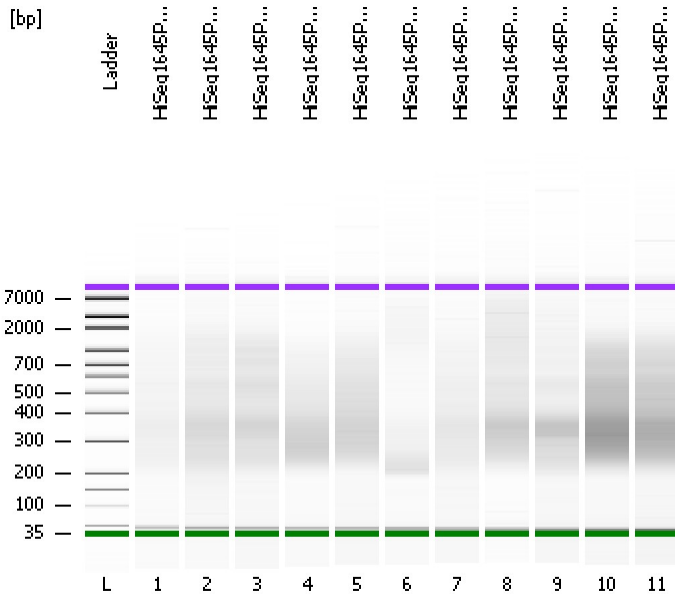


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
Modified: 12/17/2018 1:37:29 PM

Electrophoresis File Run Summary



Instrument Information:

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

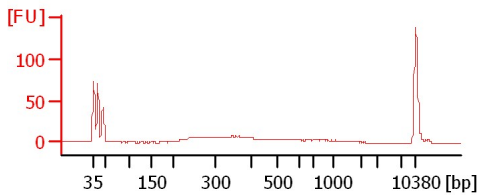
Assay Information:

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

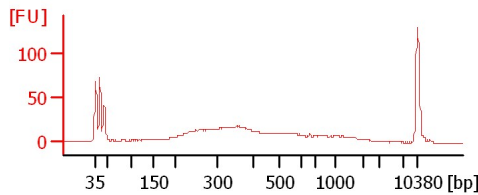
Chip Information:

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

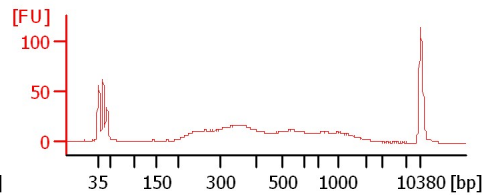
HiSeq1645P_Questa_Amp4X_6



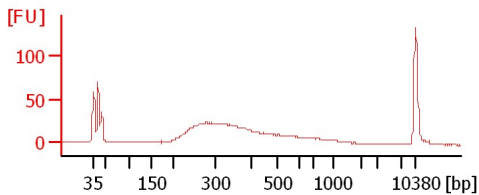
HiSeq1645P_Questa_Amp4X_8



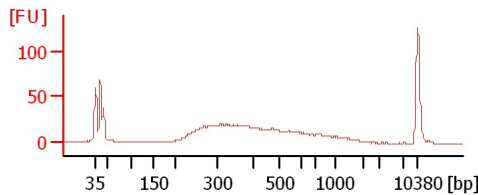
HiSeq1645P_Questa_Amp4X_17



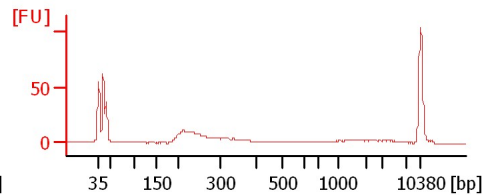
HiSeq1645P_Questa_Amp4X_23



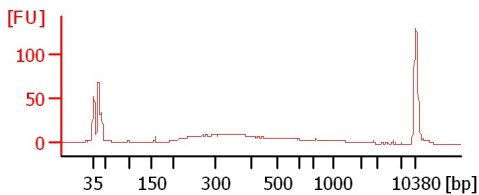
HiSeq1645P_Questa_Amp4X_31



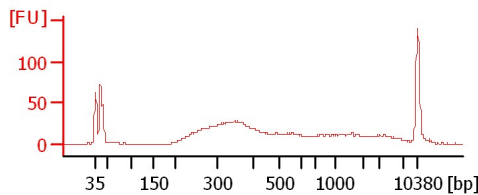
HiSeq1645P_Questa_Amp4X_11



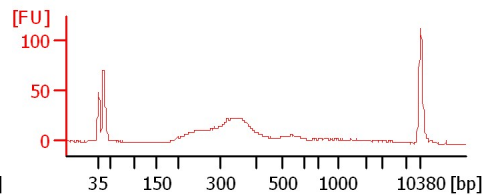
HiSeq1645P_Questa_Amp4X_14



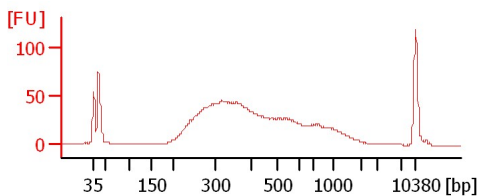
HiSeq1645P_Questa_Amp4X_28



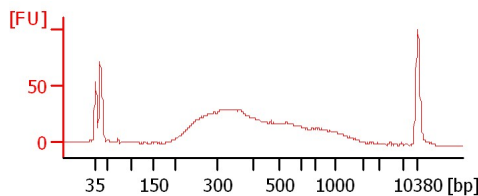
HiSeq1645P_Questa_Amp4X_2



HiSeq1645P_Questa_Amp4X_24



HiSeq1645P_Questa_Amp4X_25



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
HiSeq1645P_Questa_Amp4X_6		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_8		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_17		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_23		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_31		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_11		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_14		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_28		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_2		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_24		<input type="checkbox"/>	✓			
HiSeq1645P_Questa_Amp4X_25		<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\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
Modified: 12/17/2018 1:37:29 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

