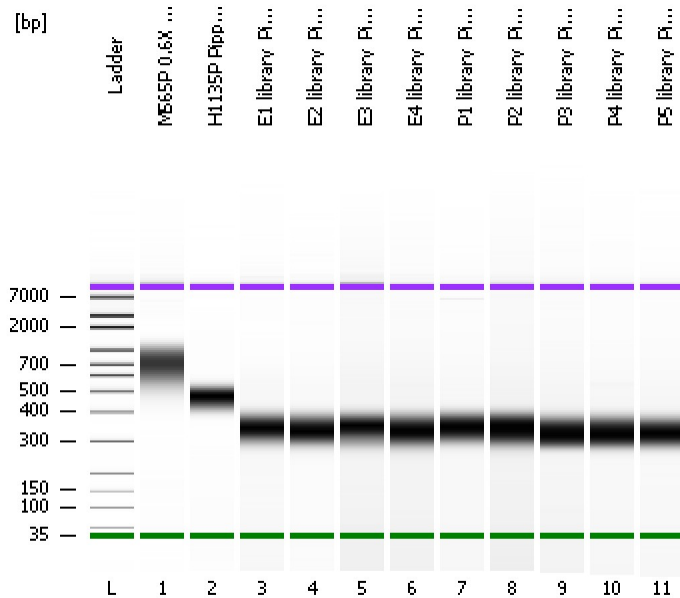


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
Modified: 9/12/2017 4:07:01 PM

Electrophoresis File Run Summary



Instrument Information:

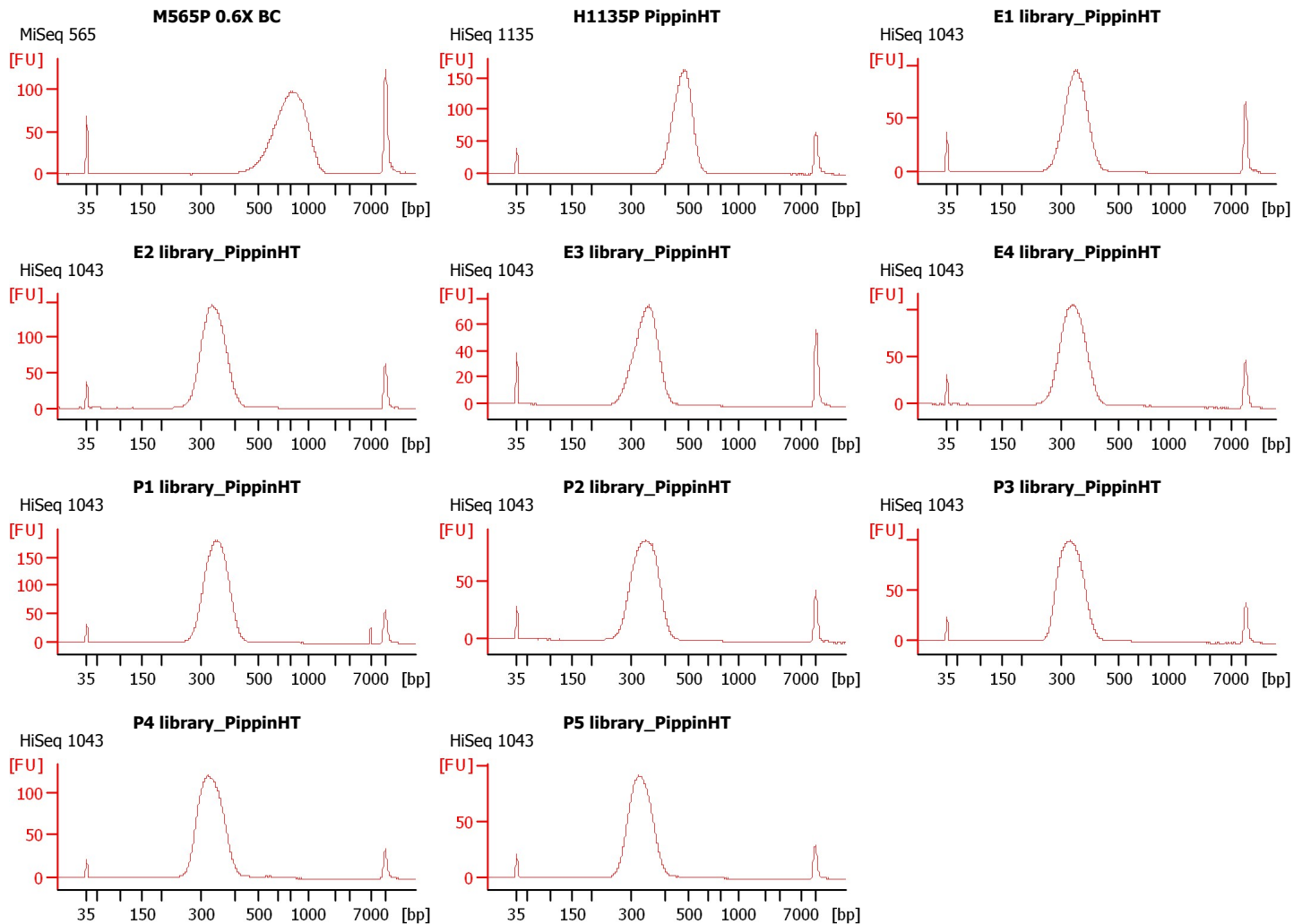
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
M565P 0.6X BC	MiSeq 565	<input type="checkbox"/>	✓			
H1135P PippinHT	HiSeq 1135	<input type="checkbox"/>	✓			
E1 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
E2 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
E3 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
E4 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
P1 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
P2 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
P3 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
P4 library_PippinHT	HiSeq 1043	<input type="checkbox"/>	✓			
P5 library_PippinHT	HiSeq 1043	<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\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
Modified: 9/12/2017 4:07:01 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

