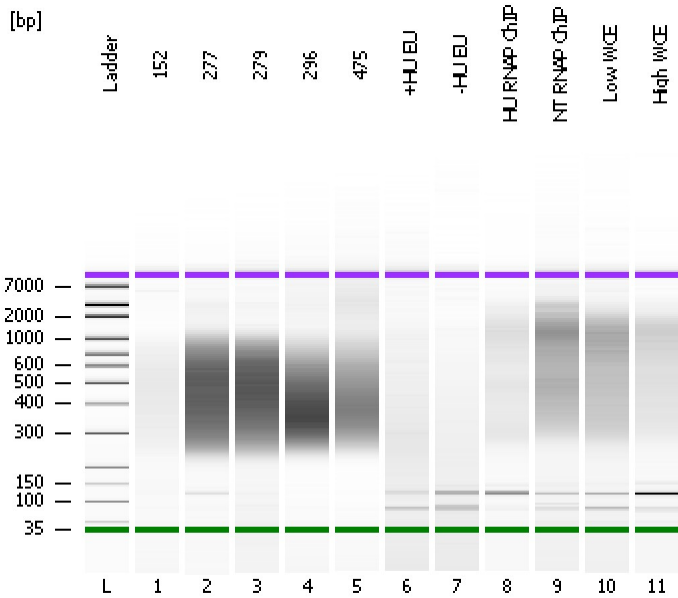


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 PM

**Electrophoresis File Run Summary**



Instrument Information:

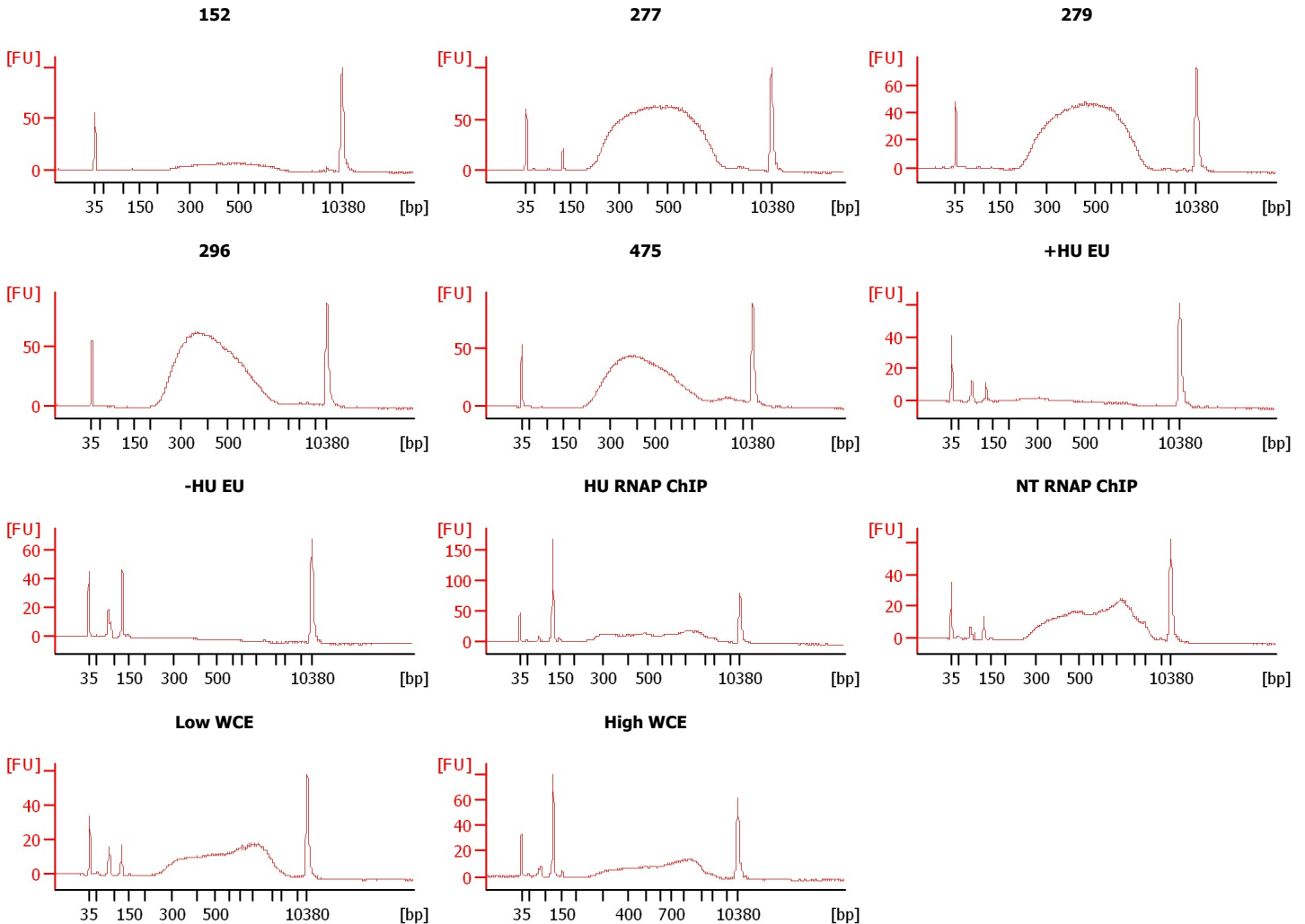
Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086      Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
152		<input type="checkbox"/>	✓			
277		<input type="checkbox"/>	✓			
279		<input type="checkbox"/>	✓			
296		<input type="checkbox"/>	✓			
475		<input type="checkbox"/>	✓			
+HU EU		<input type="checkbox"/>	✓			
-HU EU		<input type="checkbox"/>	✓			
HU RNAP ChIP		<input type="checkbox"/>	✓			
NT RNAP ChIP		<input type="checkbox"/>	✓			
Low WCE		<input type="checkbox"/>	✓			
High WCE		<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\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 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

