.54	53.0		79.39
6	473	22.35	71.6		81.04
7	499	60.93	185.1		82.51
8	538	100.26	282.3		84.35
9	559	69.69	188.8		85.34
10	588	68.75	177.2		86.66
11	600	81.62	206.1		87.23
12	641	141.61	334.8		88.55
13	709	188.19	402.4		90.58
14	819	158.24	292.6		92.09
15	920	252.13	415.3		93.46
16	975	147.86	229.7		94.21
17	1,056	150.24	215.6		94.92
18	1,299	192.95	225.1		96.53
19	1,406	165.25	178.0		97.23
20	1,649	196.30	180.3		98.84
21	1,742	758.85	660.0		99.45
22	10,380	75.00	10.9	Upper Marker	113.00


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...

... Region table for sample 4 :

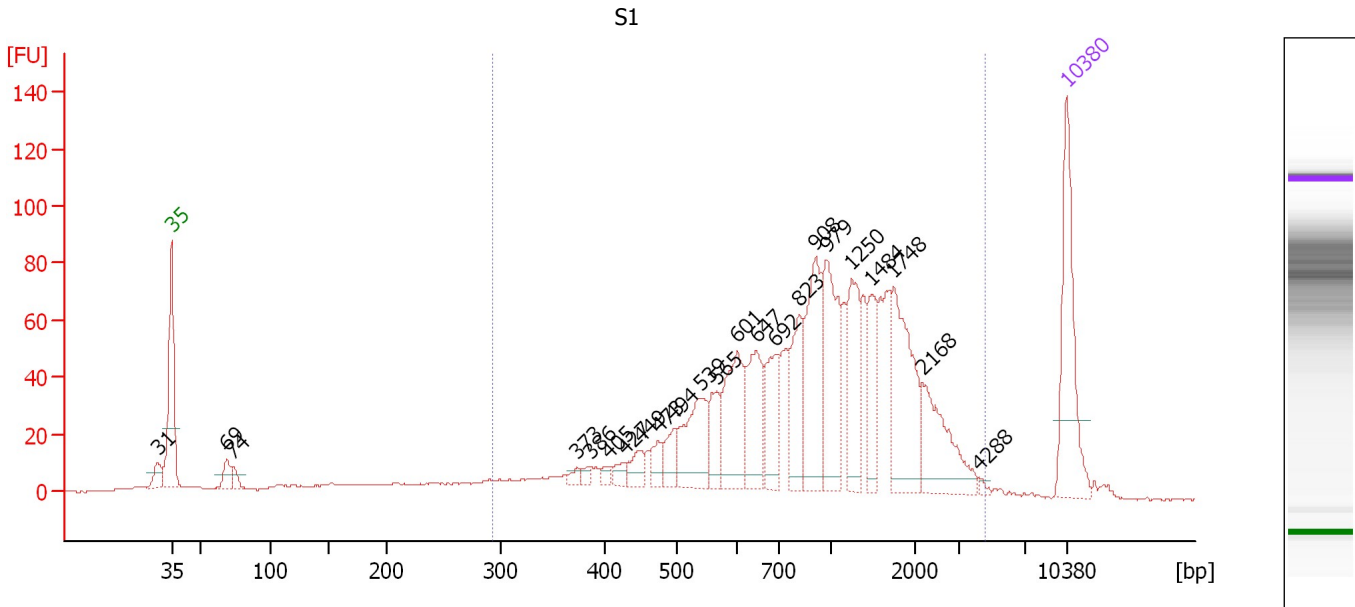
T4 S1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
378	6,088	1,401	3,562.1	4,582.7	 2,923.02	95	59.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
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Electropherogram Summary Continued ...



