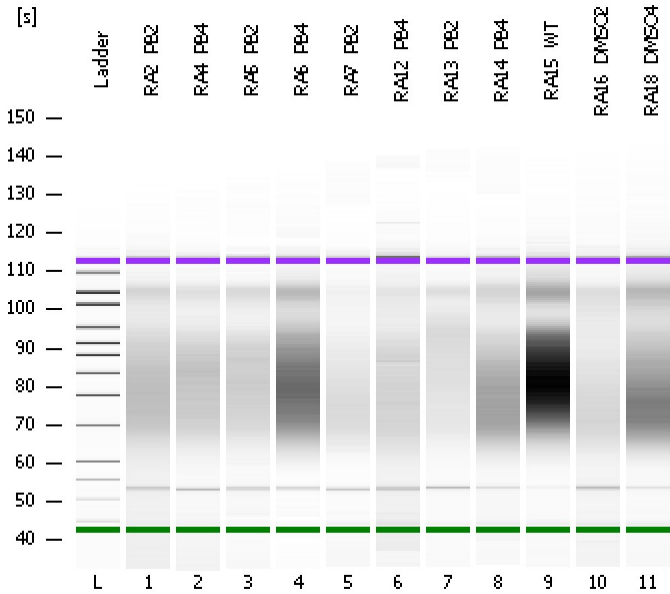


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
Modified: 1/5/2018 1:21:00 PM

Electrophoresis File Run Summary



Instrument Information:

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

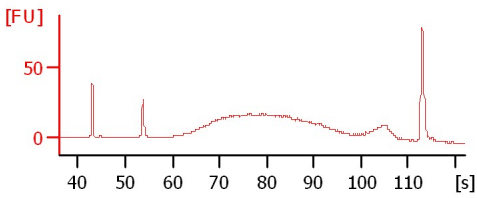
Assay Information:

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

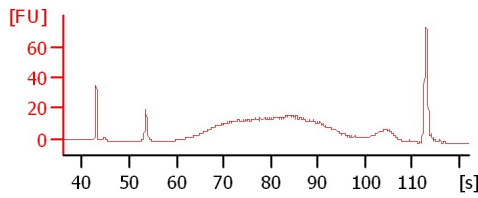
Chip Information:

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

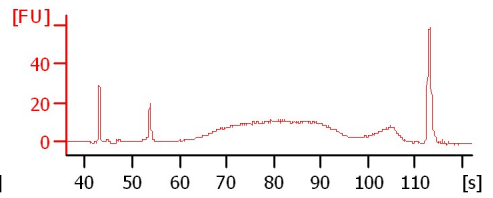
RA2_PB2



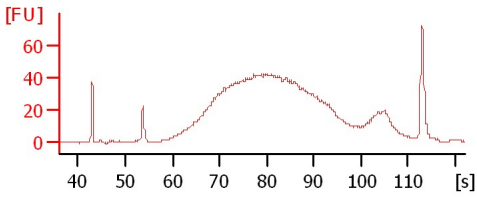
RA4_PB4



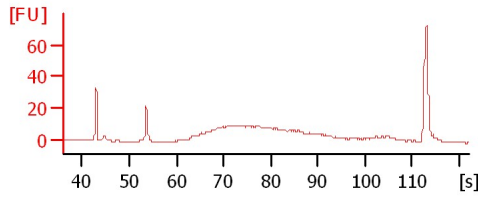
RA5_PB2



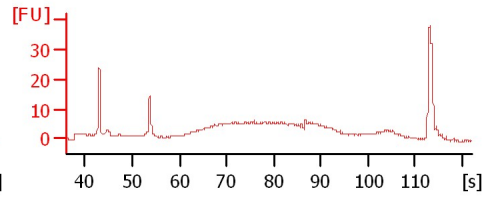
RA6_PB4



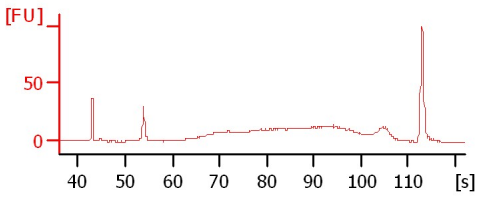
RA7_PB2



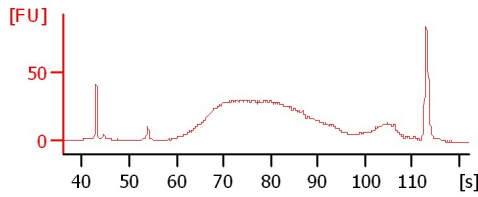
RA12_PB4



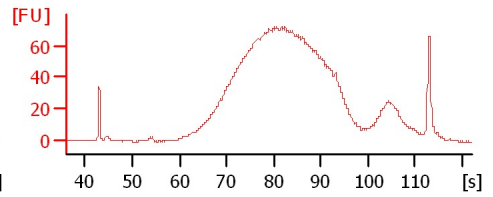
RA13_PB2



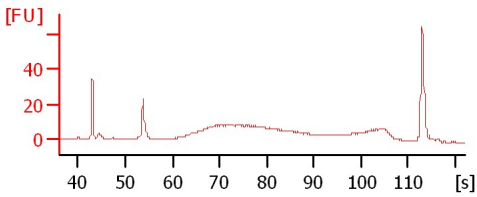
RA14_PB4



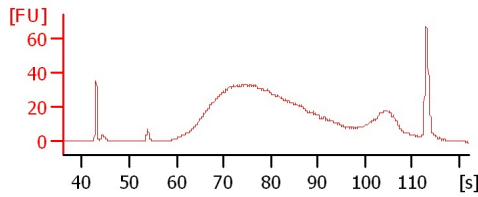
RA15_WT



RA16_DMSO2



RA18_DMSO4



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
RA2_PB2		<input type="checkbox"/>	✓			
RA4_PB4		<input type="checkbox"/>	✓			
RA5_PB2		<input type="checkbox"/>	✓			
RA6_PB4		<input type="checkbox"/>	✓			
RA7_PB2		<input type="checkbox"/>	✓			
RA12_PB4		<input type="checkbox"/>	✓			
RA13_PB2		<input type="checkbox"/>	✓			
RA14_PB4		<input type="checkbox"/>	✓			
RA15_WT		<input type="checkbox"/>	✓			
RA16_DMSO2		<input type="checkbox"/>	✓			
RA18_DMSO4		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