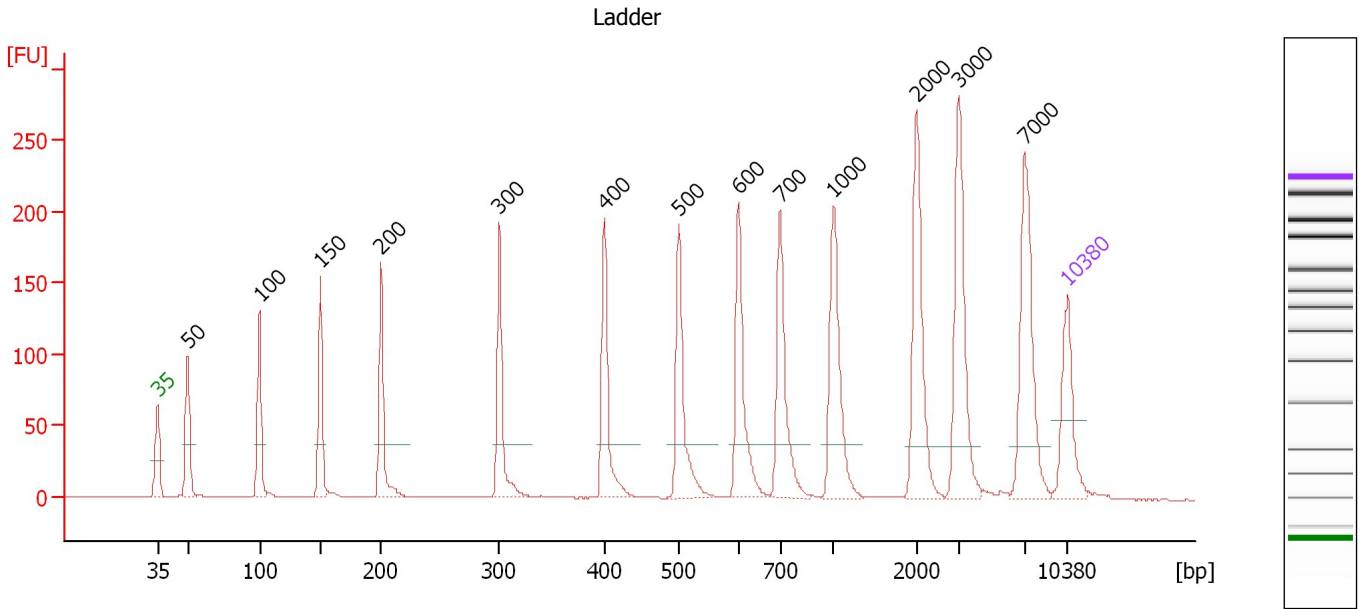
Ladder

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\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

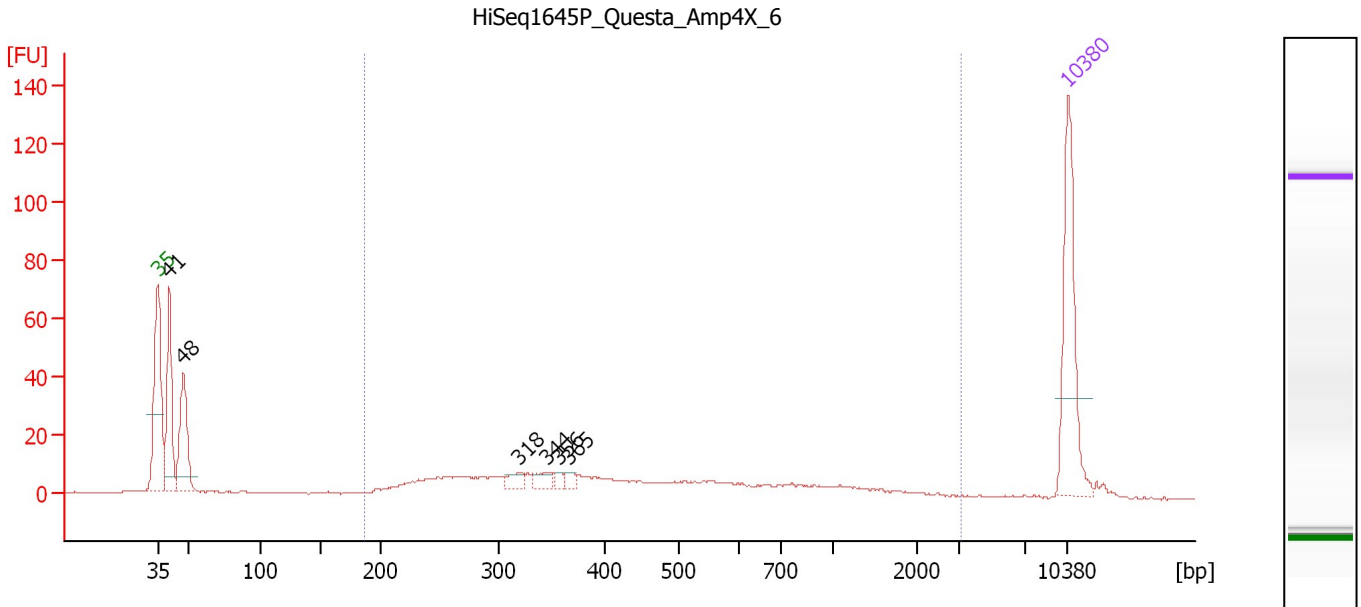
Peak table for Ladder

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	45.33
3	100	150.00	2,272.7	Ladder Peak	50.85
4	150	150.00	1,515.2	Ladder Peak	55.51
5	200	150.00	1,136.4	Ladder Peak	60.17
6	300	150.00	757.6	Ladder Peak	69.30
7	400	150.00	568.2	Ladder Peak	77.38
8	500	150.00	454.5	Ladder Peak	83.09
9	600	150.00	378.8	Ladder Peak	87.70
10	700	150.00	324.7	Ladder Peak	90.89
11	1,000	150.00	227.3	Ladder Peak	94.98
12	2,000	150.00	113.6	Ladder Peak	101.35
13	3,000	150.00	75.8	Ladder Peak	104.63
14	7,000	150.00	32.5	Ladder Peak	109.67
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : HiSeq1645P_Questa_Amp4X_6