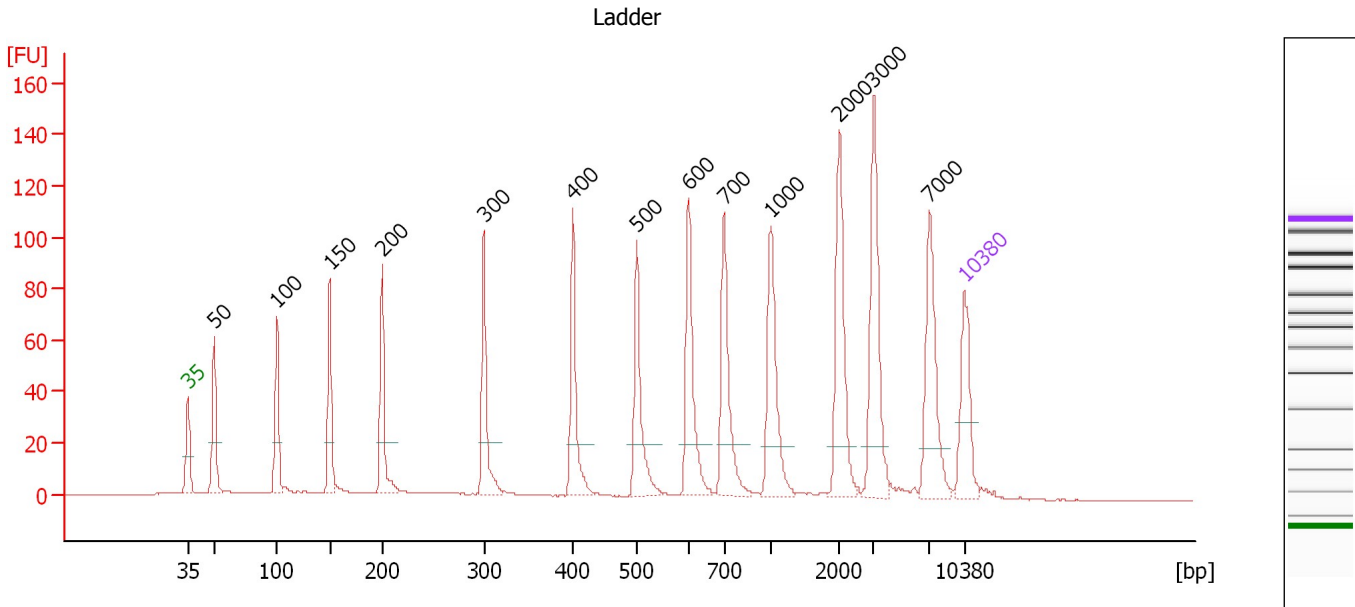
**Ladder**

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\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

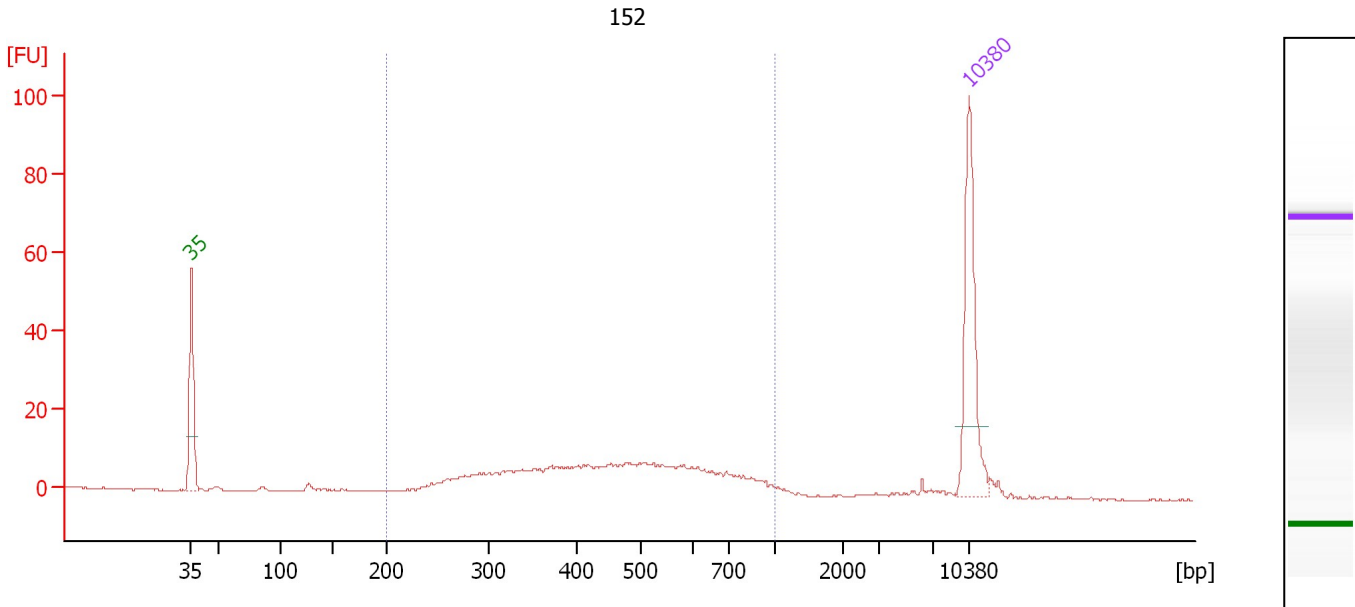
**Peak table for Ladder**

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	45.36
3	100	150.00	2,272.7	Ladder Peak	50.98
4	150	150.00	1,515.2	Ladder Peak	55.79
5	200	150.00	1,136.4	Ladder Peak	60.50
6	300	150.00	757.6	Ladder Peak	69.68
7	400	150.00	568.2	Ladder Peak	77.71
8	500	150.00	454.5	Ladder Peak	83.48
9	600	150.00	378.8	Ladder Peak	88.14
10	700	150.00	324.7	Ladder Peak	91.32
11	1,000	150.00	227.3	Ladder Peak	95.55
12	2,000	150.00	113.6	Ladder Peak	101.70
13	3,000	150.00	75.8	Ladder Peak	104.83
14	7,000	150.00	32.5	Ladder Peak	109.83
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 152**