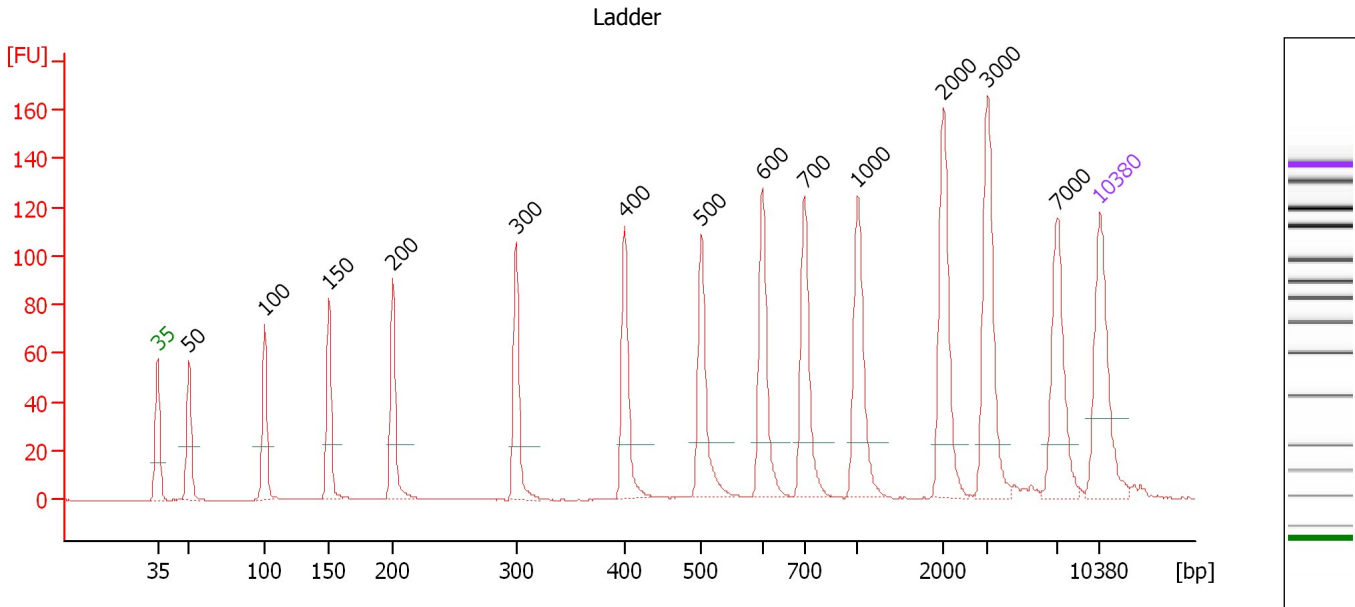
Ladder

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\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

