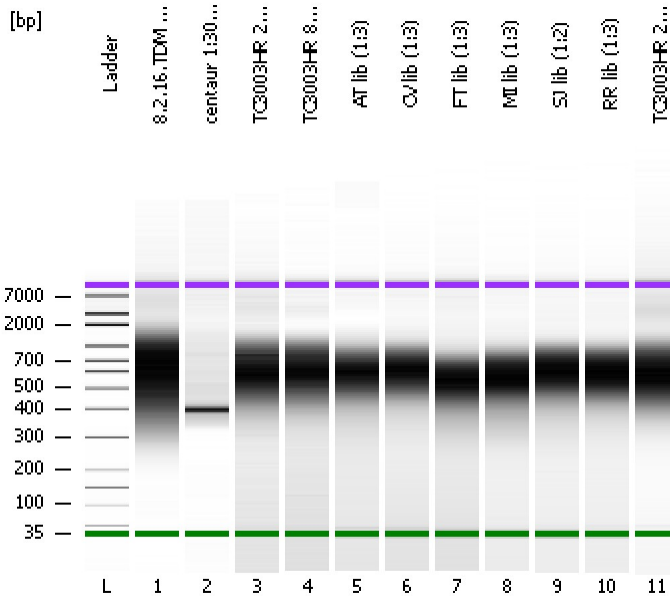


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
Modified: 8/4/2016 4:38:30 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

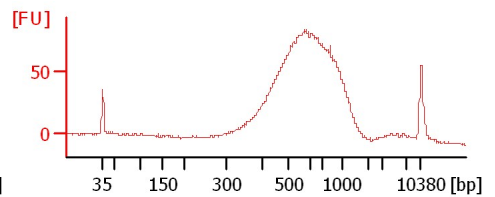
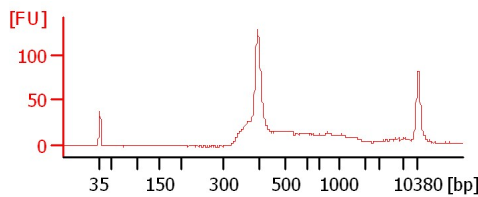
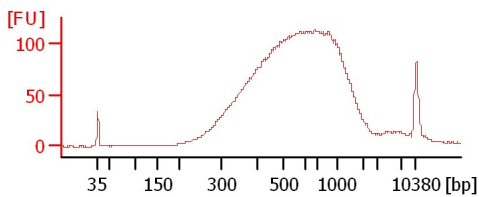
Chip Information:

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

8.2.16.TDM (1:2)

centaur 1:30 (1:2)

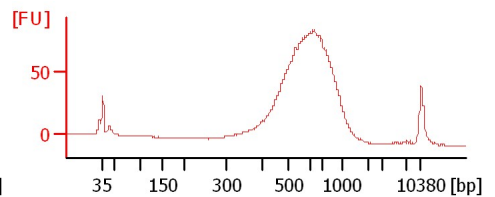
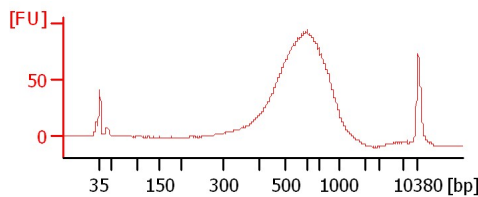
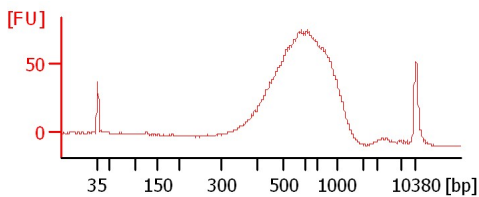
TC3003HR 2kb MP lib



TC3003HR 8kb MP lib

AT lib (1:3)