Number of peaks found: 6 Corr. Area 1: 274.7
 Noise: 0.2

Peak table for sample 1 : HiSeq1645P_Questa_Amp4X_6

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	97.57	3,624.2		43.90
3	48	83.14	2,635.5		44.99
4	318	10.83	51.6		70.73
5	344	10.33	45.5		72.86
6	356	4.93	21.0		73.81
7	365	6.17	25.6		74.57
8	10,380	75.00	10.9	Upper Marker	113.00

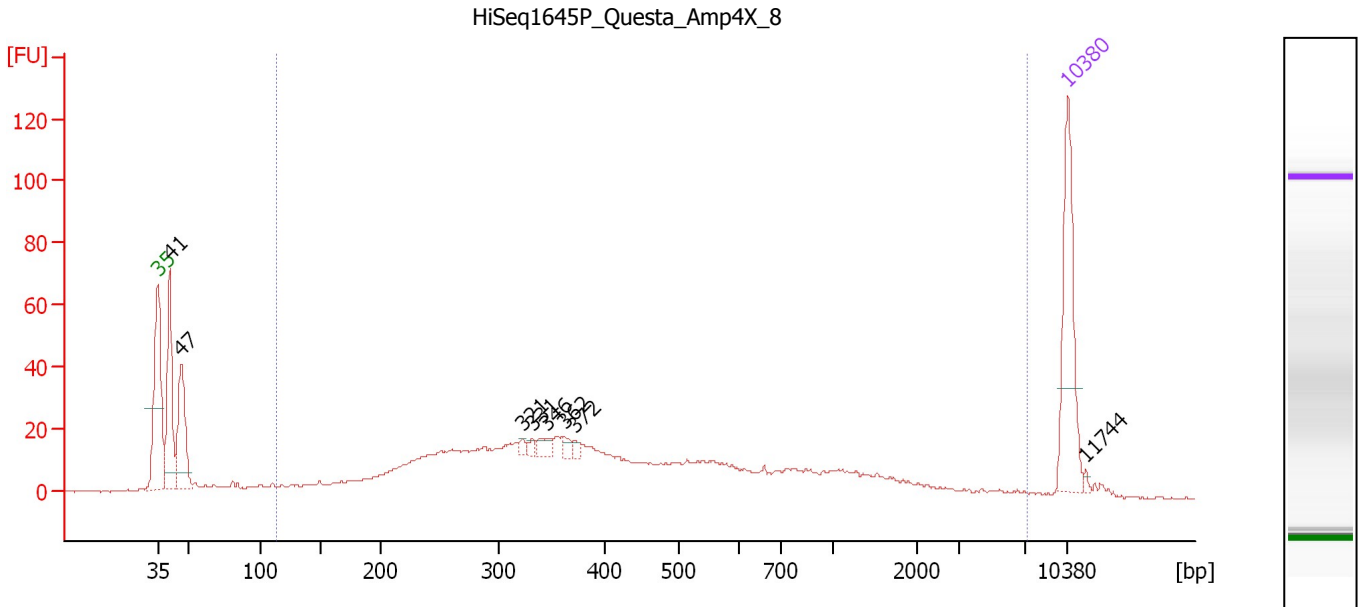
Region table for sample 1 : HiSeq1645P_Questa_Amp4X_6

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
186	3,086	548	246.73	274.7	1,023.7	62	77.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : HiSeq1645P_Questa_Amp4X_8

Number of peaks found: 8 Corr. Area 1: 690.6
 Noise: 0.3

Peak table for sample 2 : HiSeq1645P_Questa_Amp4X_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	41	116.90	4,334.1		43.91
3	47	90.46	2,932.6		44.82
4	321	4.44	20.9		71.02
5	331	4.63	21.2		71.79
6	346	10.20	44.7		72.99
7	362	6.95	29.1		74.28
8	372	4.89	19.9		75.15
9	10,380	75.00	10.9	Upper Marker	113.00
10	11,744	0.00	0.0		114.34

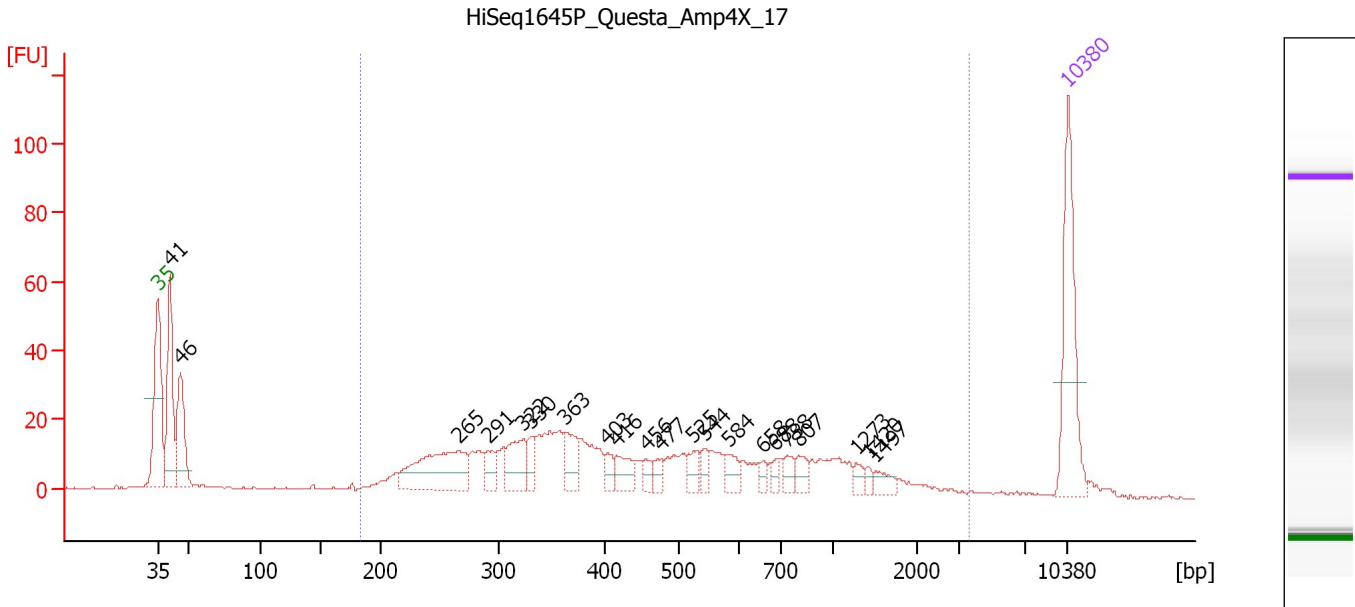
Region table for sample 2 : HiSeq1645P_Questa_Amp4X_8