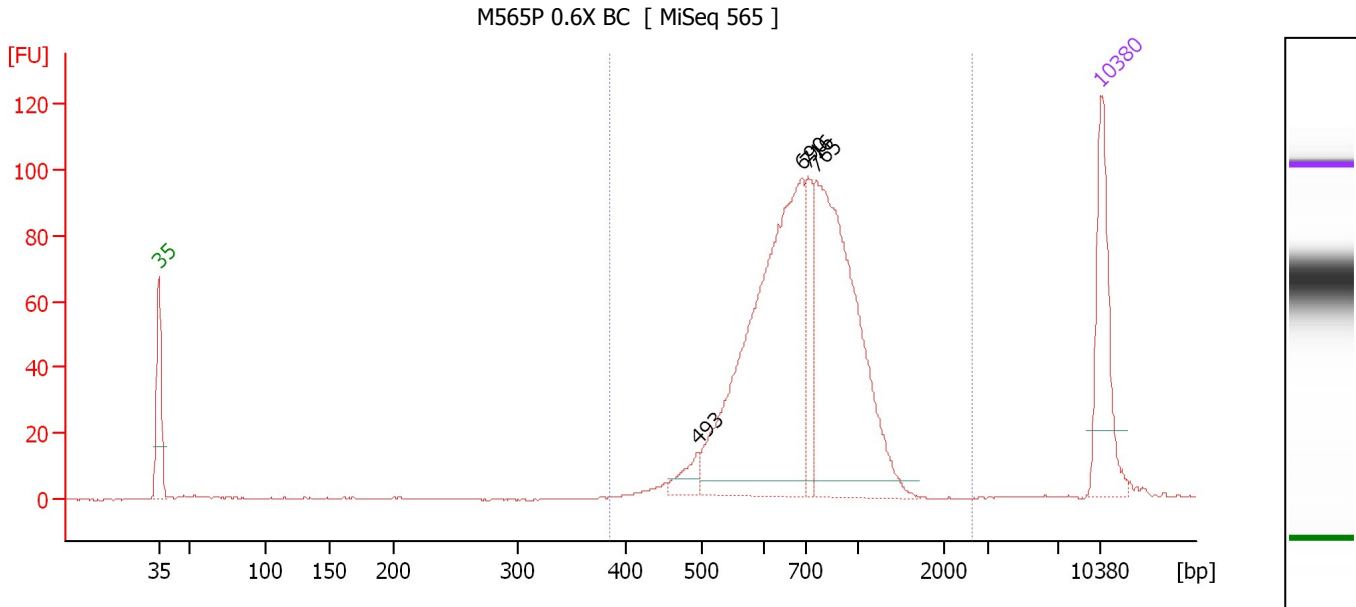
Peak table for Ladder

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	45.35
3	100	150.00	2,272.7	Ladder Peak	50.94
4	150	150.00	1,515.2	Ladder Peak	55.74
5	200	150.00	1,136.4	Ladder Peak	60.44
6	300	150.00	757.6	Ladder Peak	69.63
7	400	150.00	568.2	Ladder Peak	77.73
8	500	150.00	454.5	Ladder Peak	83.42
9	600	150.00	378.8	Ladder Peak	87.92
10	700	150.00	324.7	Ladder Peak	91.02
11	1,000	150.00	227.3	Ladder Peak	95.01
12	2,000	150.00	113.6	Ladder Peak	101.41
13	3,000	150.00	75.8	Ladder Peak	104.66
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\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : M565P 0.6X BC

Number of peaks found: 4 Corr. Area 1: 1,001.4
 Noise: 0.1

Peak table for sample 1 : M565P 0.6X BC

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	493	20.65	63.4		83.05
3	690	410.89	901.9		90.72
4	716	59.72	126.5		91.22
5	765	322.42	638.5		91.88
6	10,380	75.00	10.9	Upper Marker	113.00

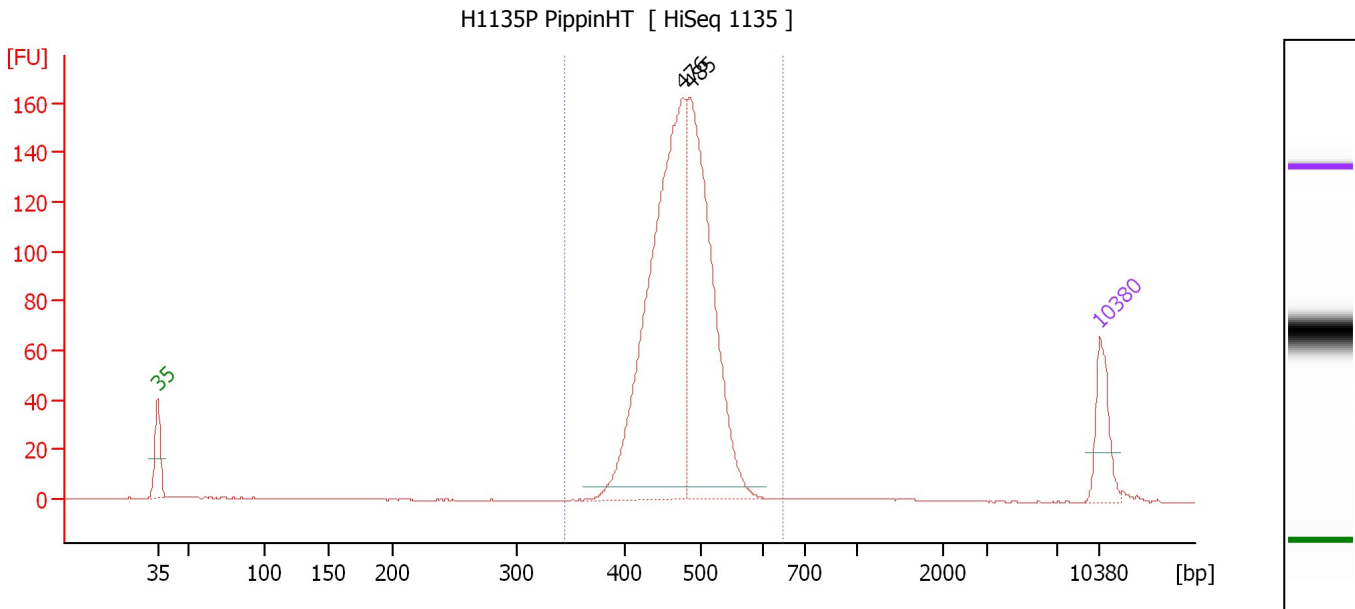
Region table for sample 1 : M565P 0.6X BC

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
385	2,617	750	1,001.4	1,795.7	825.41	96	27.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : H1135P PippinHT

Number of peaks found: 2 Corr. Area 1: 1,057.2
 Noise: 0.2

Peak table for sample 2 : H1135P PippinHT

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	476	987.95	3,146.7		82.04
3	485	706.31	2,207.7		82.55
4	10,380	75.00	10.9	Upper Marker	113.00