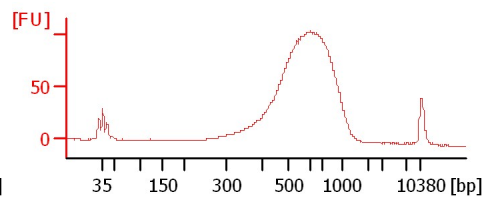
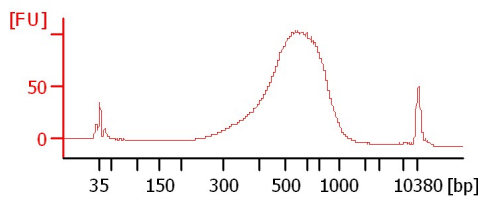
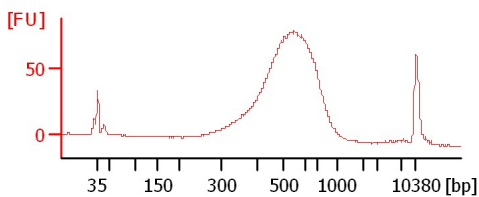
CV lib (1:3)



FT lib (1:3)

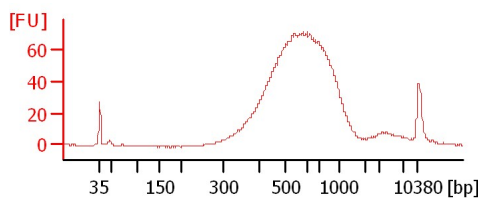
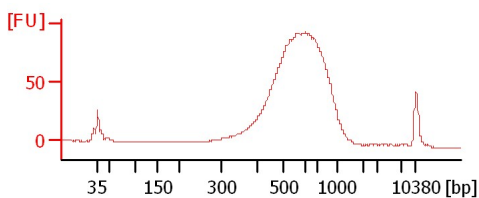
MI lib (1:3)

SJ lib (1:2)



RR lib (1:3)

TC3003HR 2kb MP lib



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
Modified: 8/4/2016 4:38:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
8.2.16.TDM (1:2)		<input type="checkbox"/>	✓			
centaur 1:30 (1:2)		<input type="checkbox"/>	✓			
TC3003HR 2kb MP lib		<input type="checkbox"/>	✓			
TC3003HR 8kb MP lib		<input type="checkbox"/>	✓			
AT lib (1:3)		<input type="checkbox"/>	✓			
CV lib (1:3)		<input type="checkbox"/>	✓			
FT lib (1:3)		<input type="checkbox"/>	✓			
MI lib (1:3)		<input type="checkbox"/>	✓			
SJ lib (1:2)		<input type="checkbox"/>	✓			
RR lib (1:3)		<input type="checkbox"/>	✓			
TC3003HR 2kb MP lib		<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\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
Modified: 8/4/2016 4:38:30 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