Overall Results for sample 5 : S1

Number of peaks found: 23 Corr. Area 1: 1,428.1
 Noise: 0.2

Peak table for sample 5 : S1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.93
2	35	125.00	5,411.3	Lower Marker	43.00
3	69	16.37	358.7		47.36
4	74	11.12	228.8		47.84
5	373	5.77	23.4		74.71
6	386	5.63	22.1		75.77
7	405	6.07	22.7		77.18
8	427	8.59	30.5		78.39
9	449	15.50	52.3		79.69
10	473	15.54	49.8		81.05
11	494	19.83	60.8		82.26
12	539	62.40	175.4		84.39
13	565	29.12	78.1		85.60
14	601	69.67	175.7		87.25
15	647	56.36	132.0		88.75
16	692	47.27	103.5		90.20
17	823	51.56	94.9		92.14
18	908	82.38	137.5		93.30
19	979	82.23	127.3		94.27
20	1,250	58.08	70.4		96.20
21	1,484	33.49	34.2		97.75
22	1,748	85.69	74.3		99.49
23	2,168	56.94	39.8		101.72
24	4,288	2.15	0.8		106.22
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

Created: 6/22/2017 9:33:59 AM
Modified: 6/22/2017 1:08:11 PM

Electropherogram Summary Continued ...

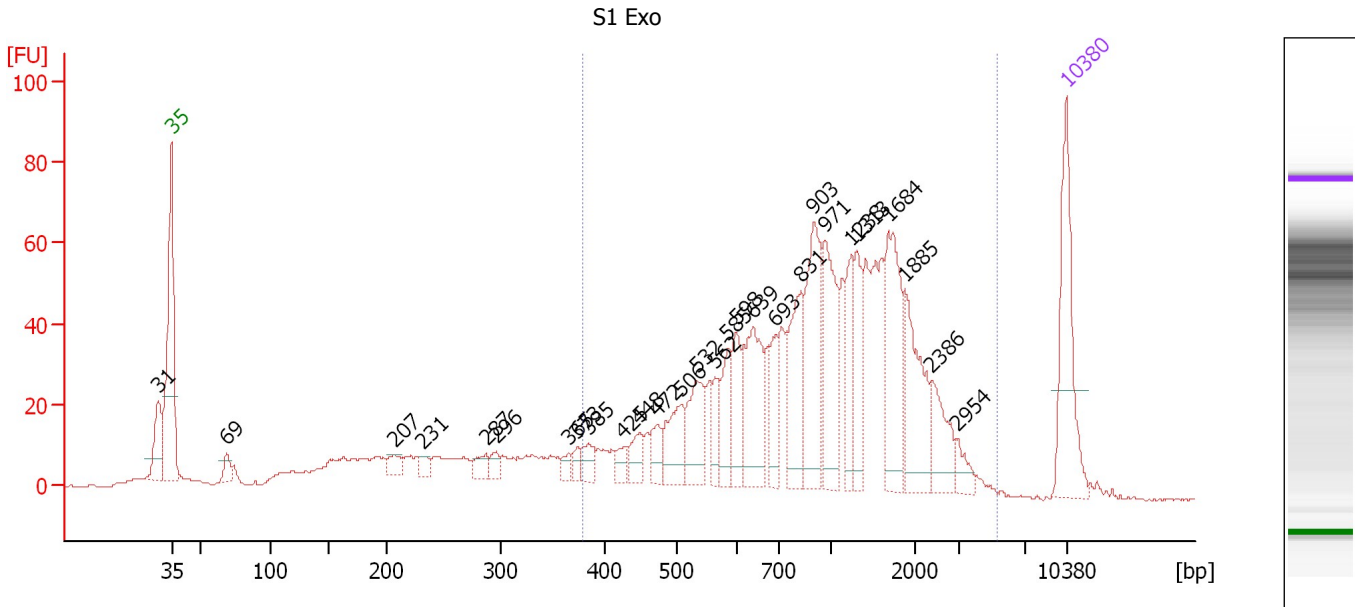
... Region table for sample 5 : S1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
293	4,656	1,132	1,428.1	2,075.7	1,046.77	88	59.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : S1 Exo