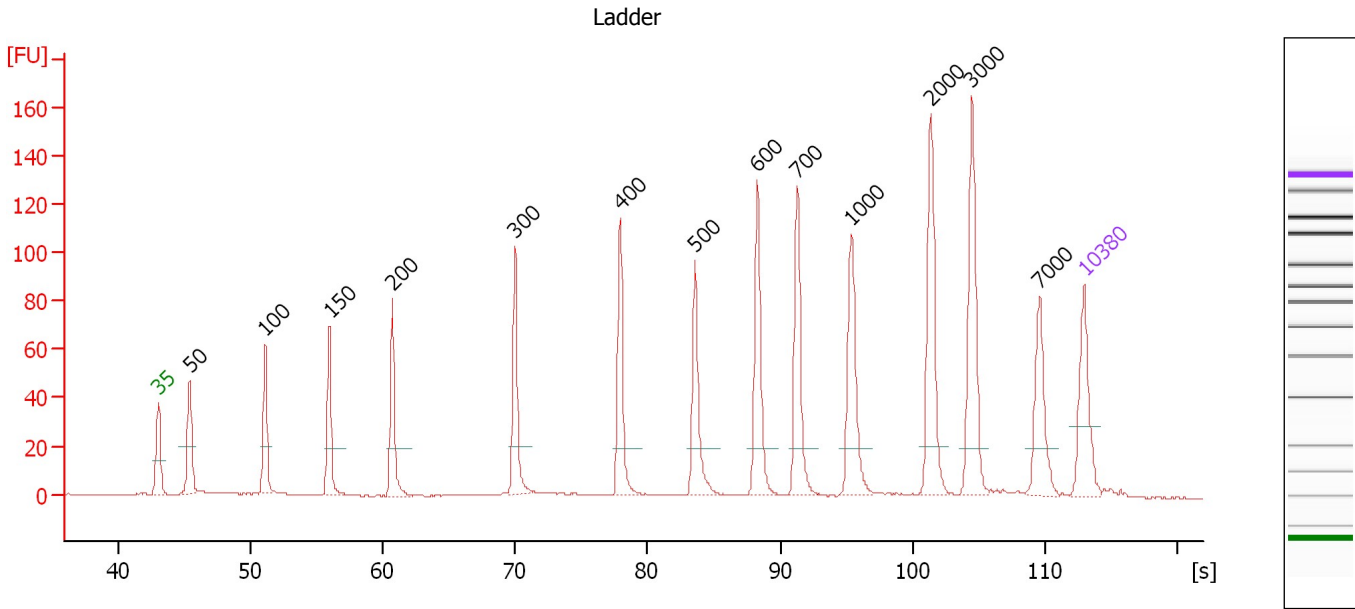
Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

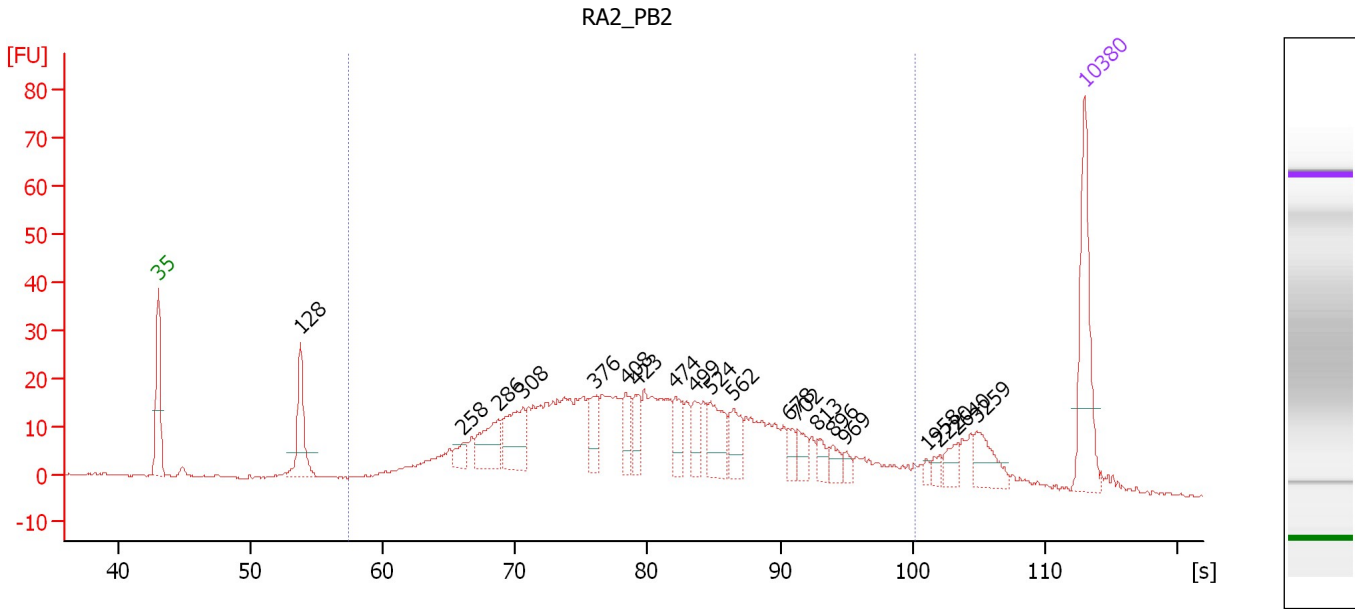
Peak table for Ladder

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	45.39
3	100	150.00	2,272.7	Ladder Peak	51.07
4	150	150.00	1,515.2	Ladder Peak	55.95
5	200	150.00	1,136.4	Ladder Peak	60.69
6	300	150.00	757.6	Ladder Peak	69.98
7	400	150.00	568.2	Ladder Peak	77.91
8	500	150.00	454.5	Ladder Peak	83.58
9	600	150.00	378.8	Ladder Peak	88.27
10	700	150.00	324.7	Ladder Peak	91.32
11	1,000	150.00	227.3	Ladder Peak	95.41
12	2,000	150.00	113.6	Ladder Peak	101.36
13	3,000	150.00	75.8	Ladder Peak	104.51
14	7,000	150.00	32.5	Ladder Peak	109.62
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : RA2_PB2

Number of peaks found: 20 Corr. Area 1: 646.4
 Noise: 0.2

Peak table for sample 1 : RA2_PB2

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	128	71.66	851.5		53.75
3	258	15.66	92.0		66.07
4	286	51.33	271.6		68.71
5	308	59.13	291.1		70.59
6	376	30.26	121.9		76.02
7	408	24.14	89.6		78.38
8	423	22.36	80.0		79.23
9	474	28.83	92.2		82.10
10	499	22.89	69.5		83.52
11	524	43.88	126.9		84.70
12	562	28.95	78.1		86.49
13	678	11.85	26.5		90.64
14	702	16.09	34.7		91.35
15	813	12.89	24.0		92.86
16	896	11.00	18.6		93.99
17	969	6.88	10.8		94.98
18	1,958	3.95	3.1		101.11
19	2,220	6.35	4.3		102.06
20	2,640	11.66	6.7		103.38
21	3,259	25.77	12.0		104.84
22	10,380	75.00	10.9	Upper Marker	113.00


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...