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
114	7,167	638	730.42	690.6	3,364.8	78	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : HiSeq1645P_Questa_Amp4X_17

Number of peaks found: 21 Corr. Area 1: 599.5
 Noise: 0.2

Peak table for sample 3 : HiSeq1645P_Questa_Amp4X_17


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	109.83	4,057.0		43.93
3	46	78.19	2,553.3		44.77
4	265	95.06	543.1		66.12
5	291	16.97	88.3		68.48
6	322	38.11	179.2		71.09
7	330	16.17	74.2		71.73
8	363	24.34	101.6		74.38
9	403	10.26	38.6		77.53
10	416	19.91	72.6		78.27
11	456	9.68	32.1		80.58
12	477	9.77	31.1		81.76
13	525	13.52	39.0		84.22
14	544	10.34	28.8		85.11
15	584	13.49	35.0		86.98
16	658	6.94	16.0		89.54
17	688	6.82	15.0		90.52
18	738	9.67	19.9		91.40
19	807	11.45	21.5		92.34
20	1,273	6.62	7.9		96.72
21	1,420	4.13	4.4		97.65
22	1,497	9.00	9.1		98.14
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...

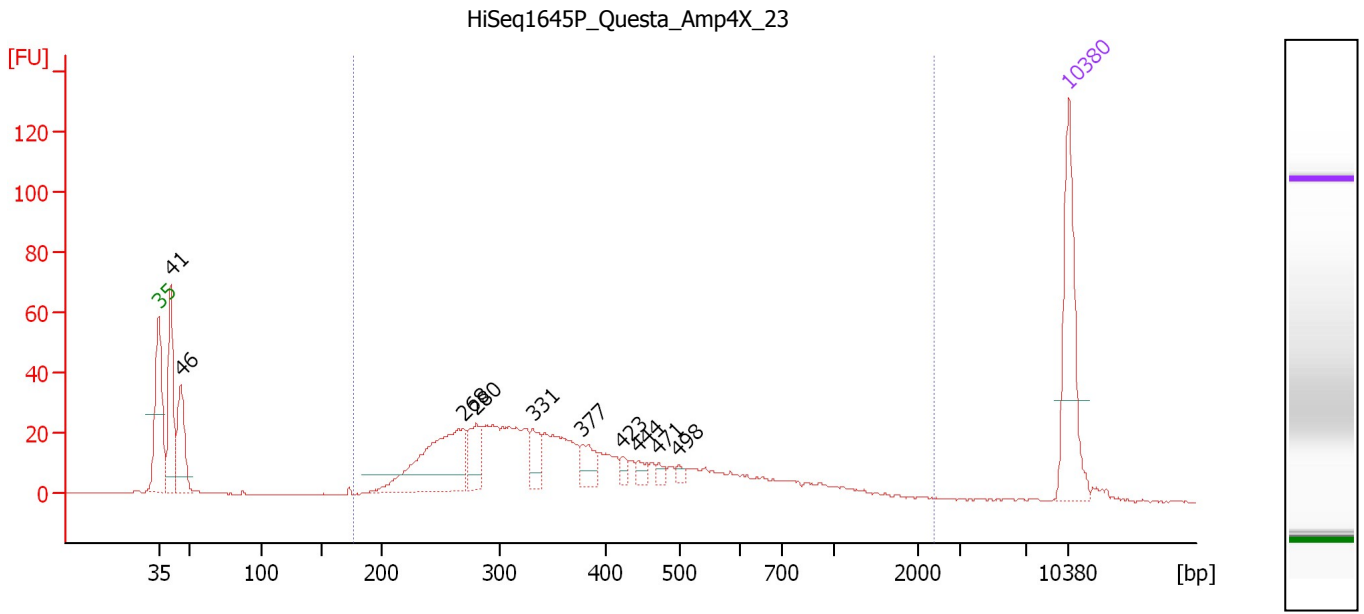
... Region table for sample 3 : HiSeq1645P_Questa_Amp4X_17

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
183	3,627	584	649.39	599.5	2,580.8	 78	76.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HiSeq1645P Questa Amp4X 23

Number of peaks found: 10 Corr. Area 1: 648.8
 Noise: 0.2

Peak table for sample 4 : HiSeq1645P Questa Amp4X 23

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	106.02	3,902.2		43.96
3	46	70.57	2,323.0		44.71
4	268	135.06	763.3		66.38
5	280	33.98	184.1		67.44
6	331	24.06	110.3		71.78
7	377	20.87	83.8		75.56
8	423	7.01	25.1		78.68
9	444	7.85	26.8		79.89
10	471	5.05	16.2		81.45
11	498	4.64	14.1		82.96
12	10,380	75.00	10.9	Upper Marker	113.00