Number of peaks found: 28 Corr. Area 1: 1,103.7
 Noise: 0.2

Peak table for sample 6 : S1 Exo

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.97
2	35	125.00	5,411.3	Lower Marker	43.00
3	69	14.57	321.0		47.32
4	207	13.54	99.1		60.41
5	231	9.23	60.5		62.57
6	287	13.53	71.5		67.58
7	296	10.66	54.6		68.41
8	365	7.27	30.2		74.05
9	373	8.36	33.9		74.69
10	385	15.79	62.2		75.62
11	425	11.51	41.0		78.32
12	448	19.58	66.3		79.59
13	472	18.91	60.7		80.97
14	506	44.60	133.4		82.88
15	532	48.93	139.4		84.06
16	562	23.71	63.9		85.48
17	585	34.57	89.6		86.51
18	598	39.42	99.8		87.15
19	639	77.88	184.8		88.47
20	693	29.46	64.4		90.24
21	831	68.28	124.5		92.25
22	903	93.34	156.6		93.23
23	971	76.48	119.3		94.16
24	1,238	38.69	47.3		96.13
25	1,313	36.15	41.7		96.62

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...

... Peak table for sample 6 : S1 Exo

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,684	79.81	71.8		99.07
27	1,885	64.33	51.7		100.39
28	2,386	35.37	22.5		102.45
29	2,954	11.85	6.1		104.37
30	10,380	75.00	10.9	Upper Marker	113.00

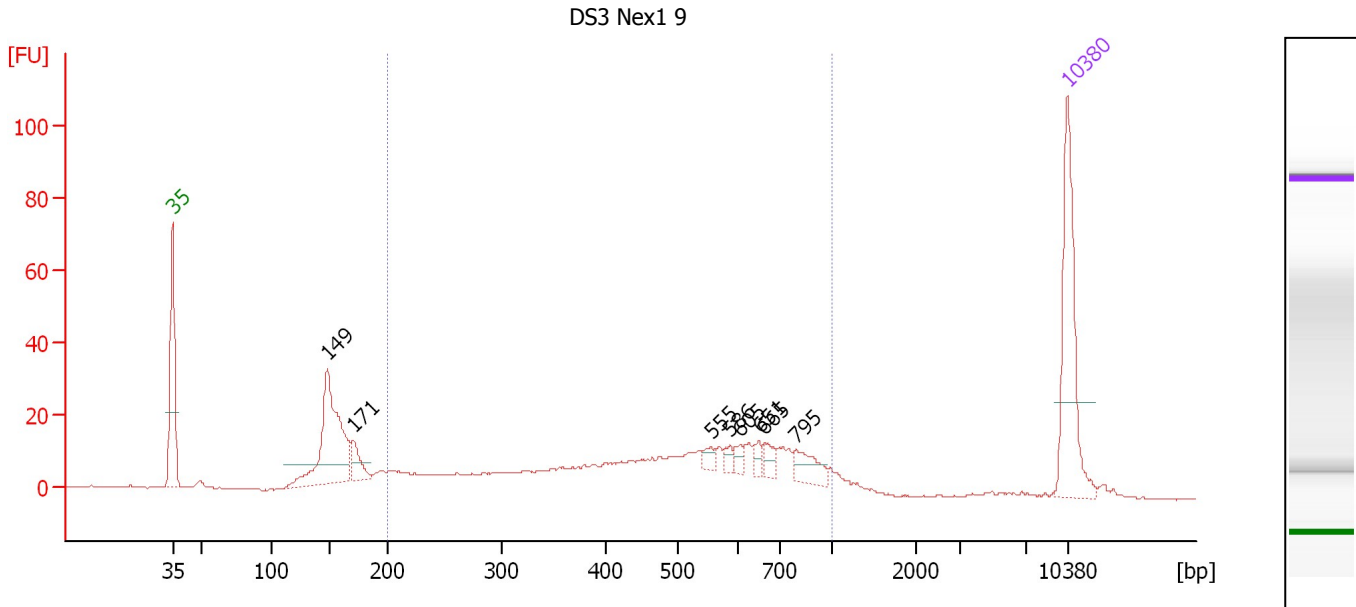
Region table for sample 6 : S1 Exo

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
379	5,312	1,151	1,103.7	2,153.1	1,143.87	75	58.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : DS3 Nex1 9