Number of peaks found: 0                      Corr. Area 1: 217.2  
 Noise: 0.2

**Peak table for sample 1 : 152**

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

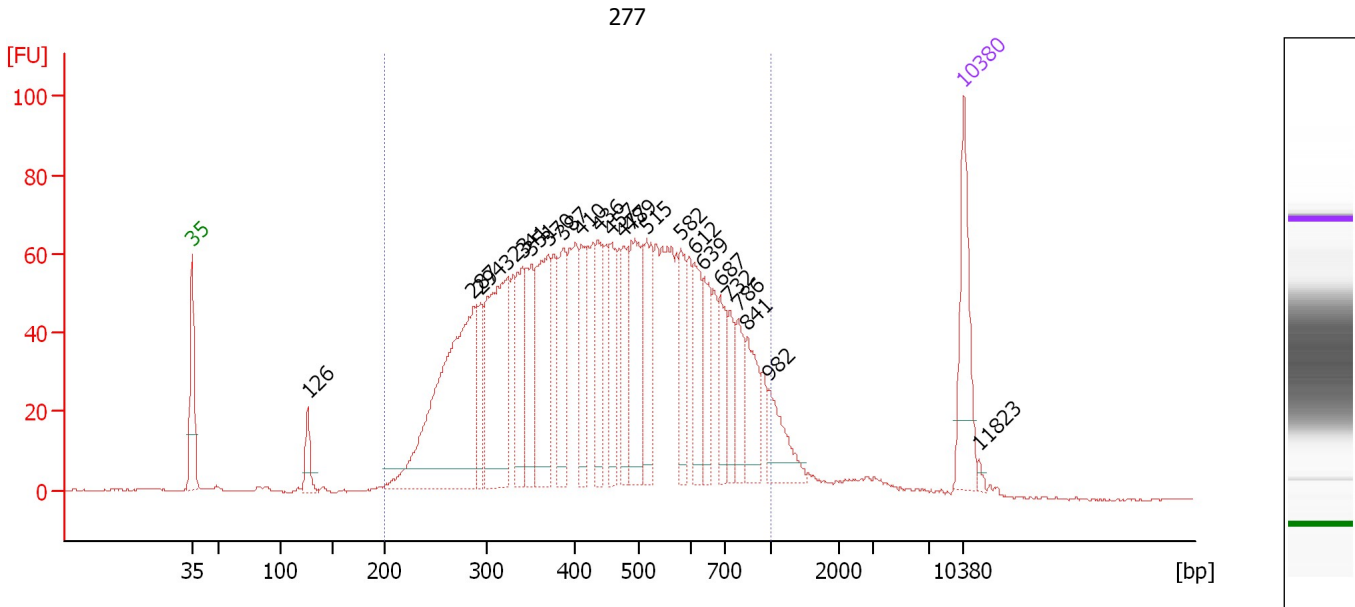
**Region table for sample 1 : 152**

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	481	217.2	908.8	248.22	93	33.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 277**

Number of peaks found: 23                      Corr. Area 1: 2,128.2  
 Noise: 0.2

**Peak table for sample 2 : 277**

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	34.54	416.4		53.45
3	287	375.84	1,983.8		68.49
4	294	59.92	308.9		69.12
5	323	224.01	1,052.3		71.49
6	341	82.57	366.6		72.99
7	351	82.14	354.7		73.77
8	370	142.64	584.8		75.27
9	387	85.49	334.7		76.67
10	410	60.81	224.5		78.31
11	436	60.69	210.7		79.81
12	457	63.40	210.0		81.02
13	477	70.65	224.6		82.14
14	489	110.81	343.1		82.86
15	515	94.21	277.3		84.17
16	582	55.55	144.6		87.31
17	612	57.58	142.6		88.52
18	639	54.00	128.0		89.39
19	687	51.05	112.6		90.89
20	732	36.43	75.4		91.76
21	786	42.26	81.4		92.54
22	841	54.60	98.3		93.31
23	982	47.34	73.0		95.29
24	10,380	75.00	10.9	Upper Marker	113.00
25	11,823	0.00	0.0		114.35

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**

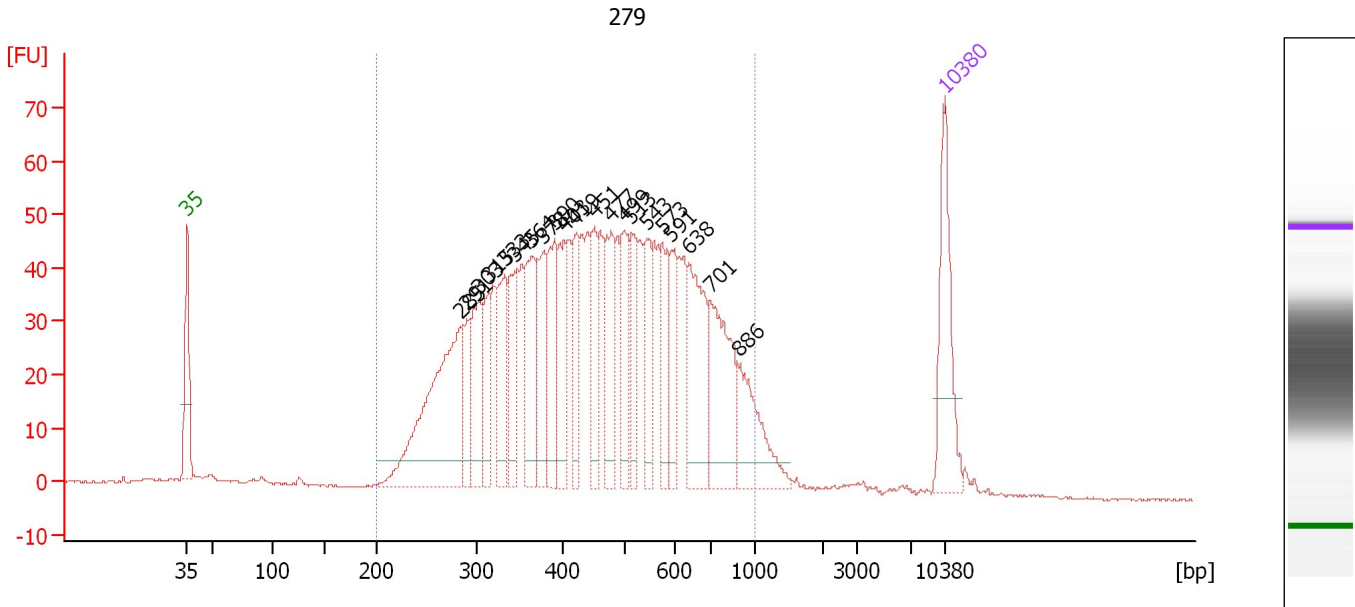
... Region table for sample 2 : 277

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	472	2,128.2	10,252.6	2,698.25	94	35.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 279**

Number of peaks found: 21                      Corr. Area 1: 1,525.6  
 Noise: 0.2