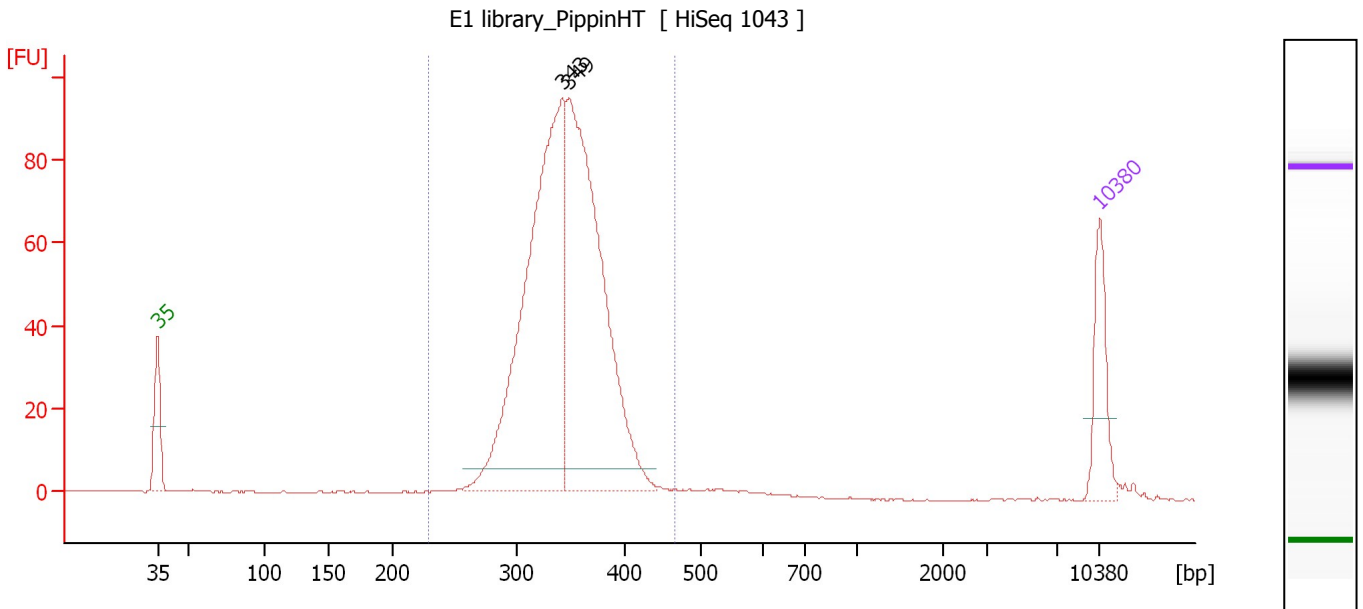
Region table for sample 2 : H1135P PippinHT

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
345	648	472	1,057.2	5,518.9	1,711.25	97	8.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : E1 library PippinHT

Number of peaks found: 2 Corr. Area 1: 853.5
 Noise: 0.1

Peak table for sample 3 : E1 library PippinHT

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	343	768.57	3,397.8		73.09
3	349	722.10	3,139.4		73.56
4	10,380	75.00	10.9	Upper Marker	113.00

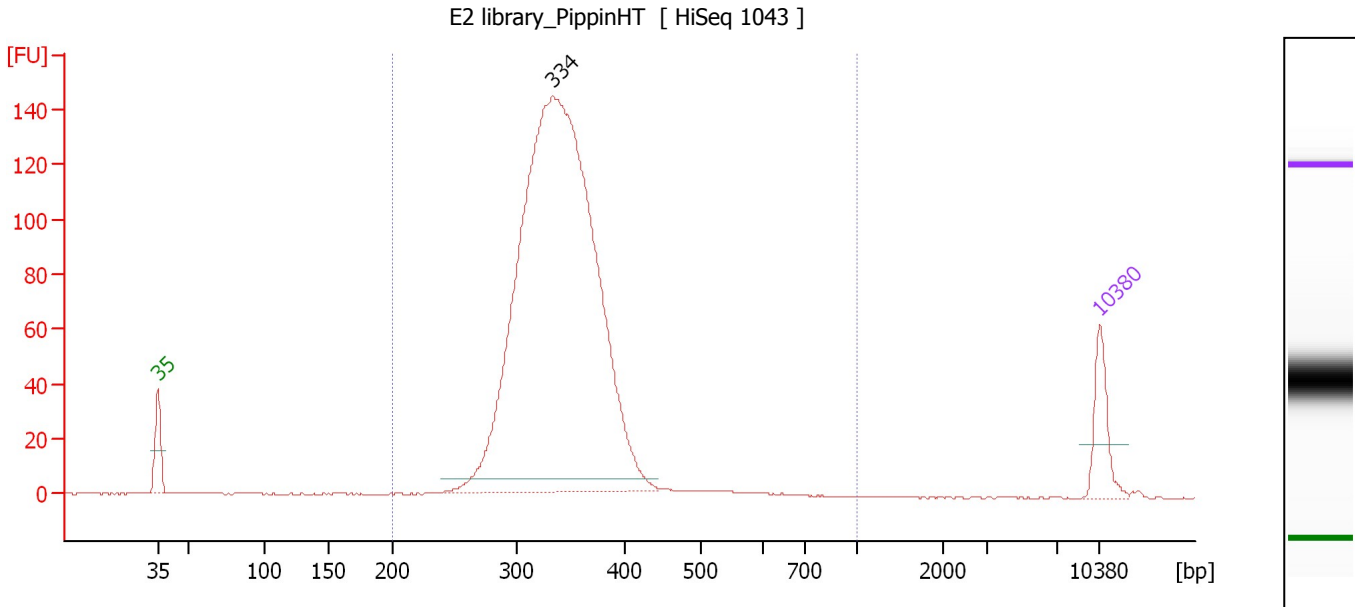
Region table for sample 3 : E1 library PippinHT

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
229	464	344	853.5	6,863.6	1,548.19	97	9.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : E2 library PippinHT

Number of peaks found: 1 Corr. Area 1: 1,403.9
 Noise: 0.2

Peak table for sample 4 : E2 library PippinHT

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	334	2,485.37	11,289.9		72.35
3	10,380	75.00	10.9	Upper Marker	113.00