Number of peaks found: 8 Corr. Area 1: 360.4
 Noise: 0.2

Peak table for sample 7 : DS3 Nex1 9

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	149	137.96	1,406.9		55.14
3	171	22.83	202.1		57.18
4	555	6.94	19.0		85.14
5	586	6.14	15.9		86.58
6	605	7.22	18.1		87.38
7	651	7.48	17.4		88.87
8	665	9.80	22.3		89.32
9	795	19.79	37.7		91.76
10	10,380	75.00	10.9	Upper Marker	113.00

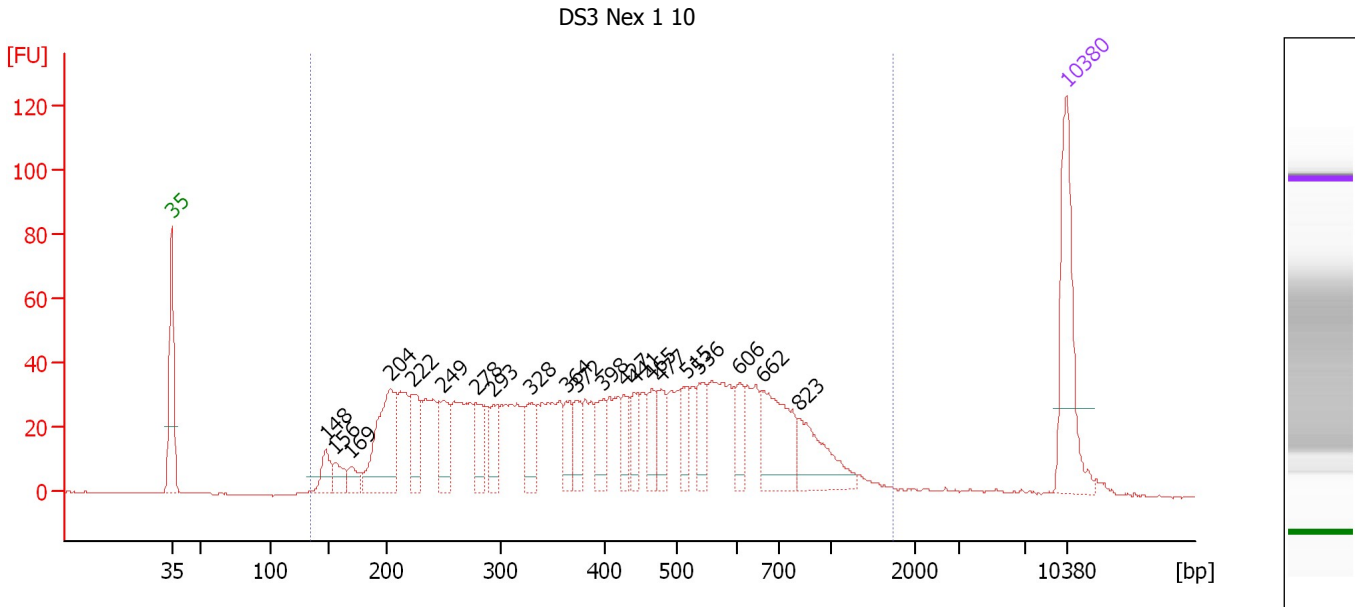
Region table for sample 7 : DS3 Nex1 9

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	509	360.4	1,440.8	385.25	66	37.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : DS3 Nex 1 10