**Peak table for sample 3 : 279**

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	258.07	1,373.8		68.27
3	291	45.49	236.8		68.86
4	303	81.39	406.6		69.95
5	315	49.35	237.4		70.88
6	333	83.02	378.1		72.31
7	345	71.67	314.8		73.30
8	364	85.28	355.0		74.82
9	378	68.41	274.6		75.91
10	390	81.01	314.4		76.94
11	403	68.15	256.3		77.88
12	419	57.17	206.7		78.81
13	451	56.88	191.2		80.64
14	477	78.24	248.5		82.16
15	499	62.22	188.8		83.44
16	513	49.13	145.1		84.08
17	543	53.05	148.2		85.46
18	573	58.87	155.6		86.89
19	591	47.30	121.2		87.73
20	638	103.79	246.4		89.35
21	701	114.28	247.2		91.33
22	886	72.98	124.8		93.94
23	10,380	75.00	10.9	Upper Marker	113.00




Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**

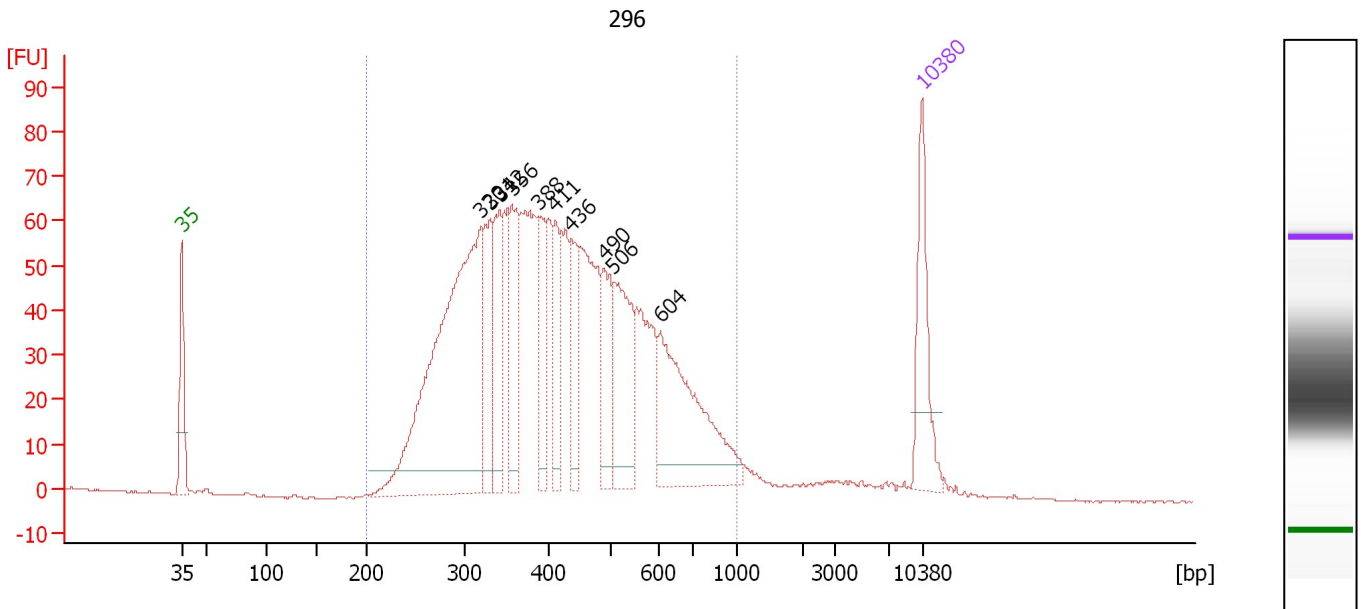
... Region table for sample 3 : 279

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
200	1,000	476	1,525.6	8,061.0	 2,157.17	95	34.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 296**

Number of peaks found: 10                      Corr. Area 1: 1,734.9  
 Noise: 0.2

**Peak table for sample 4 : 296**

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	320	559.71	2,651.8		71.27
3	331	102.96	471.2		72.18
4	342	97.52	431.5		73.09
5	356	113.14	481.2		74.20
6	388	81.63	319.1		76.71
7	411	71.41	263.4		78.33
8	436	65.71	228.3		79.79
9	490	80.09	247.6		82.91
10	506	133.72	400.2		83.77
11	604	196.07	492.3		88.26
12	10,380	75.00	10.9	Upper Marker	113.00

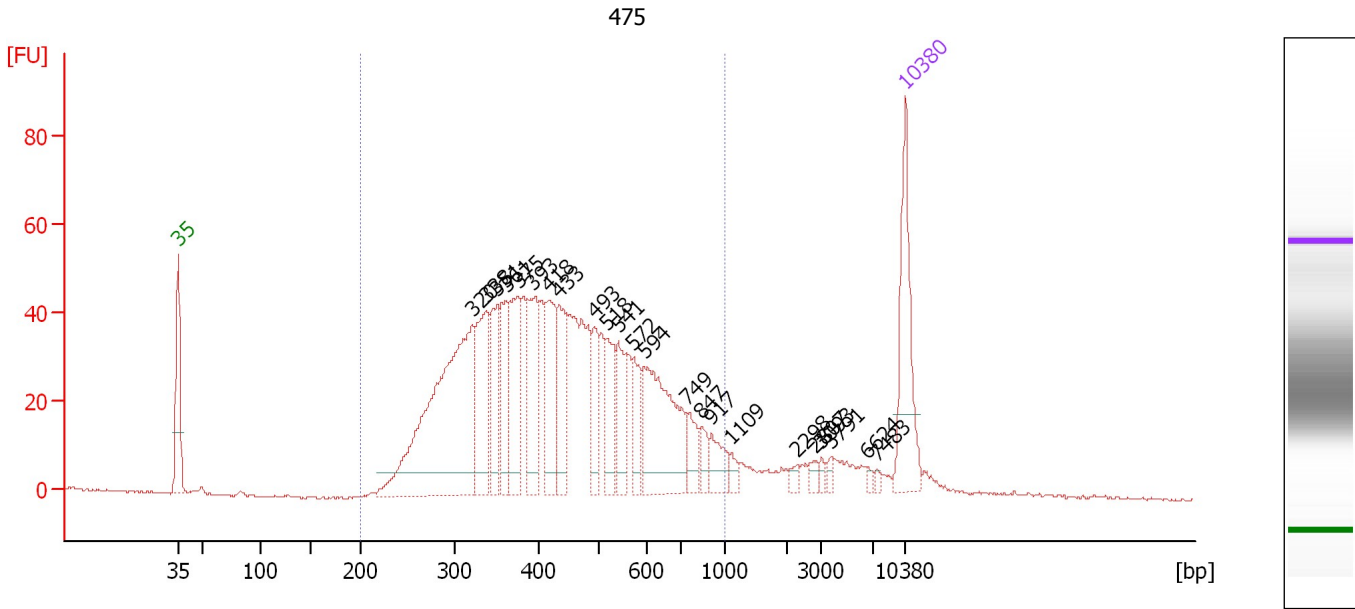
**Region table for sample 4 : 296**

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	435	1,734.9	8,921.3	2,258.73	96	32.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 475**