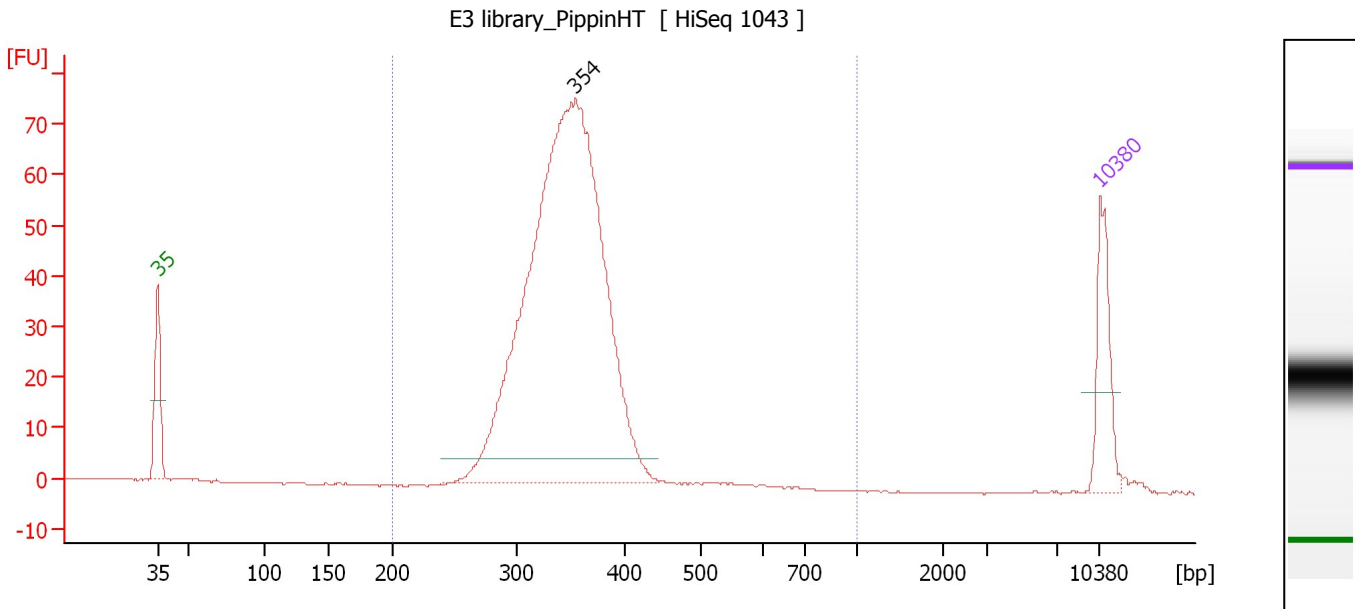
Region table for sample 4 : E2 library PippinHT

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	343	1,403.9	11,570.2	2,576.18	98	13.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : E3 library PippinHT

Number of peaks found: 1 Corr. Area 1: 695.5
 Noise: 0.1

Peak table for sample 5 : E3 library PippinHT

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	354	1,331.71	5,699.3		74.00
3	10,380	75.00	10.9	Upper Marker	113.00

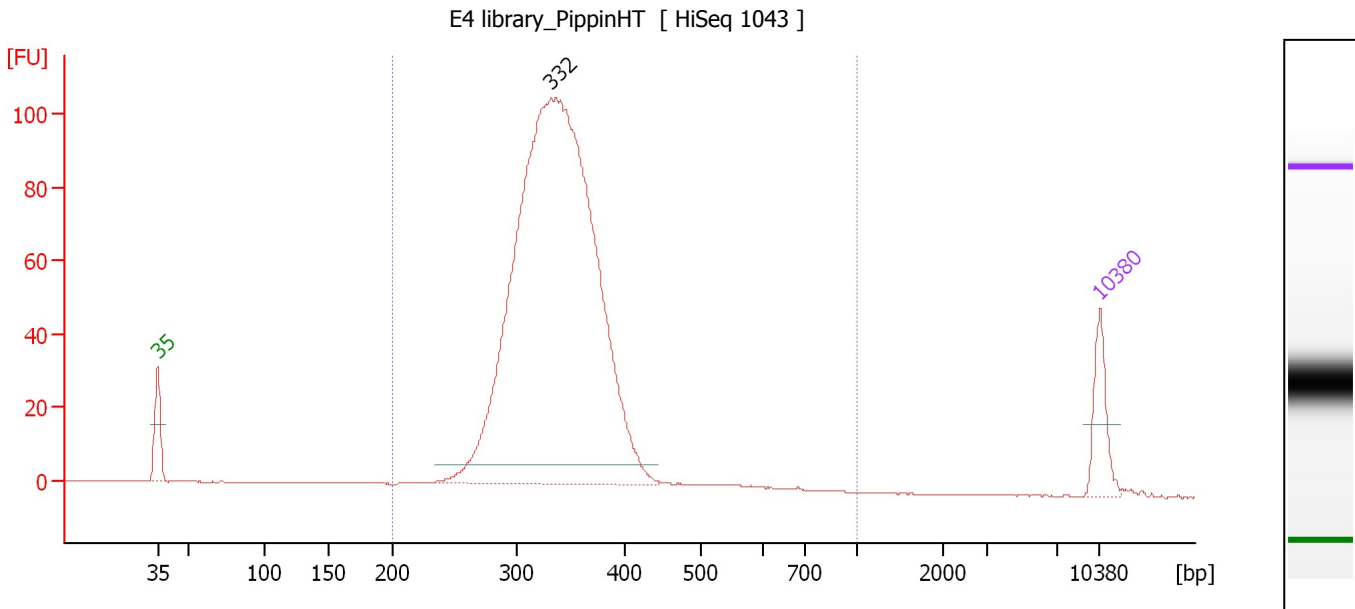
Region table for sample 5 : E3 library PippinHT