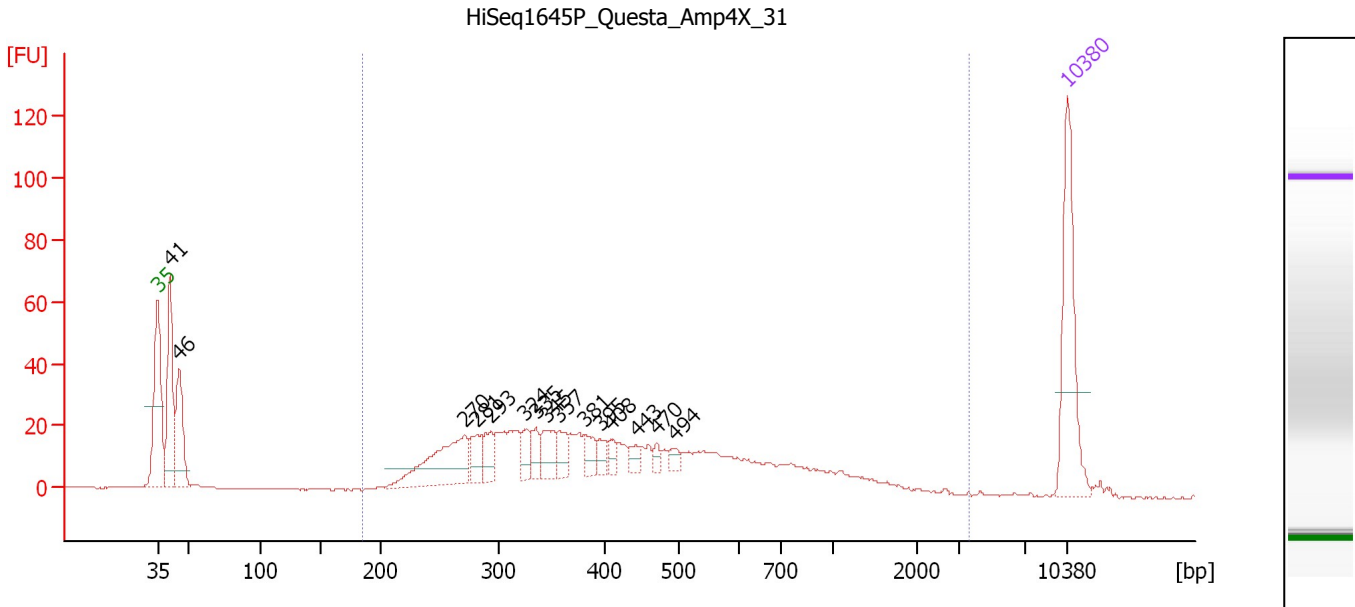
Region table for sample 4 : HiSeq1645P Questa Amp4X 23

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
177	2,394	415	640.94	648.8	2,906.7	82	55.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : HiSeq1645P Questa Amp4X 31

Number of peaks found: 15 Corr. Area 1: 663.7
 Noise: 0.2

Peak table for sample 5 : HiSeq1645P Questa Amp4X 31

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	106.64	3,940.2		43.93
3	46	71.07	2,357.2		44.66
4	270	85.82	481.0		66.59
5	281	22.32	120.3		67.58
6	293	24.93	128.9		68.67
7	324	16.45	77.0		71.21
8	335	16.89	76.5		72.09
9	345	24.60	108.1		72.92
10	357	19.05	80.8		73.90
11	381	14.53	57.7		75.87
12	395	9.24	35.4		77.01
13	408	9.41	34.9		77.84
14	443	8.32	28.5		79.81
15	470	6.14	19.8		81.37
16	494	6.78	20.8		82.77
17	10,380	75.00	10.9	Upper Marker	113.00

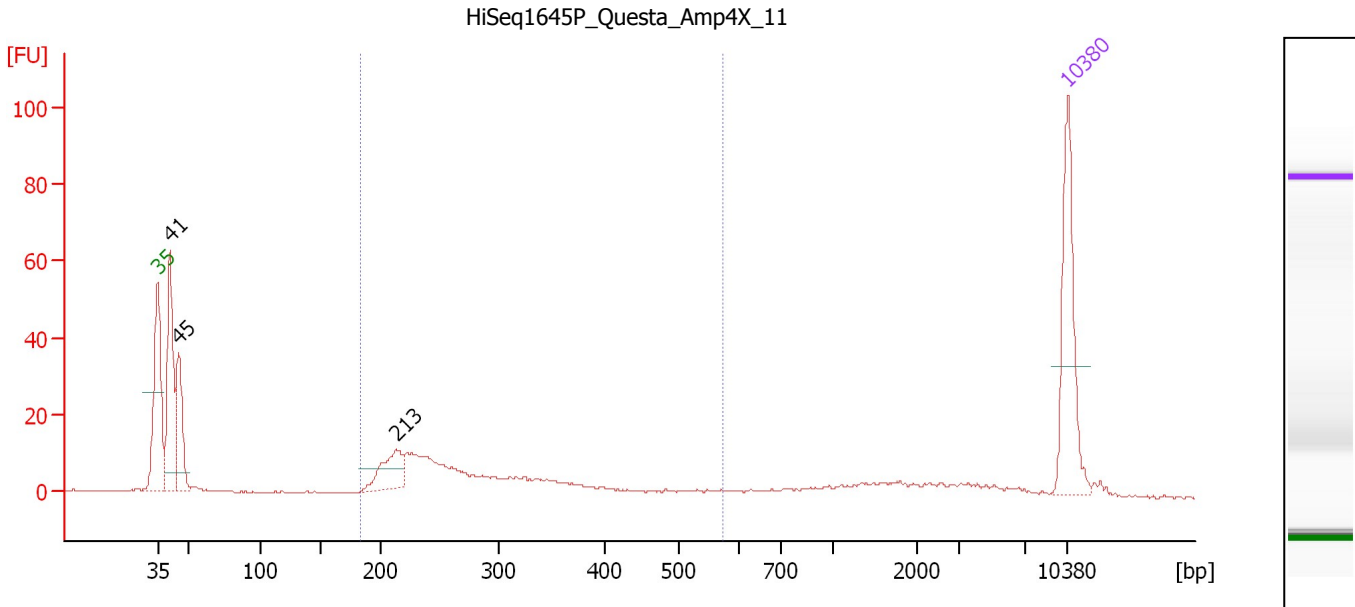
Region table for sample 5 : HiSeq1645P Questa Amp4X 31

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
185	3,639	506	631.21	663.7	2,573.1	82	69.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : HiSeq1645P_Questa_Amp4X_11

Number of peaks found: 3 Corr. Area 1: 163.4
 Noise: 0.2

Peak table for sample 6 : HiSeq1645P_Questa_Amp4X_11

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	126.63	4,661.6		43.96
3	45	77.20	2,584.0		44.59
4	213	43.71	311.2		61.34
5	10,380	75.00	10.9	Upper Marker	113.00