Number of peaks found: 23                      Corr. Area 1: 1,208.4  
 Noise: 0.2

**Peak table for sample 5 : 475**

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	320	347.47	1,645.9		71.28
3	338	98.64	442.5		72.71
4	351	65.73	284.1		73.74
5	361	66.02	276.8		74.61
6	375	76.75	309.7		75.74
7	393	81.05	312.2		77.18
8	418	77.18	279.5		78.77
9	433	66.91	233.9		79.64
10	493	42.43	130.4		83.08
11	518	48.52	142.0		84.31
12	541	42.24	118.3		85.39
13	572	34.37	91.1		86.83
14	594	130.30	332.5		87.85
15	749	25.69	52.0		92.01
16	847	14.65	26.2		93.40
17	917	24.00	39.7		94.37
18	1,109	7.00	9.6		96.22
19	2,298	5.38	3.5		102.63
20	2,807	5.64	3.0		104.22
21	3,093	4.19	2.1		104.94
22	3,791	4.71	1.9		105.82
23	6,624	3.19	0.7		109.36
24	7,483	2.64	0.5		110.28
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**

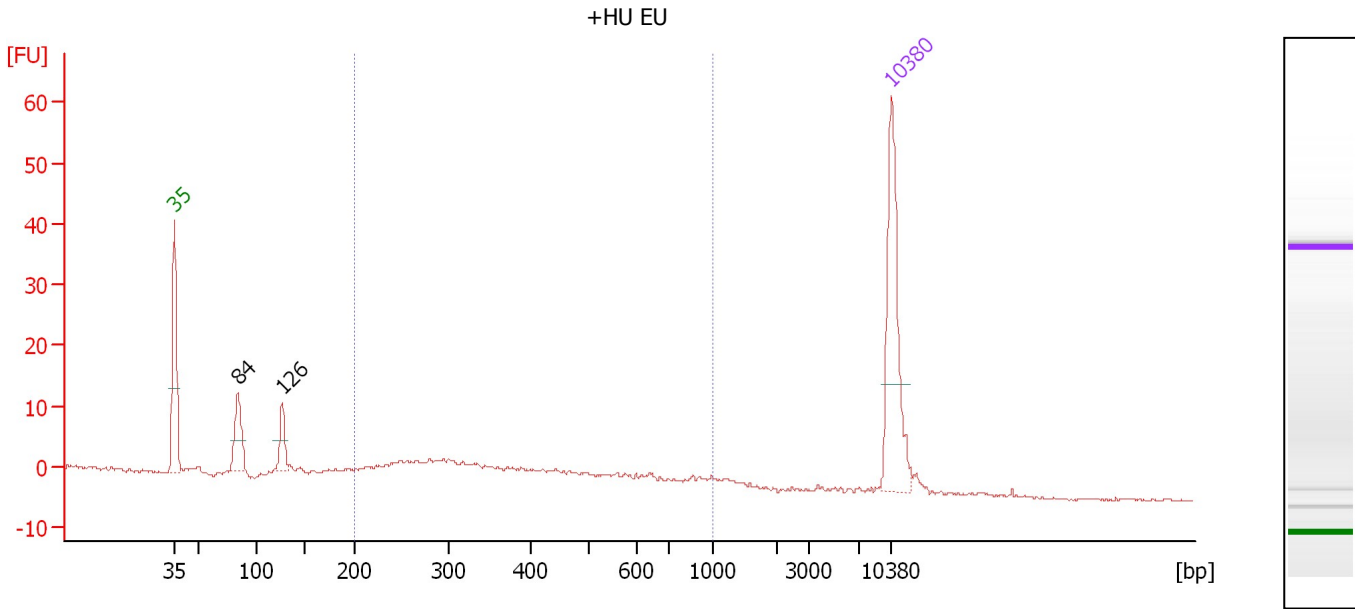
... Region table for sample 5 : 475

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	454	1,208.4	5,906.5	■ 1,550.26	90	32.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : +HU EU**

Number of peaks found: 2                      Corr. Area 1: 76.6  
 Noise: 0.1

**Peak table for sample 6 : +HU EU**

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	84	43.98	792.6		49.19
3	126	23.45	281.5		53.51
4	10,380	75.00	10.9	Upper Marker	113.00

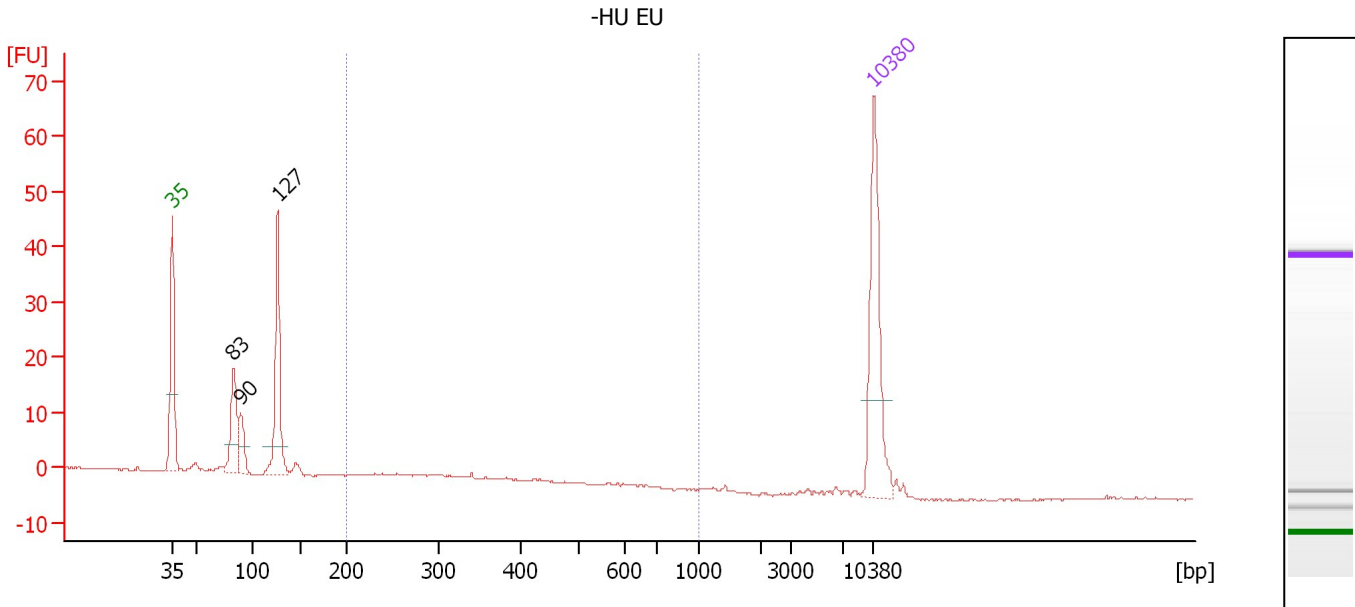
**Region table for sample 6 : +HU EU**

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	408	76.6	568.6	125.90	63	42.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : -HU EU**