... Region table for sample 1 :

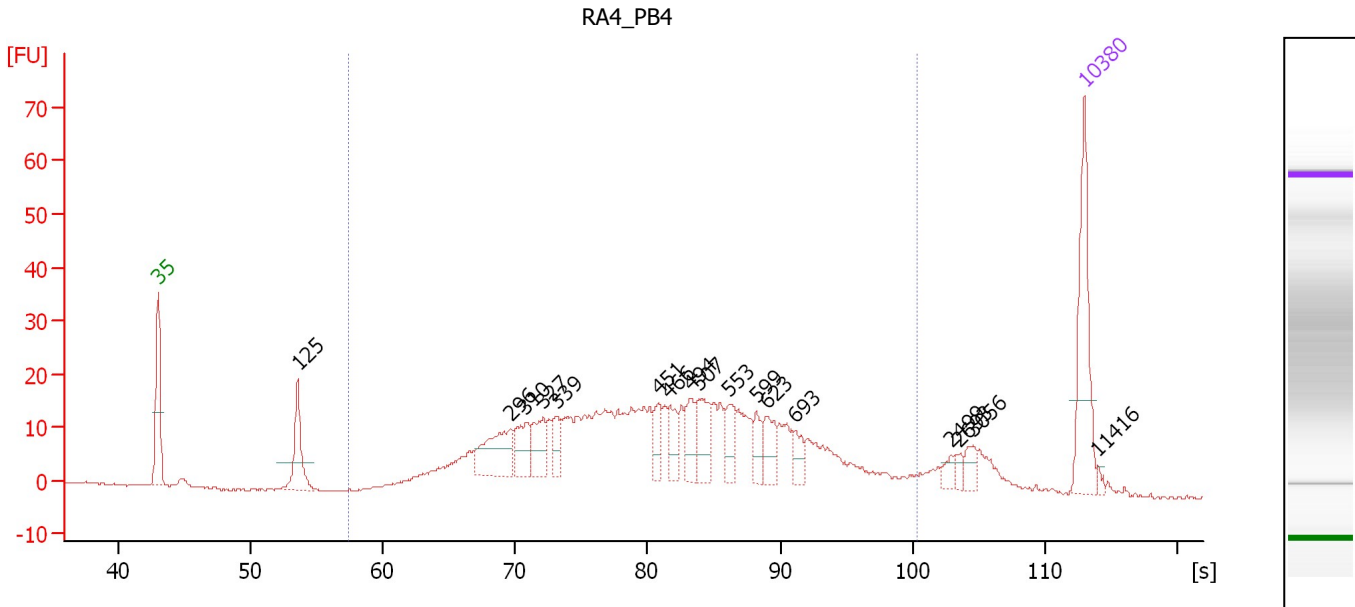
RA2 PB2

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/ μ l]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
57.44	100.20	505	646.4	1,124.51	4,351.7	 84	53.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : RA4 PB4

Number of peaks found: 17 Corr. Area 1: 514.2
 Noise: 0.2

Peak table for sample 2 : RA4 PB4

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	125	65.90	796.9		53.54
3	296	68.14	348.5		69.63
4	310	36.75	179.7		70.76
5	327	37.03	171.6		72.12
6	339	20.22	90.2		73.11
7	451	21.25	71.5		80.78
8	466	26.22	85.2		81.67
9	494	28.81	88.4		83.22
10	507	38.85	116.0		83.93
11	553	26.55	72.7		86.09
12	599	16.82	42.6		88.21
13	623	27.81	67.7		88.96
14	693	18.17	39.7		91.13
15	2,499	7.70	4.7		102.93
16	2,693	5.77	3.2		103.54
17	3,056	11.37	5.6		104.58
18	10,380	75.00	10.9	Upper Marker	113.00
19	11,416	0.00	0.0		114.03