Number of peaks found: 21 Corr. Area 1: 1,470.4
 Noise: 0.2

Peak table for sample 8 : DS3 Nex 1 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	148	25.61	262.2		55.08
3	156	18.84	183.3		55.79
4	169	16.55	148.4		56.99
5	204	104.31	774.6		60.15
6	222	47.01	320.2		61.80
7	249	41.23	250.7		64.21
8	278	33.28	181.2		66.82
9	293	25.63	132.4		68.17
10	328	38.45	177.4		71.08
11	364	24.14	100.5		73.94
12	372	23.90	97.2		74.64
13	398	27.60	105.0		76.75
14	427	21.61	76.7		78.40
15	441	21.86	75.1		79.20
16	465	23.75	77.5		80.56
17	477	26.88	85.4		81.26
18	515	23.33	68.7		83.27
19	536	28.41	80.2		84.27
20	606	24.06	60.1		87.43
21	662	70.28	160.9		89.23
22	823	51.21	94.3		92.14
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...

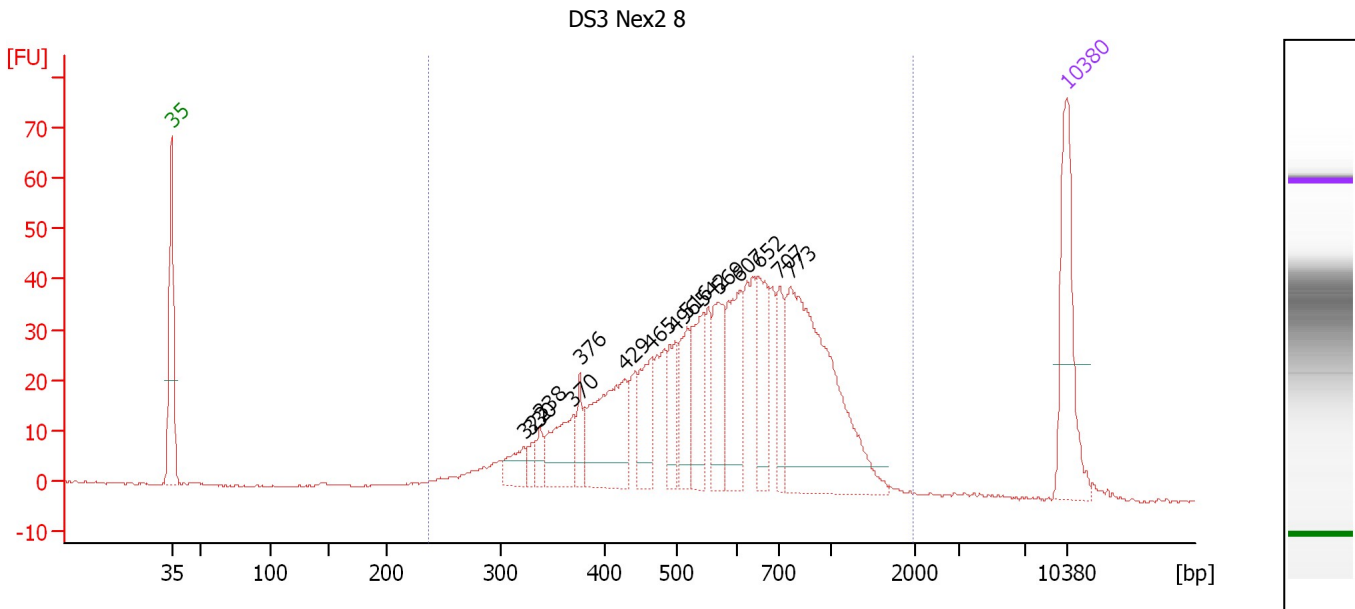
... Region table for sample 8 : DS3 Nex 1 10

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
135	1,735	455	1,470.4	6,402.4	1,391.17	98	51.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : DS3 Nex2 8

Number of peaks found: 15 Corr. Area 1: 862.3
 Noise: 0.2

Peak table for sample 9 : DS3 Nex2 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	322	24.30	114.3		70.57
3	330	11.02	50.7		71.17
4	338	14.32	64.3		71.82
5	370	53.01	217.1		74.43
6	376	22.51	90.7		74.94
7	429	107.51	380.1		78.50
8	465	50.58	164.9		80.56
9	496	36.90	112.6		82.37
10	516	46.11	135.4		83.32
11	542	53.26	148.9		84.53
12	569	56.13	149.5		85.78
13	607	74.23	185.4		87.44
14	652	58.26	135.5		88.90
15	707	35.98	77.1		90.55
16	773	234.55	459.7		91.46
17	10,380	75.00	10.9	Upper Marker	113.00