Number of peaks found: 3                      Corr. Area 1: 9.6  
 Noise: 0.2

**Peak table for sample 7 : -HU EU**

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	83	58.70	1,066.0		49.12
3	90	26.69	449.1		49.86
4	127	100.68	1,205.6		53.53
5	10,380	75.00	10.9	Upper Marker	113.00

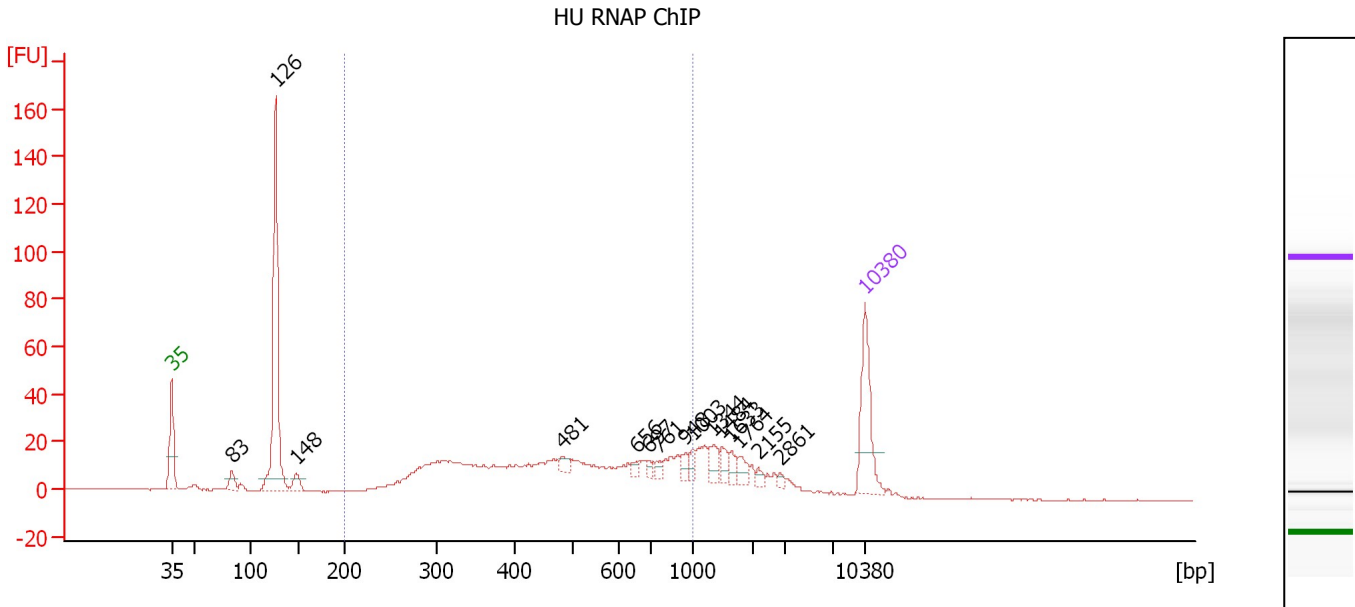
**Region table for sample 7 : -HU EU**

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	296	9.6	87.9	16.51	9	18.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : HU RNAP ChIP**

Number of peaks found: 15      Corr. Area 1: 467.5  
 Noise: 0.2

**Peak table for sample 8 : HU RNAP ChIP**

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	83	20.66	376.7		49.08
3	126	331.64	3,984.3		53.49
4	148	20.19	206.8		55.59
5	481	11.04	34.8		82.39
6	656	7.62	17.6		89.92
7	697	7.25	15.8		91.21
8	761	8.06	16.0		92.18
9	948	12.10	19.3		94.81
10	1,003	10.98	16.6		95.57
11	1,344	19.40	21.9		97.67
12	1,484	12.91	13.2		98.53
13	1,633	12.56	11.7		99.44
14	1,764	13.11	11.3		100.25
15	2,155	6.18	4.3		102.19
16	2,861	4.17	2.2		104.39
17	10,380	75.00	10.9	Upper Marker	113.00

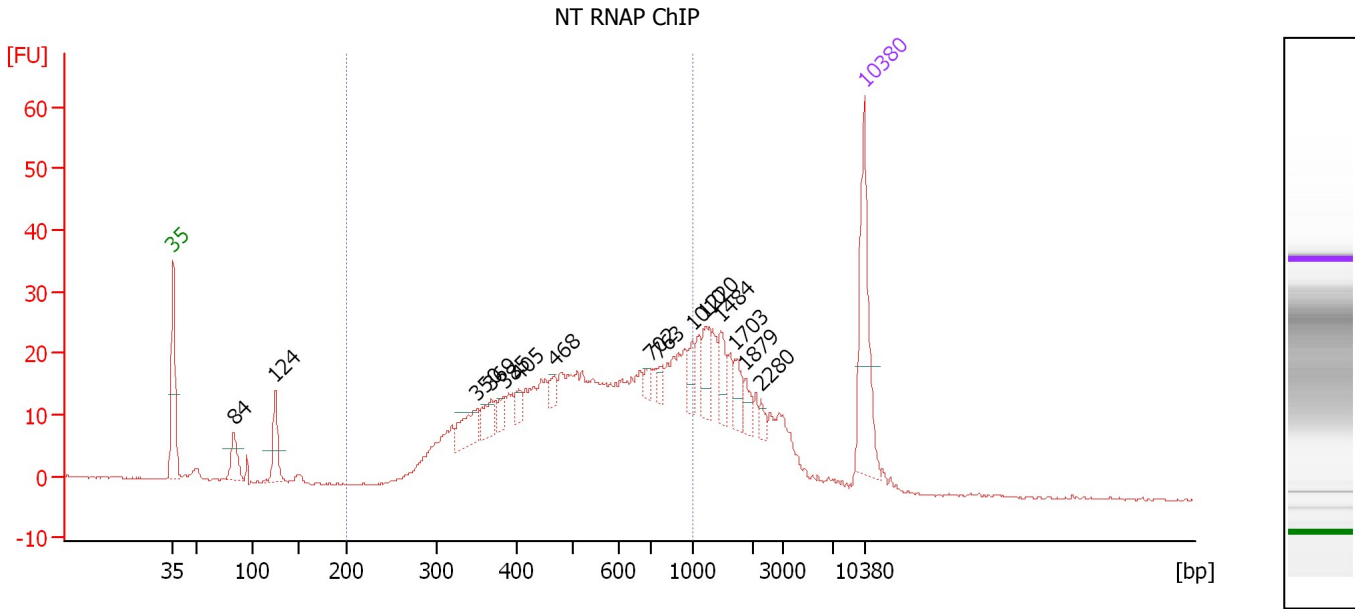
**Region table for sample 8 : HU RNAP ChIP**