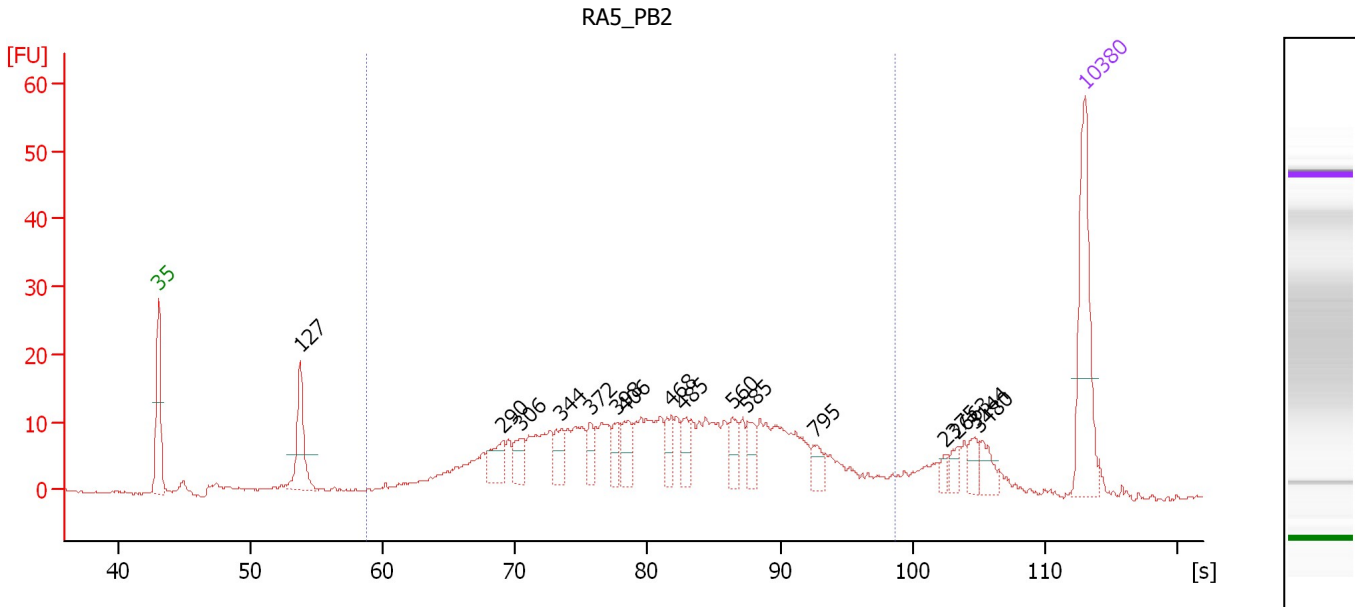
Region table for sample 2 : RA4 PB4

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
57.43	100.28	512	514.2	1,010.93	3,722.0	88	48.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : RA5_PB2

Number of peaks found: 16 Corr. Area 1: 371.0
 Noise: 0.2

Peak table for sample 3 : RA5_PB2

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	127	63.65	756.9		53.74
3	290	28.25	147.5		69.07
4	306	20.61	102.1		70.46
5	344	20.86	91.9		73.46
6	372	16.48	67.2		75.66
7	398	16.10	61.4		77.71
8	406	22.94	85.6		78.24
9	468	18.37	59.5		81.77
10	485	17.86	55.8		82.73
11	560	18.40	49.8		86.40
12	585	16.02	41.5		87.55
13	795	12.81	24.4		92.61
14	2,375	5.26	3.4		102.54
15	2,663	8.19	4.7		103.45
16	3,144	10.47	5.0		104.69
17	3,480	13.33	5.8		105.12
18	10,380	75.00	10.9	Upper Marker	113.00

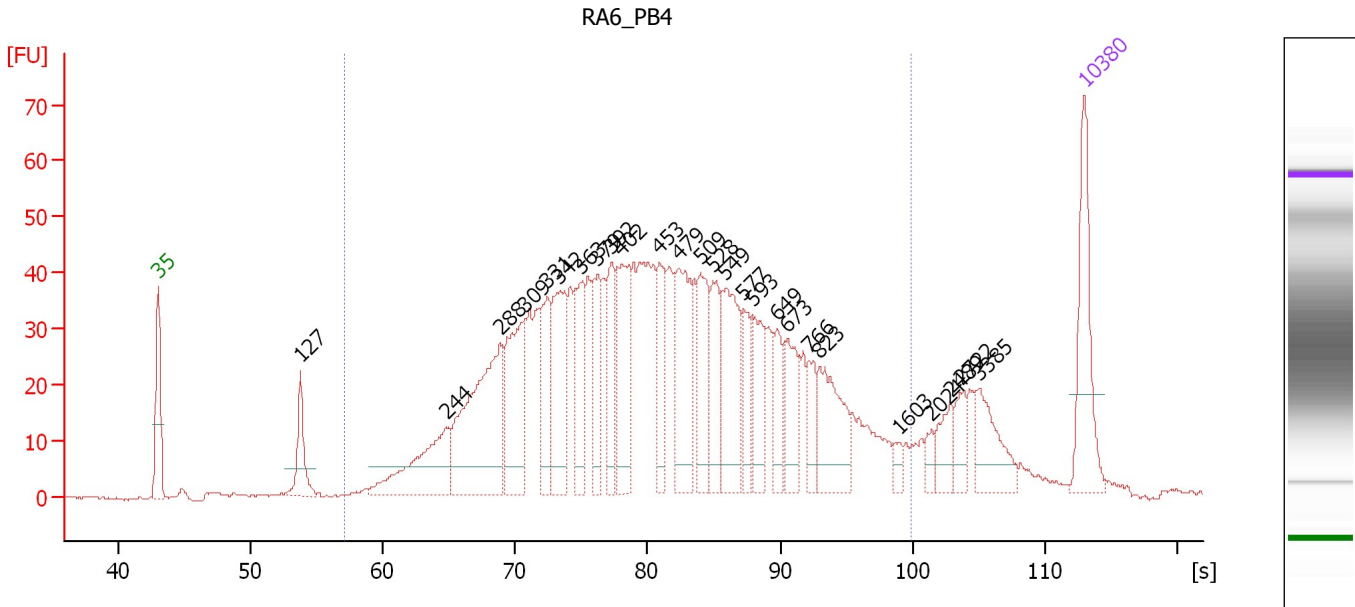
Region table for sample 3 : RA5_PB2

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
58.68	98.71	500	371.0	804.32	3,004.4	81	43.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : RA6_PB4

Number of peaks found: 26 Corr. Area 1: 1,381.1
 Noise: 0.2

Peak table for sample 4 : RA6_PB4

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	127	59.41	707.2		53.73
3	244	116.04	721.2		64.76
4	288	224.26	1,180.2		68.85
5	309	125.19	614.7		70.66
6	331	74.60	341.7		72.41
7	342	106.99	473.4		73.34
8	363	71.03	296.7		74.95
9	379	66.36	265.5		76.22
10	392	62.79	242.9		77.24
11	402	93.49	352.3		78.02
12	453	59.79	200.1		80.90
13	479	116.99	370.4		82.37
14	509	72.86	216.7		84.02
15	528	64.90	186.2		84.90
16	549	102.47	282.8		85.88
17	577	40.88	107.3		87.20
18	593	57.06	145.9		87.93
19	649	37.21	86.8		89.78
20	673	55.35	124.5		90.51
21	766	29.45	58.3		92.22
22	823	76.99	141.7		93.00
23	1,603	9.43	8.9		99.00
24	2,024	10.43	7.8		101.44
25	2,489	22.41	13.6		102.90

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : RA6_PB4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,722	20.82	11.6		103.63
27	3,385	44.94	20.1		105.00
28	10,380	75.00	10.9	Upper Marker	113.00