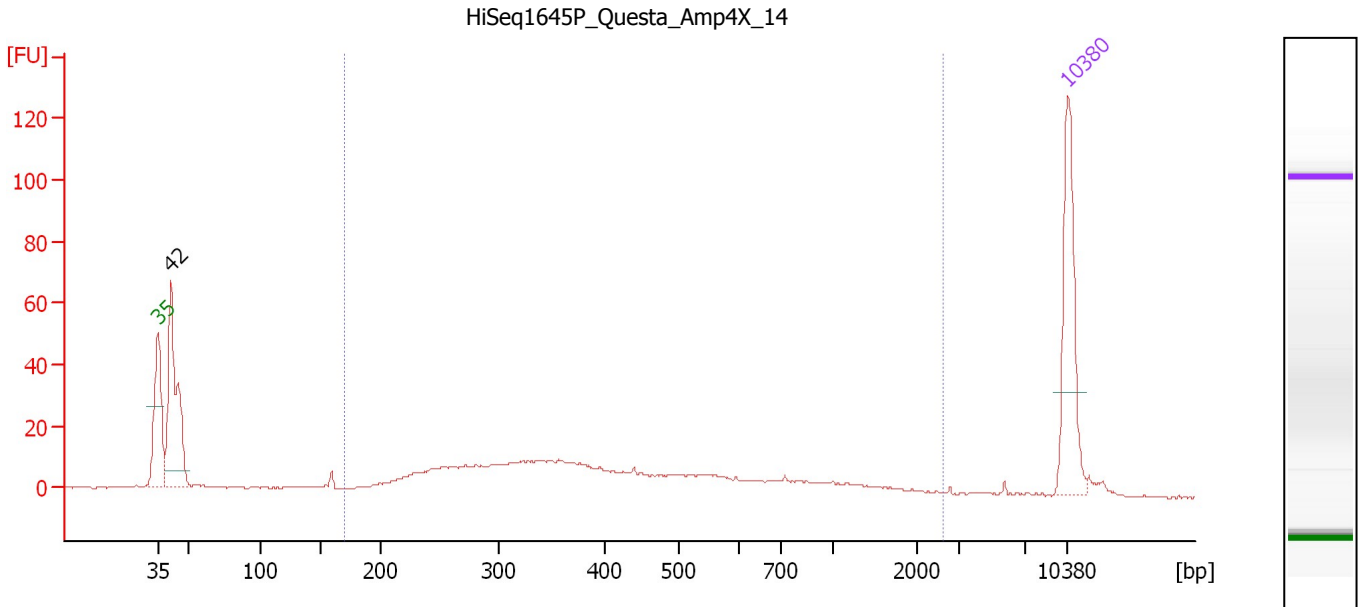
Region table for sample 6 : HiSeq1645P_Questa_Amp4X_11

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
183	575	286	214.02	163.4	1,236.7	46	28.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : HiSeq1645P_Questa_Amp4X_14

Number of peaks found: 1 Corr. Area 1: 321.3
 Noise: 0.2

Peak table for sample 7 : HiSeq1645P_Questa_Amp4X_14

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	42	164.49	5,987.1		44.03
3	10,380	75.00	10.9	Upper Marker	113.00

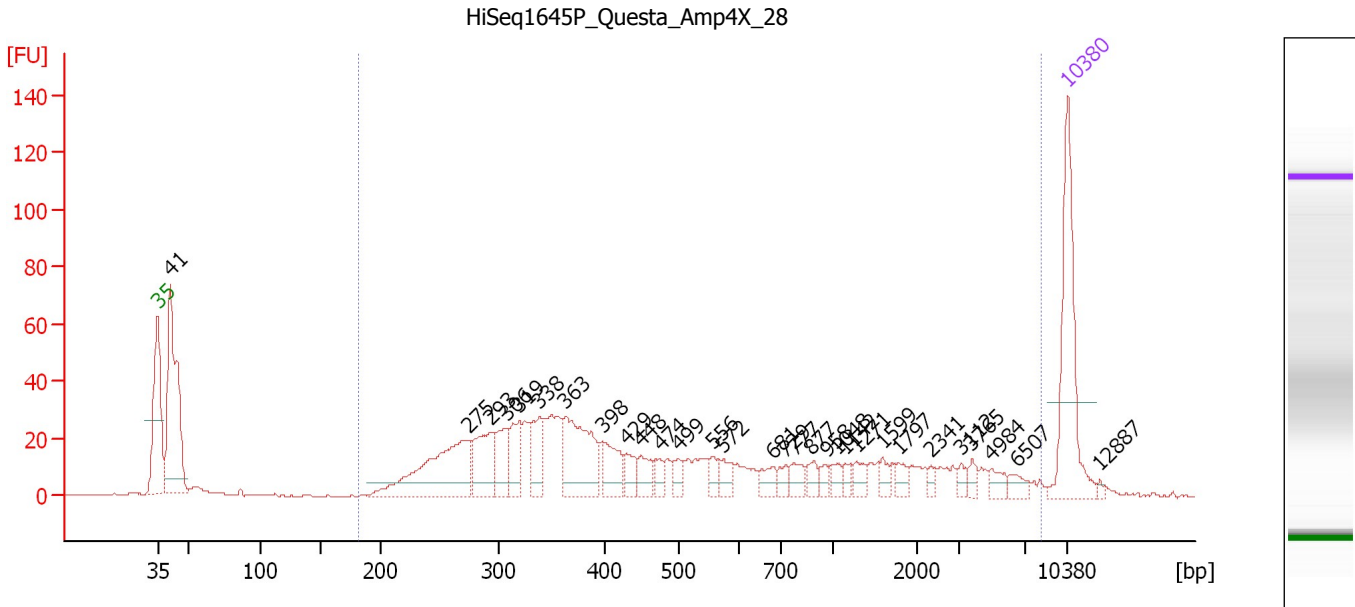
Region table for sample 7 : HiSeq1645P_Questa_Amp4X_14

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
171	2,603	494	311.36	321.3	1,336.8	67	69.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : HiSeq1645P_Questa_Amp4X_28

Number of peaks found: 30 Corr. Area 1: 889.0
 Noise: 0.3

Peak table for sample 8 : HiSeq1645P_Questa_Amp4X_28

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	176.76	6,475.8		43.99
3	275	114.14	629.9		66.97
4	293	49.20	254.8		68.62
5	306	31.94	158.2		69.77
6	319	31.86	151.4		70.81
7	338	34.03	152.7		72.35
8	363	77.10	322.0		74.38
9	398	30.07	114.4		77.23
10	429	13.98	49.4		79.04
11	448	15.90	53.7		80.14
12	474	10.00	31.9		81.62
13	499	9.70	29.4		83.05
14	556	9.44	25.7		85.68
15	572	11.42	30.3		86.39
16	681	12.11	26.9		90.29
17	729	8.36	17.4		91.28
18	777	10.76	21.0		91.93
19	877	8.76	15.1		93.31
20	958	5.79	9.2		94.40
21	1,048	7.60	11.0		95.28
22	1,142	5.83	7.7		95.88
23	1,271	8.92	10.6		96.71
24	1,599	7.42	7.0		98.79
25	1,797	7.08	6.0		100.05

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...