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	512	467.5	2,444.3	677.62	56	38.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : NT RNAP ChIP**

Number of peaks found: 15                      Corr. Area 1: 500.9  
 Noise: 0.2

**Peak table for sample 9 : NT RNAP ChIP**

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	84	34.65	626.6		49.16
3	124	41.45	505.3		53.32
4	350	34.90	150.9		73.73
5	369	22.65	92.9		75.25
6	385	11.77	46.3		76.54
7	405	11.15	41.7		78.00
8	468	9.96	32.2		81.62
9	702	7.86	17.0		91.34
10	763	8.06	16.0		92.21
11	1,010	13.59	20.4		95.61
12	1,220	24.62	30.6		96.90
13	1,484	19.82	20.2		98.52
14	1,703	16.18	14.4		99.88
15	1,879	10.58	8.5		100.96
16	2,280	5.90	3.9		102.58
17	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 9 : NT RNAP ChIP**

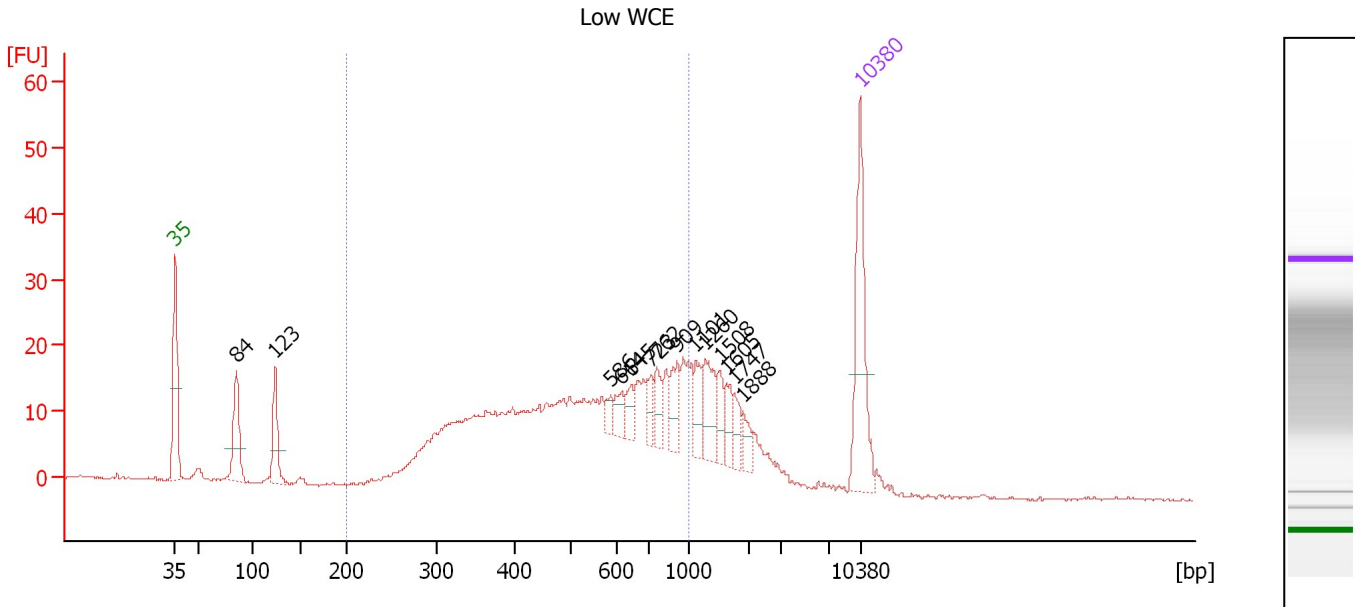
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	553	500.9	3,187.1	999.38	68	33.8



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

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 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : Low WCE**

Number of peaks found: 14                      Corr. Area 1: 418.6  
 Noise: 0.2

**Peak table for sample 10 : Low WCE**

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	84	64.21	1,154.7		49.21
3	123	47.11	580.5		53.19
4	586	8.84	22.8		87.51
5	614	14.30	35.3		88.60
6	645	13.25	31.1		89.58
7	723	14.43	30.2		91.65
8	762	18.36	36.5		92.19
9	909	23.09	38.5		94.26
10	1,101	20.34	28.0		96.17
11	1,260	32.58	39.2		97.15
12	1,508	15.64	15.7		98.67
13	1,605	14.66	13.8		99.27
14	1,747	12.26	10.6		100.14
15	1,888	10.55	8.5		101.02
16	10,380	75.00	10.9	Upper Marker	113.00