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	348	695.5	6,234.9	1,412.71	98	12.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : E4 library PippinHT

Number of peaks found: 1 Corr. Area 1: 1,082.9
 Noise: 0.1

Peak table for sample 6 : E4 library PippinHT

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	332	2,462.40	11,228.7		72.24
3	10,380	75.00	10.9	Upper Marker	113.00

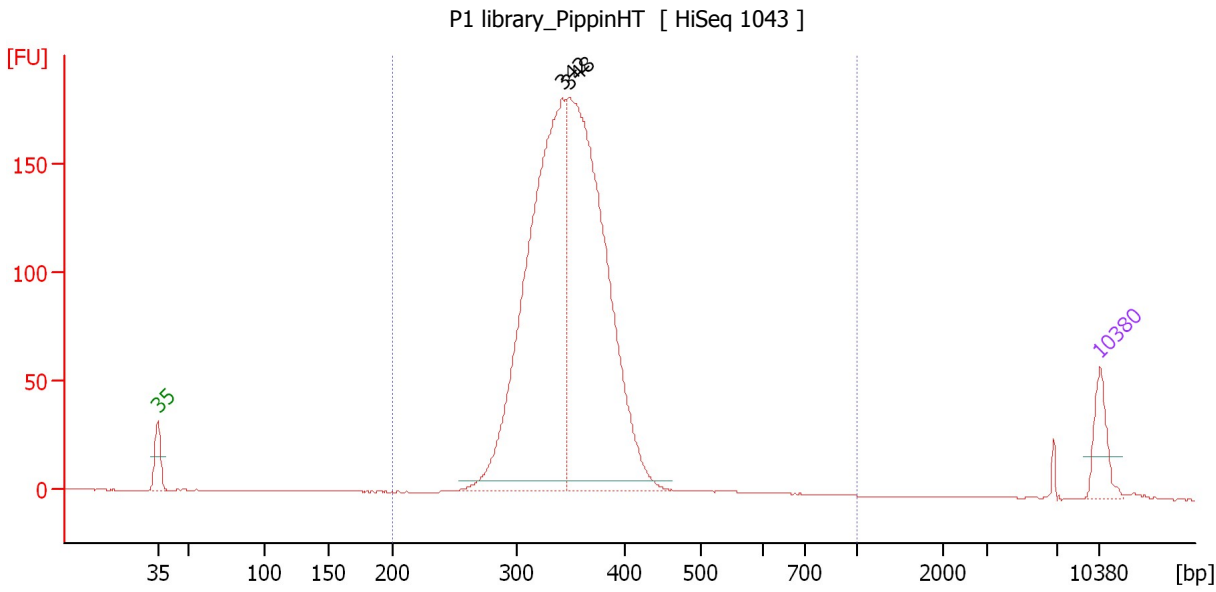
Region table for sample 6 : E4 library PippinHT

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	341	1,082.9	11,688.5	2,580.18	98	14.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : P1 library PippinHT

Number of peaks found: 2 Corr. Area 1: 1,697.7
 Noise: 0.1

Peak table for sample 7 : P1 library PippinHT

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	342	1,566.06	6,932.2		73.05
3	348	1,617.51	7,036.4		73.54
4	10,380	75.00	10.9	Upper Marker	113.00

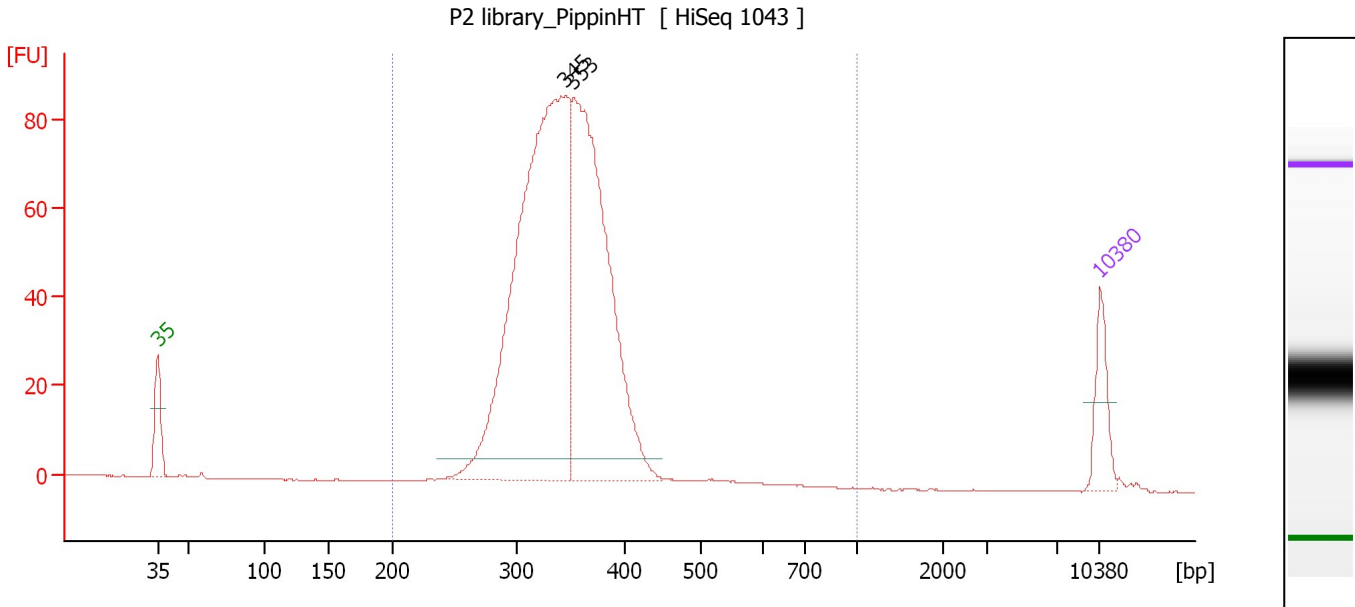
Region table for sample 7 : P1 library PippinHT