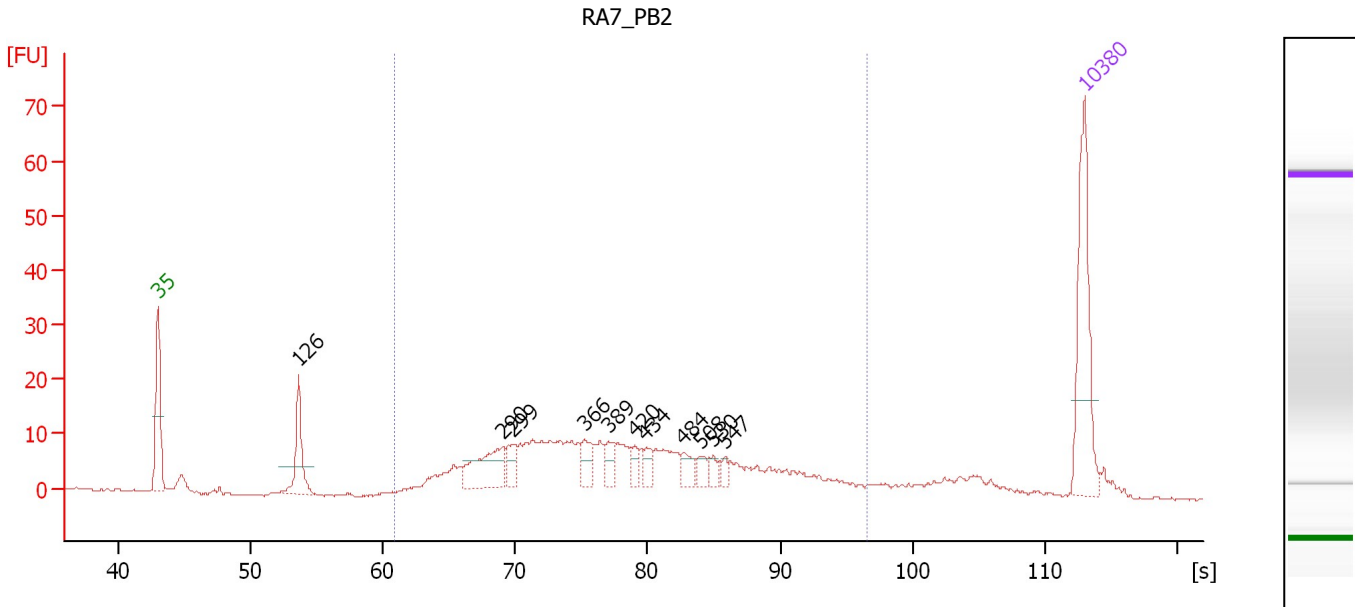
Region table for sample 4 : RA6_PB4

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
57.02	99.91	504	1,381.1	2,398.28	9,135.6	89	50.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : RA7_PB2

Number of peaks found: 11 Corr. Area 1: 268.3
 Noise: 0.2

Peak table for sample 5 : RA7_PB2

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	126	56.28	675.6		53.63
3	290	52.17	272.5		69.05
4	299	16.52	83.8		69.84
5	366	18.08	74.9		75.18
6	389	13.05	50.9		77.01
7	420	10.42	37.6		79.04
8	434	11.32	39.5		79.83
9	484	11.85	37.1		82.70
10	508	10.35	30.8		83.98
11	530	6.55	18.7		84.97
12	547	6.13	17.0		85.81
13	10,380	75.00	10.9	Upper Marker	113.00

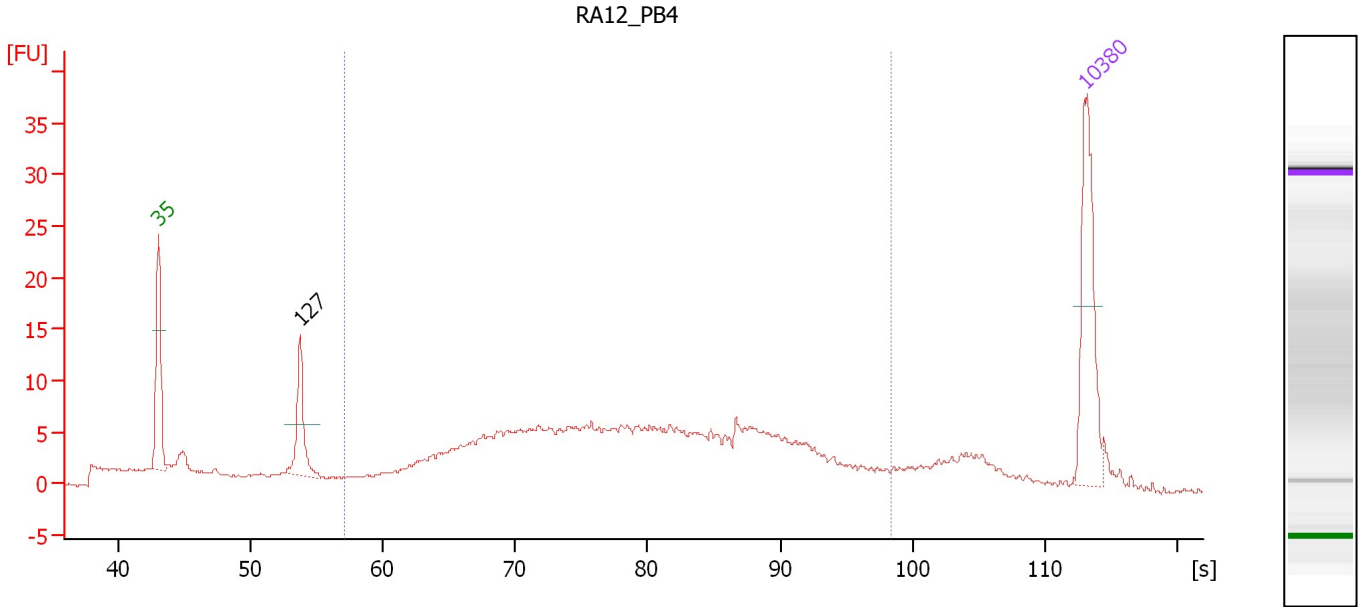
Region table for sample 5 : RA7_PB2

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
60.92	96.54	431	268.3	484.09	1,975.4	84	36.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : RA12_PB4

Number of peaks found: 1 Corr. Area 1: 216.7
 Noise: 0.2

Peak table for sample 6 : RA12_PB4

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	127	65.12	778.4		53.68
3	10,380	75.00	10.9	Upper Marker	113.00