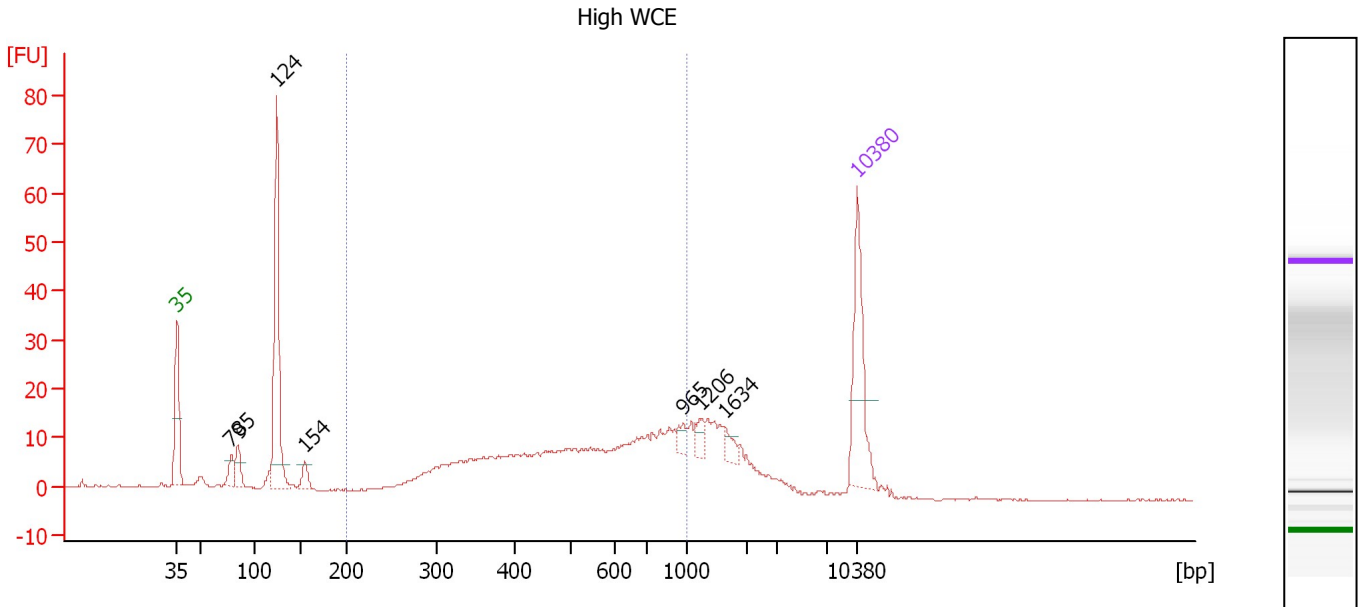
**Region table for sample 10 : Low WCE**

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	548	418.6	2,735.0	828.12	71	35.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : High WCE**

Number of peaks found: 7      Corr. Area 1: 273.5  
 Noise: 0.1

**Peak table for sample 11 : High WCE**

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	79	20.46	393.3		48.60
3	85	26.76	476.2		49.31
4	124	203.03	2,487.1		53.26
5	154	17.20	169.2		56.17
6	965	10.59	16.6		95.06
7	1,206	11.32	14.2		96.82
8	1,634	10.99	10.2		99.45
9	10,380	75.00	10.9	Upper Marker	113.00

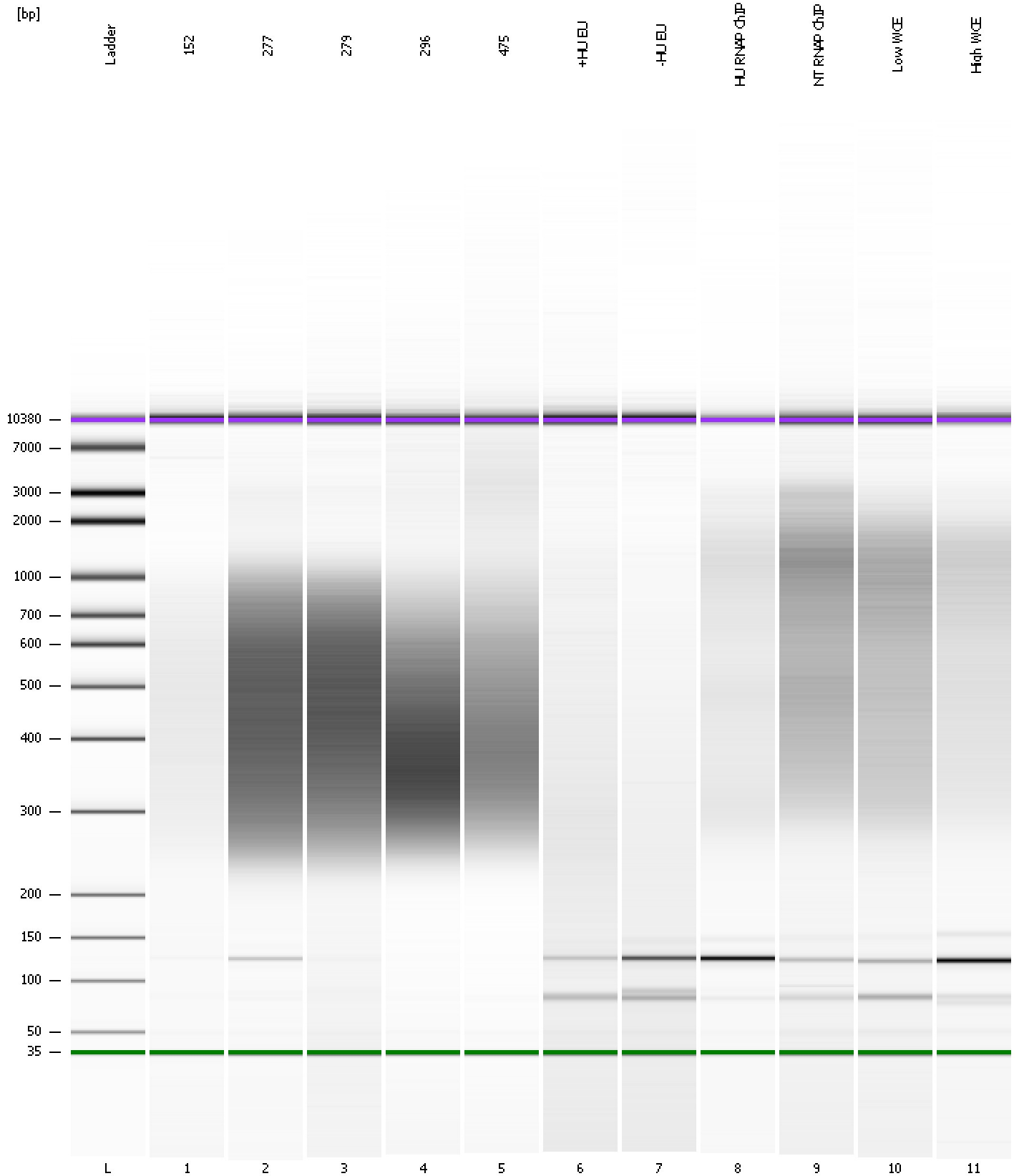
**Region table for sample 11 : High WCE**

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	557	273.5	1,623.1	500.08	56	35.3

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
Modified: 12/16/2016 4:39:22 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-12-16\2016-12-16\_001.xad

Created: 12/16/2016 3:59:58 PM  
 Modified: 12/16/2016 4:39:22 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		12/16/2016 4:41:14 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-12-16\2016-12-16_001.xad)		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/16/2016 4:00:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1