... Peak table for sample 8 : HiSeq1645P_Questa_Amp4X_28

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,341	4.46	2.9		102.47
27	3,112	4.88	2.4		104.77
28	3,765	5.55	2.2		105.59
29	4,984	7.61	2.3		107.13
30	6,507	7.01	1.6		109.05
31	10,380	75.00	10.9	Upper Marker	113.00
32	12,887	0.00	0.0		115.47

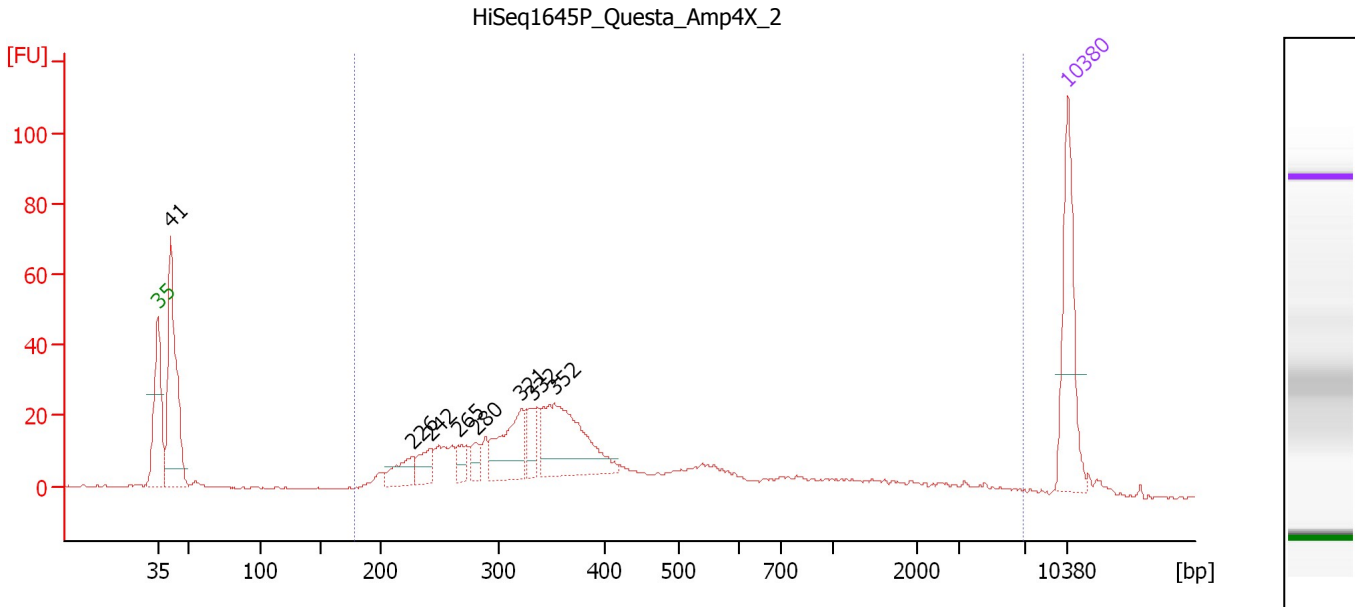
Region table for sample 8 : HiSeq1645P_Questa_Amp4X_28

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
181	8,295	1,010	733.20	889.0	2,768.2	83	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : HiSeq1645P_Questa_Amp4X_2

Number of peaks found: 8 Corr. Area 1: 525.9
 Noise: 0.3

Peak table for sample 9 : HiSeq1645P_Questa_Amp4X_2

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	197.33	7,221.5		43.99
3	226	28.47	190.7		62.56
4	242	26.15	163.8		63.99
5	265	15.32	87.6		66.09
6	280	15.08	81.6		67.48
7	321	66.25	313.2		70.96
8	332	28.50	130.0		71.90
9	352	118.66	510.8		73.50
10	10,380	75.00	10.9	Upper Marker	113.00