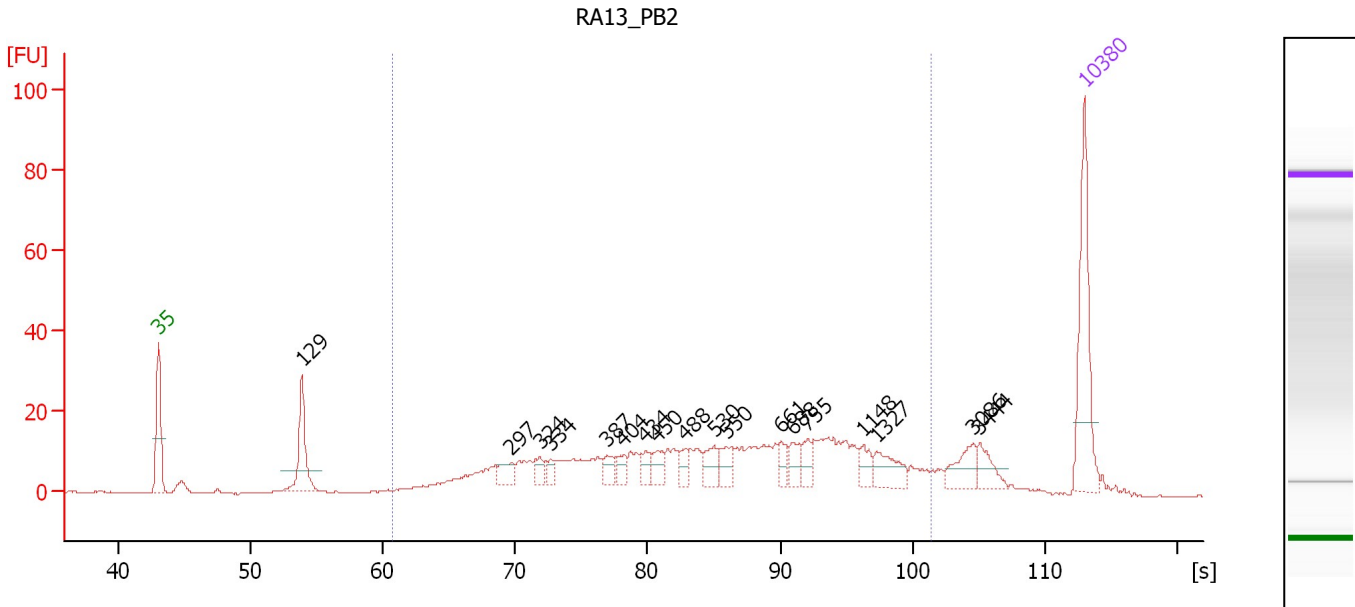
Region table for sample 6 : RA12_PB4

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
57.03	98.39	466	216.7	659.95	2,743.9	71	46.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : RA13 PB2

Number of peaks found: 18 Corr. Area 1: 432.7
 Noise: 0.2

Peak table for sample 7 : RA13 PB2

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	129	67.98	800.3		53.87
3	297	16.07	82.0		69.67
4	324	11.86	55.5		71.85
5	334	9.16	41.5		72.67
6	387	13.23	51.8		76.88
7	404	11.69	43.8		78.15
8	434	10.44	36.5		79.83
9	450	16.29	54.8		80.74
10	488	13.54	42.1		82.88
11	530	18.96	54.2		85.01
12	550	15.67	43.2		85.92
13	661	11.38	26.1		90.14
14	688	15.62	34.4		90.95
15	755	14.98	30.1		92.07
16	1,148	12.44	16.4		96.29
17	1,327	23.30	26.6		97.35
18	3,086	20.82	10.2		104.62
19	3,444	15.86	7.0		105.08
20	10,380	75.00	10.9	Upper Marker	113.00

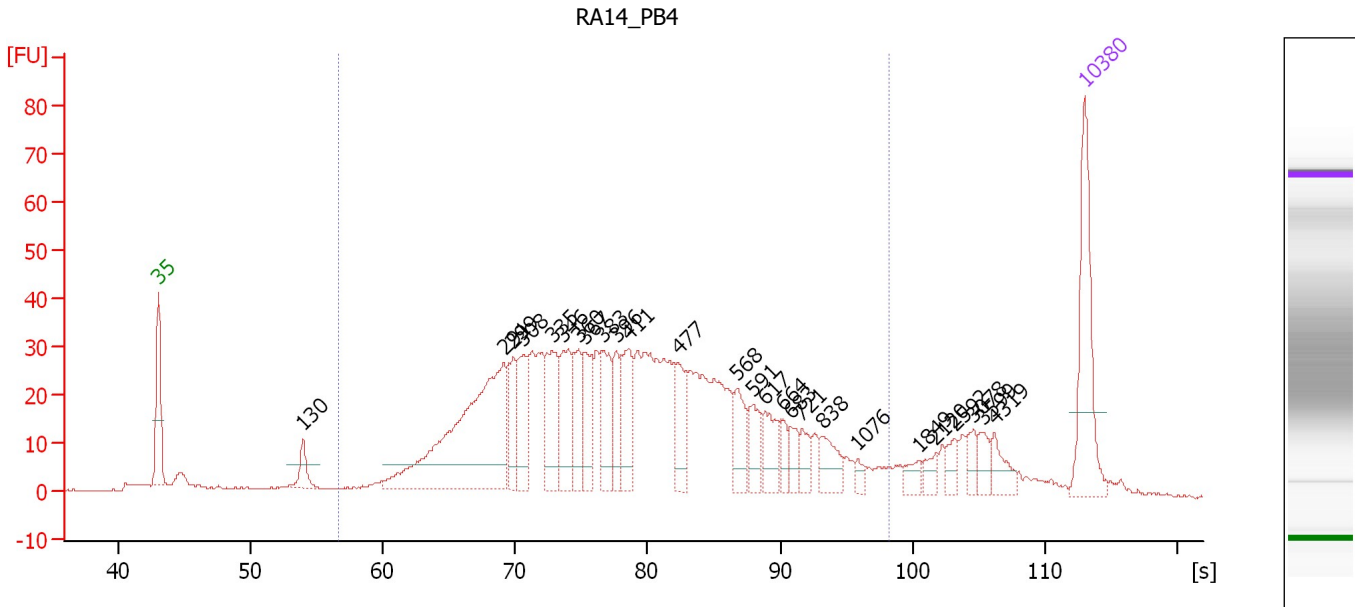
Region table for sample 7 : RA13 PB2

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
60.69	101.39	654	432.7	641.26	2,119.8	82	58.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : RA14 PB4