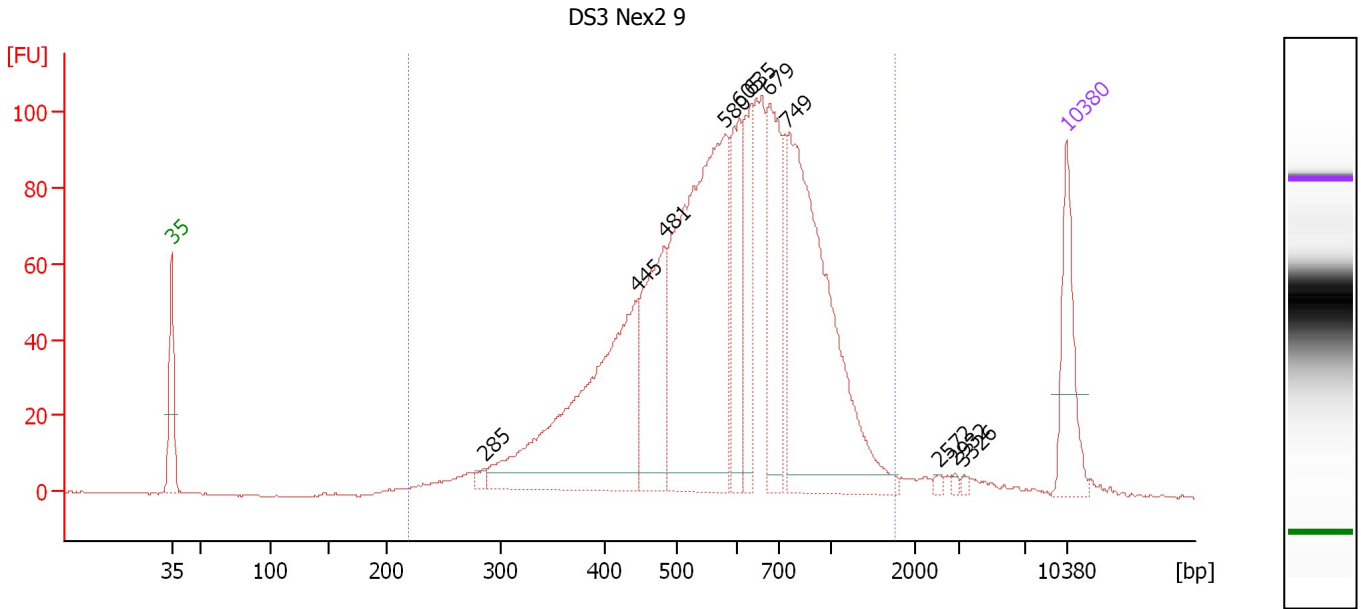
Region table for sample 9 : DS3 Nex2 8

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
238	1,964	624	862.3	3,218.5	1,113.75	99	38.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : DS3 Nex2 9

Number of peaks found: 11 Corr. Area 1: 1,920.4
 Noise: 0.2

Peak table for sample 10 : DS3 Nex2 9

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	285	9.00	47.8		67.45
3	445	416.80	1,419.0		79.44
4	481	194.23	611.3		81.52
5	580	512.48	1,339.7		86.28
6	605	124.66	312.2		87.39
7	635	103.84	247.9		88.35
8	679	161.06	359.6		89.77
9	749	463.15	936.5		91.13
10	2,572	3.69	2.2		103.08
11	2,932	2.92	1.5		104.29
12	3,326	2.74	1.2		104.95
13	10,380	75.00	10.9	Upper Marker	113.00