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	350	1,697.7	14,402.0	3,286.57	99	12.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : P2 library PippinHT

Number of peaks found: 2 Corr. Area 1: 914.7
 Noise: 0.1

Peak table for sample 8 : P2 library PippinHT

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	345	1,459.96	6,408.1		73.29
3	353	1,002.75	4,308.4		73.89
4	10,380	75.00	10.9	Upper Marker	113.00

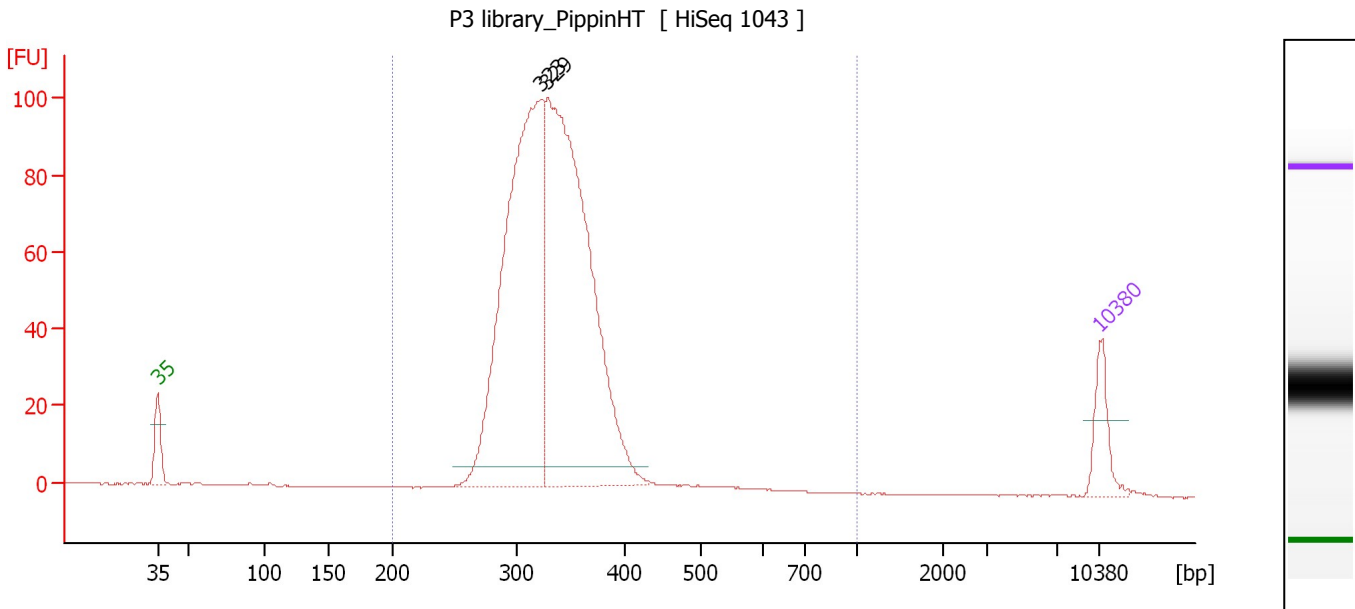
Region table for sample 8 : P2 library PippinHT

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	344	914.7	11,546.2	2,590.36	99	11.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : P3 library PippinHT

Number of peaks found: 2 Corr. Area 1: 996.6
 Noise: 0.1

Peak table for sample 9 : P3 library PippinHT

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	323	1,295.83	6,069.7		71.53
3	329	1,471.04	6,775.9		71.97
4	10,380	75.00	10.9	Upper Marker	113.00