Number of peaks found: 26 Corr. Area 1: 970.9
 Noise: 0.2

Peak table for sample 8 : RA14 PB4

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	24.30	284.2		53.95
3	291	282.38	1,469.9		69.15
4	299	43.00	218.1		69.86
5	308	50.27	247.6		70.58
6	335	67.75	306.1		72.78
7	346	60.01	262.5		73.65
8	360	43.08	181.4		74.73
9	367	41.88	172.9		75.29
10	383	48.95	193.6		76.57
11	396	36.34	139.0		77.59
12	411	50.05	184.7		78.51
13	477	43.09	136.8		82.30
14	568	33.85	90.4		86.75
15	591	26.22	67.2		87.88
16	617	29.72	73.0		88.80
17	664	15.90	36.3		90.23
18	683	13.68	30.4		90.79
19	721	15.59	32.8		91.61
20	838	25.88	46.8		93.20
21	1,076	6.62	9.3		95.86
22	1,849	9.81	8.0		100.46
23	2,120	8.75	6.3		101.74
24	2,592	10.74	6.3		103.23
25	3,078	11.76	5.8		104.61

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...

... Peak table for sample 8 : RA14 PB4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	3,599	12.03	5.1		105.27
27	4,319	17.18	6.0		106.19
28	10,380	75.00	10.9	Upper Marker	113.00

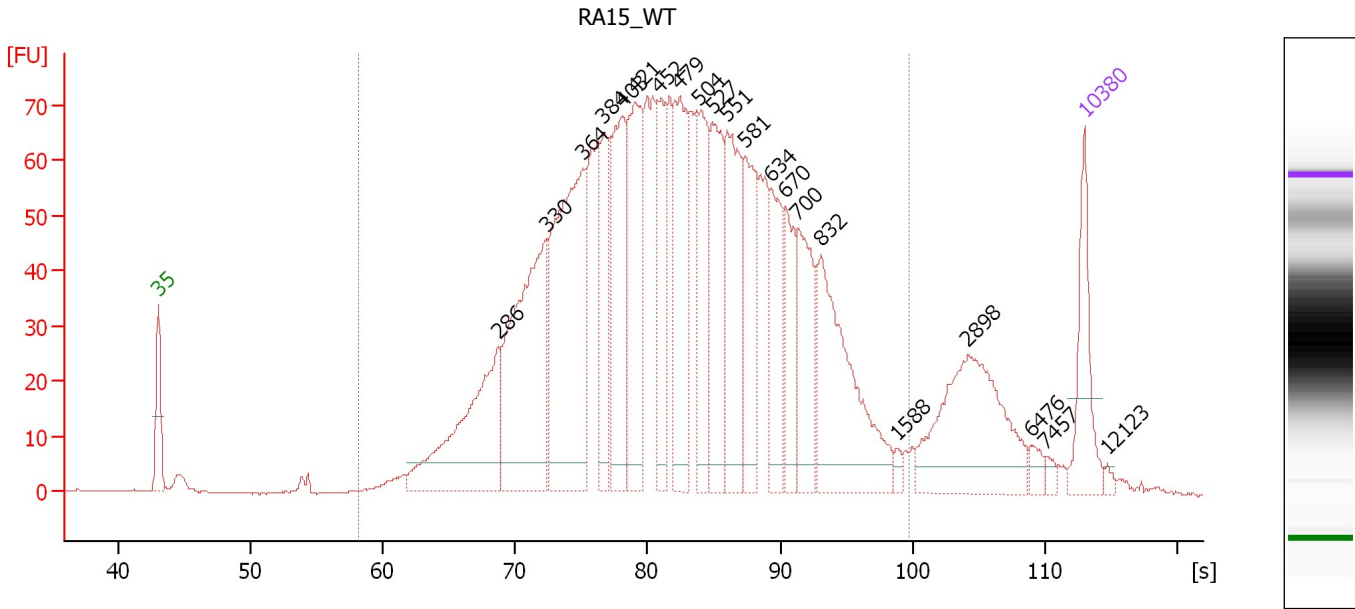
Region table for sample 8 : RA14 PB4

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
56.70	98.21	449	970.9	1,480.41	6,074.8	86	43.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : RA15_WT

Number of peaks found: 21 Corr. Area 1: 2,025.8
 Noise: 0.2


Peak table for sample 9 : RA15_WT

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	269.15	1,426.1		68.67
3	330	365.12	1,676.0		72.36
4	364	398.95	1,659.1		75.08
5	384	107.44	423.5		76.67
6	403	197.89	743.1		78.10
7	421	203.46	732.9		79.08
8	452	109.95	368.4		80.87
9	479	181.02	572.3		82.41
10	504	139.85	420.1		83.79
11	527	159.87	459.3		84.87
12	551	181.72	499.3		85.99
13	581	133.35	347.8		87.38
14	634	101.30	241.9		89.33
15	670	87.99	199.1		90.40
16	700	126.49	273.8		91.32
17	832	236.30	430.4		93.12
18	1,588	7.69	7.3		98.91
19	2,898	190.25	99.5		104.19
20	6,476	11.98	2.8		108.95
21	7,457	7.78	1.6		110.08
22	10,380	75.00	10.9	Upper Marker	113.00
23	12,123	0.00	0.0		114.74

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
Modified: 1/5/2018 1:21:00 PM