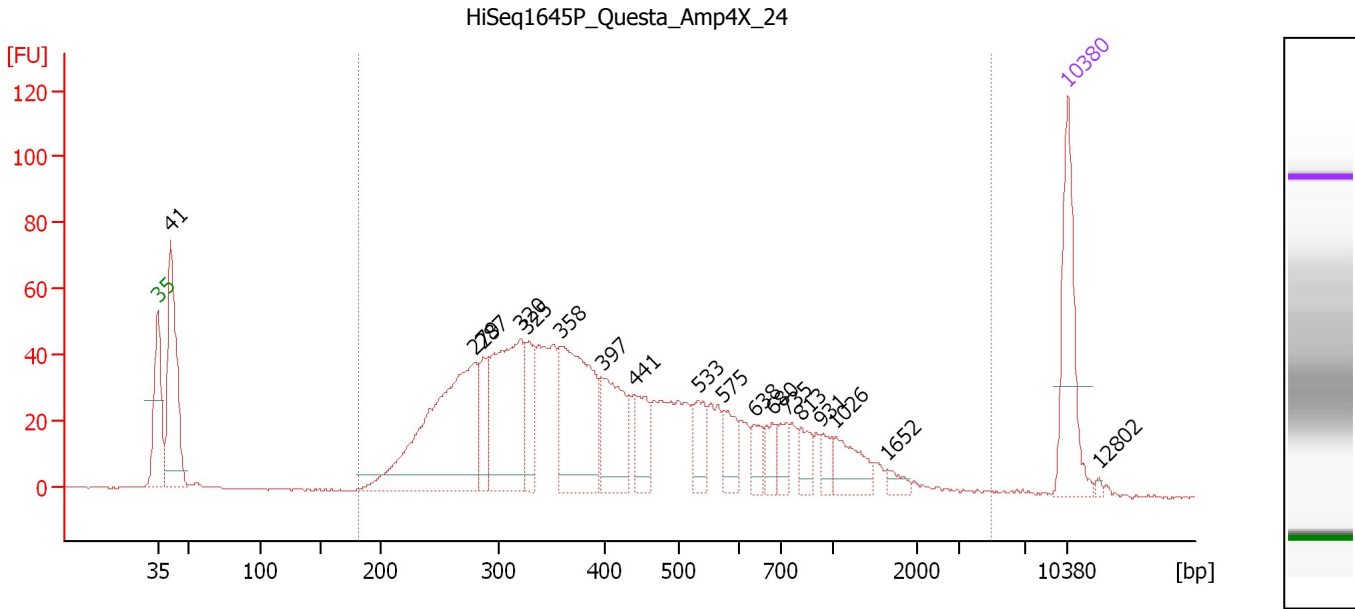
Region table for sample 9 : HiSeq1645P_Questa_Amp4X_2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
178	6,934	621	613.72	525.9	2,691.1	80	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : HiSeq1645P_Questa_Amp4X_24

Number of peaks found: 18 Corr. Area 1: 1,396.5
 Noise: 0.2

Peak table for sample 10 : HiSeq1645P_Questa_Amp4X_24

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	185.81	6,795.7		44.00
3	279	279.82	1,518.1		67.41
4	287	45.50	240.6		68.07
5	320	174.62	827.3		70.90
6	325	46.66	217.3		71.34
7	358	170.62	723.0		73.95
8	397	85.25	325.1		77.17
9	441	41.39	142.2		79.72
10	533	34.16	97.2		84.60
11	575	29.43	77.6		86.54
12	638	20.16	47.8		88.93
13	680	19.25	42.9		90.26
14	735	18.19	37.5		91.37
15	813	20.40	38.0		92.42
16	931	13.55	22.1		94.03
17	1,026	33.91	50.1		95.14
18	1,652	8.23	7.6		99.13
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,802	0.00	0.0		115.39

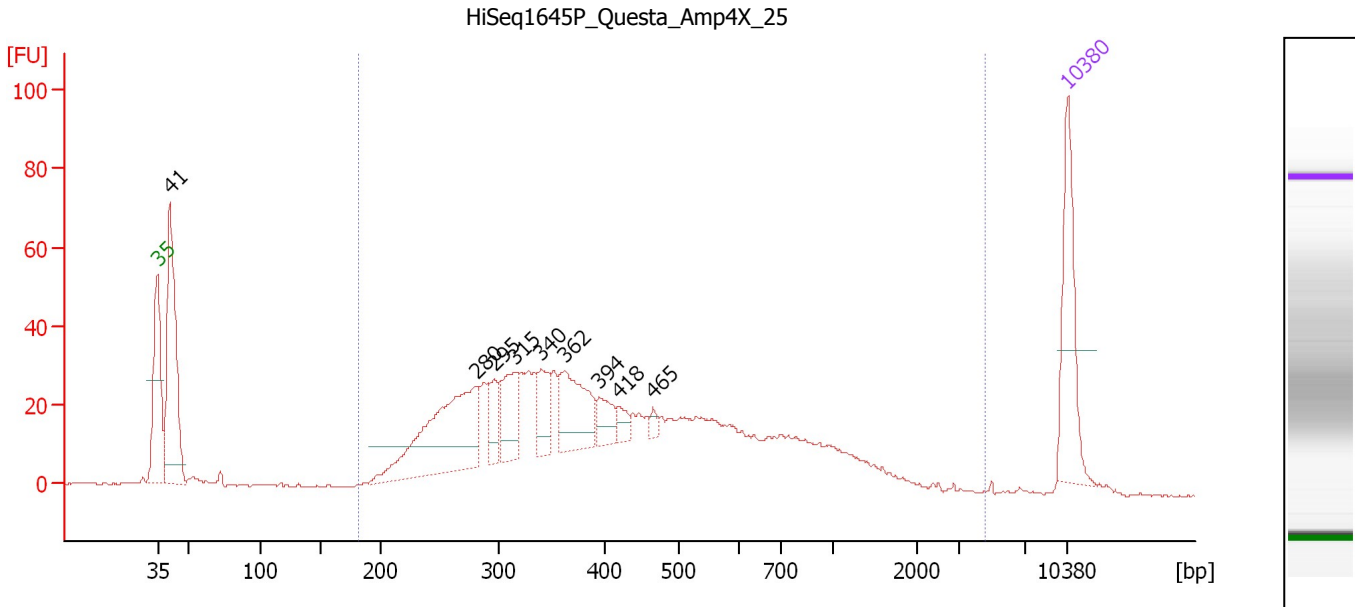
Region table for sample 10 : HiSeq1645P_Questa_Amp4X_24

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
182	4,996	505	1,378.45	1,396.5	5,685.1	90	74.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

Created: 12/17/2018 12:45:44 PM
 Modified: 12/17/2018 1:37:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : HiSeq1645P_Questa_Amp4X_25

Number of peaks found: 9 Corr. Area 1: 937.0
 Noise: 0.2

Peak table for sample 11 : HiSeq1645P_Questa_Amp4X_25

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	225.83	8,322.6		43.95
3	280	190.66	1,031.8		67.47
4	295	31.93	163.9		68.87
5	315	61.28	295.0		70.49
6	340	35.57	158.7		72.50
7	362	78.30	328.0		74.28
8	394	24.63	94.7		76.91
9	418	12.50	45.3		78.42
10	465	6.22	20.2		81.10
11	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : HiSeq1645P_Questa_Amp4X_25

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
181	4,552	503	1,182.82	937.0	4,905.0	86	71.7

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-12-17\2018-12-17_002.xad

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Gel Image