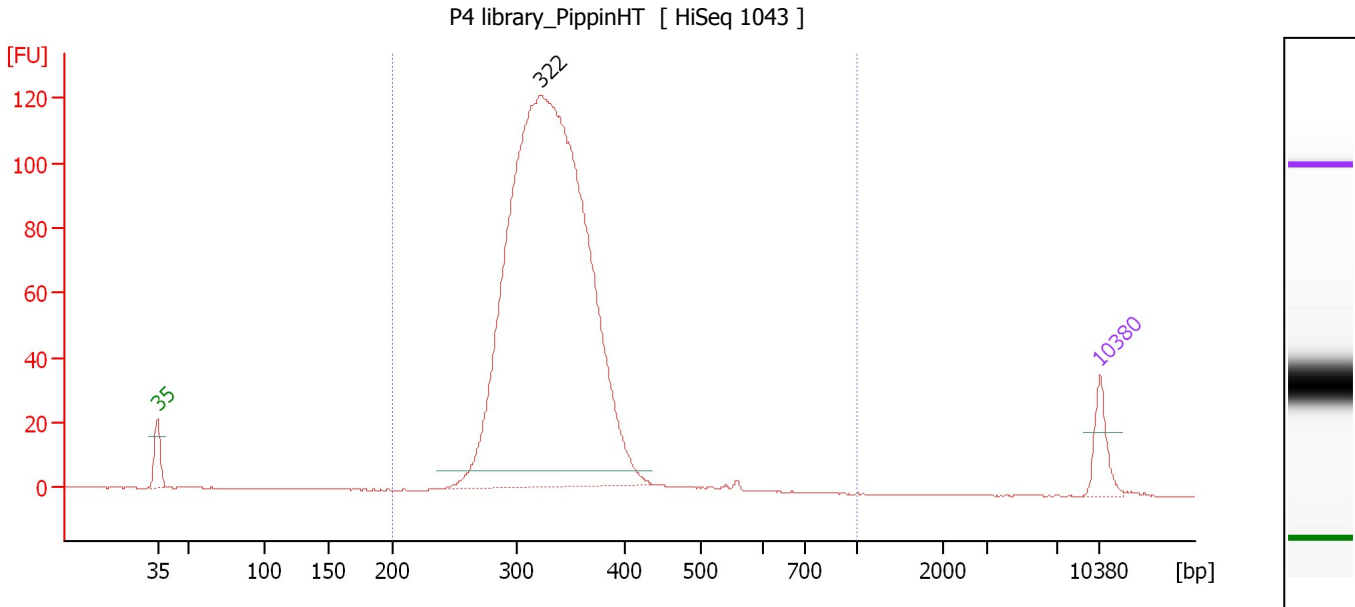
Region table for sample 9 : P3 library PippinHT

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	334	996.6	13,142.5	2,856.29	99	12.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : P4 library PippinHT

Number of peaks found: 1 Corr. Area 1: 1,191.9
 Noise: 0.1

Peak table for sample 10 : P4 library PippinHT

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	4,016.00	18,872.1		71.45
3	10,380	75.00	10.9	Upper Marker	113.00

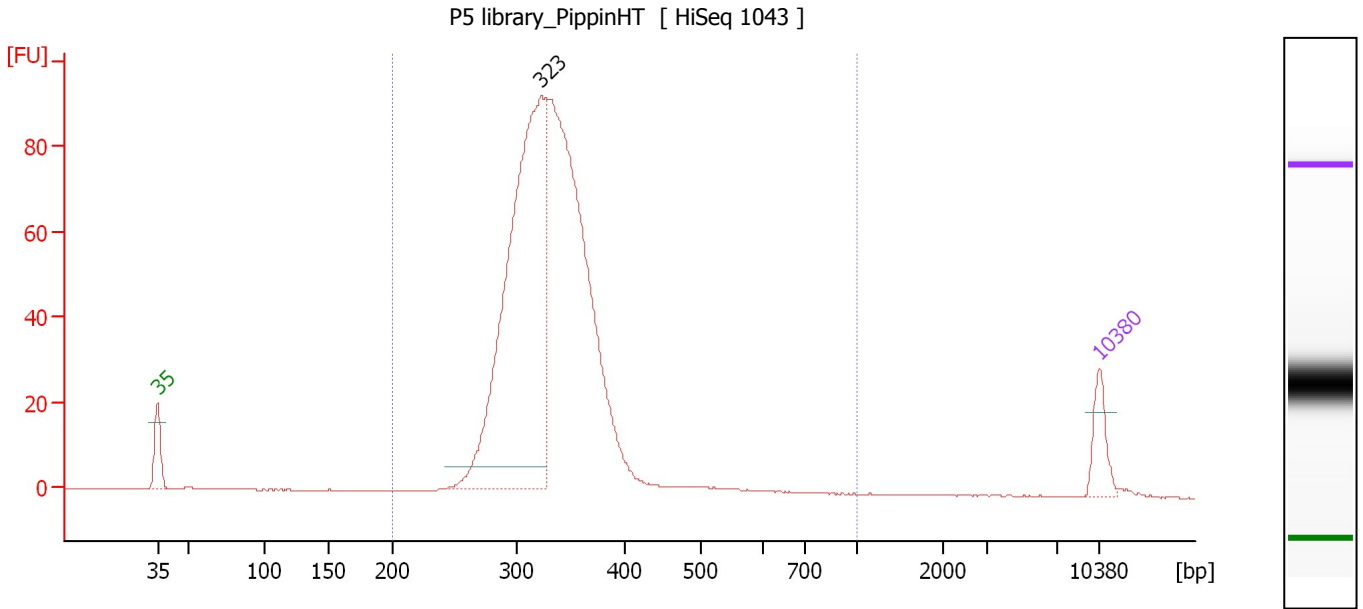
Region table for sample 10 : P4 library PippinHT

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	335	1,191.9	18,877.0	4,107.00	99	13.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
 Modified: 9/12/2017 4:07:01 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : P5 library PippinHT

Number of peaks found: 1 Corr. Area 1: 835.9
 Noise: 0.1

Peak table for sample 11 : P5 library PippinHT

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	323	1,647.86	7,729.6		71.49
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : P5 library PippinHT

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	334	835.9	15,891.1	3,449.88	99	13.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-09-12\2017-09-12_002.xad

Created: 9/12/2017 3:05:16 PM
Modified: 9/12/2017 4:07:01 PM

Gel Image