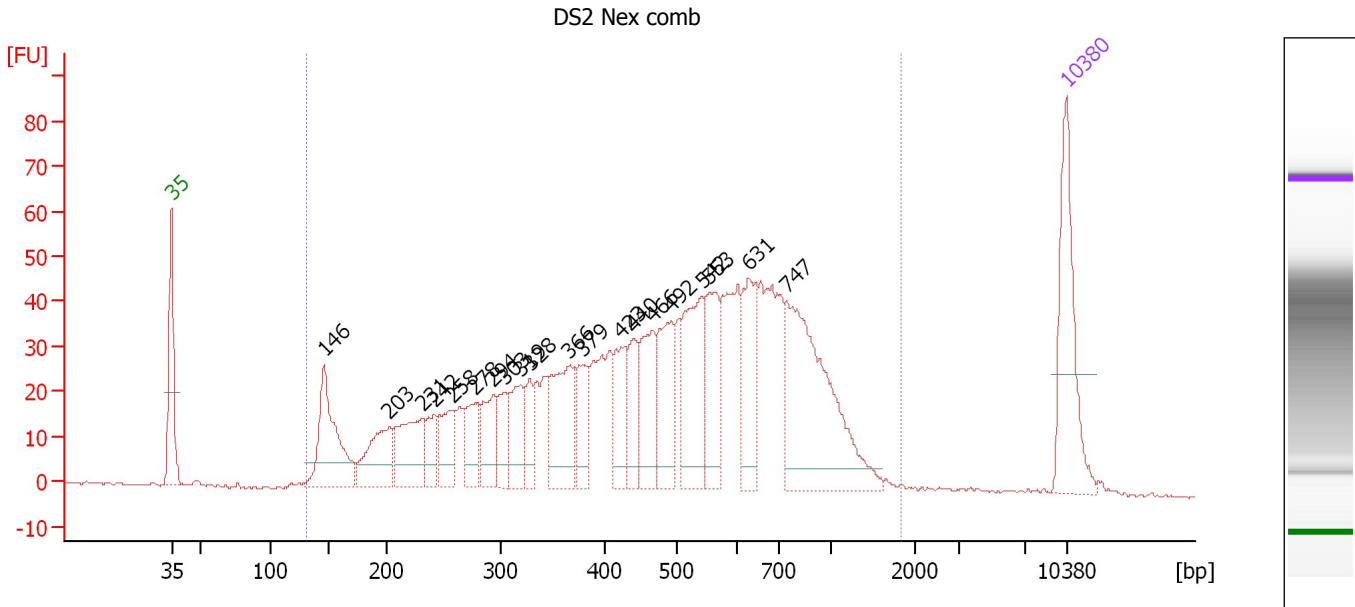
Region table for sample 10 : DS3 Nex2 9

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
220	1,763	630	1,920.4	6,425.6	2,282.08	97	36.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : DS2 Nex comb

Number of peaks found: 20 Corr. Area 1: 1,418.2
 Noise: 0.2

Peak table for sample 11 : DS2 Nex comb

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	146	111.30	1,154.3		54.91
3	203	63.40	473.8		60.03
4	231	72.02	471.4		62.62
5	242	25.71	161.2		63.53
6	258	42.89	251.9		65.00
7	278	41.64	227.2		66.77
8	294	44.67	230.6		68.19
9	303	29.78	148.7		69.05
10	319	45.70	217.0		70.32
11	328	25.44	117.6		71.03
12	366	77.97	322.7		74.12
13	379	38.71	154.7		75.19
14	423	48.01	172.0		78.18
15	440	38.55	132.8		79.14
16	466	60.50	196.5		80.66
17	492	67.70	208.4		82.13
18	542	91.17	255.0		84.51
19	553	63.90	175.2		85.02
20	631	65.54	157.5		88.21
21	747	185.54	376.3		91.10
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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Electropherogram Summary Continued ...

... Region table for sample 11 : DS2 Nex comb

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
131	1,840	518	1,418.2	6,764.1	■ 1,619.53	99	47.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-06-22\2017-06-22_001.xad

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