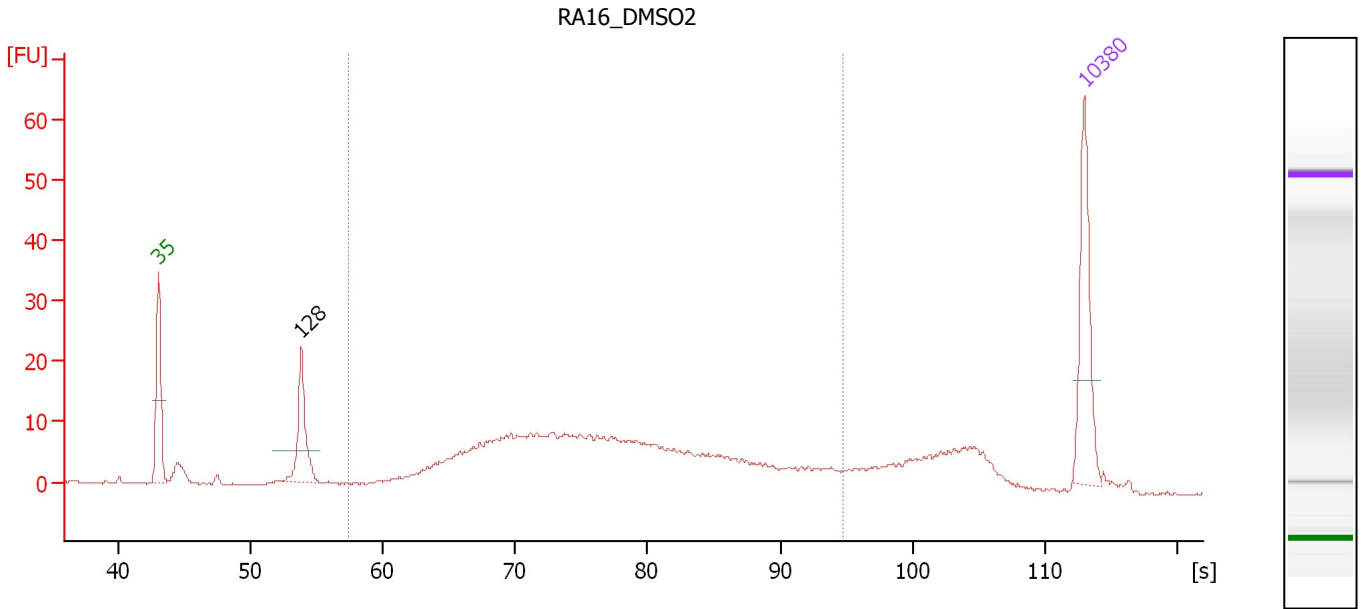
Electropherogram Summary Continued ...**... Region table for sample 9 : RA15 WT**

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
58.11	99.68	513	2,025.8	3,790.68	13,431.1	 91	41.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrilib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : RA16_DMSO2

Number of peaks found: 1 Corr. Area 1: 278.5
 Noise: 0.2

Peak table for sample 10 : RA16_DMSO2

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	128	82.17	973.9		53.79
3	10,380	75.00	10.9	Upper Marker	113.00

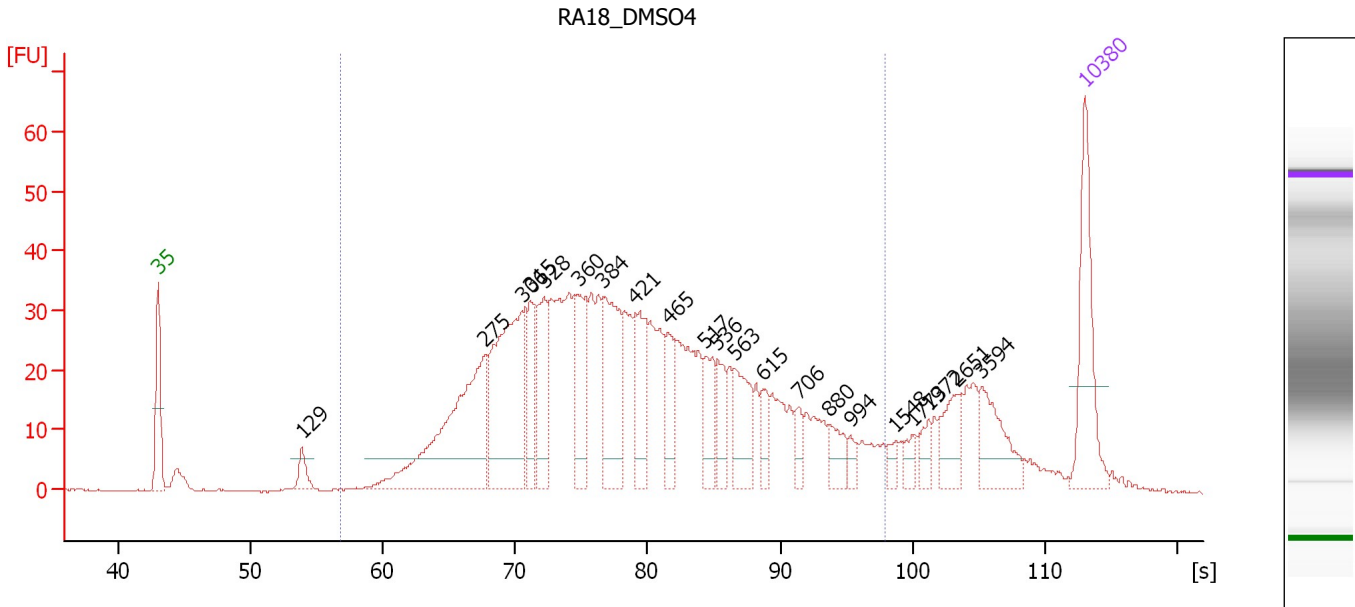
Region table for sample 10 : RA16_DMSO2

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
57.34	94.82	420	278.5	638.91	2,735.8	68	37.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrplib.xad

Created: 1/5/2018 12:36:04 PM
 Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : RA18 DMSO4

Number of peaks found: 21 Corr. Area 1: 995.3
 Noise: 0.1

Peak table for sample 11 : RA18 DMSO4


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	129	19.75	232.4		53.88
3	275	235.60	1,296.2		67.69
4	306	206.09	1,019.6		70.47
5	315	60.67	292.1		71.14
6	328	68.56	317.0		72.18
7	360	69.40	291.9		74.75
8	384	103.92	409.8		76.66
9	421	62.75	226.0		79.08
10	465	36.88	120.1		81.61
11	517	41.16	120.6		84.39
12	536	31.96	90.3		85.27
13	563	50.99	137.1		86.56
14	615	19.88	49.0		88.72
15	706	15.96	34.3		91.40
16	880	22.14	38.1		93.77
17	994	8.63	13.2		95.32
18	1,548	7.13	7.0		98.67
19	1,773	10.46	8.9		100.01
20	1,972	13.17	10.1		101.20
21	2,651	31.50	18.0		103.41
22	3,594	44.58	18.8		105.27
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdriplib.xad

Created: 1/5/2018 12:36:04 PM
Modified: 1/5/2018 1:21:00 PM

Electropherogram Summary Continued ...

... Region table for sample 11 : RA18_DMSO4

From [s]	To [s]	Average Size [bp]	Corr. Area	Conc. [pg/ μ l]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
56.77	97.89	453	995.3	1,802.62	7,327.3	 85	44.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... expert\data\2018-01-05\2018-01-05_001_chedin_pbdrplib.xad

Created: 1/5/2018 12:36:04 PM
Modified: 1/5/2018 1:21:00 PM

Gel Image