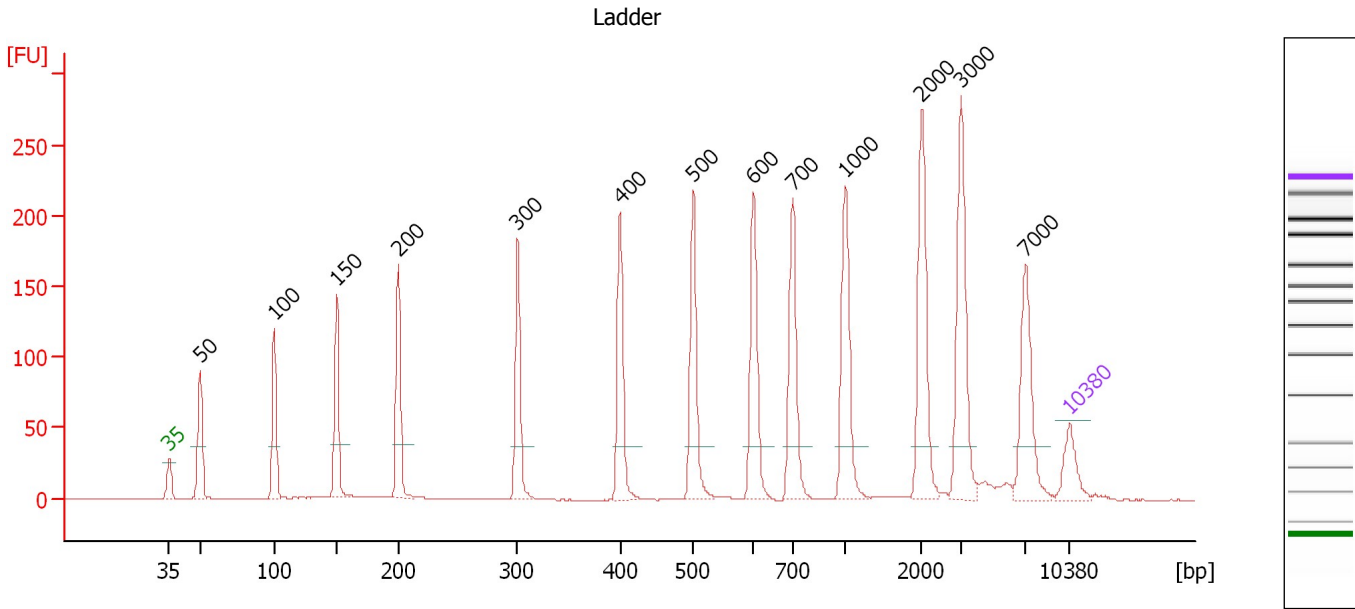
Ladder

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\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

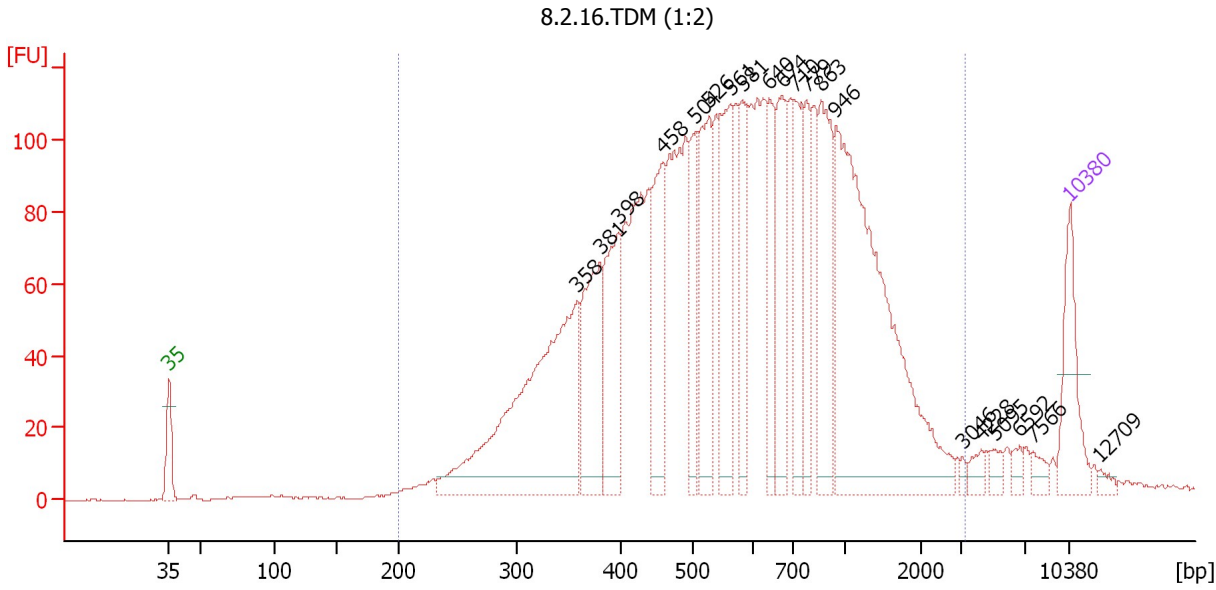
Peak table for Ladder

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	45.45
3	100	150.00	2,272.7	Ladder Peak	51.17
4	150	150.00	1,515.2	Ladder Peak	56.06
5	200	150.00	1,136.4	Ladder Peak	60.85
6	300	150.00	757.6	Ladder Peak	70.10
7	400	150.00	568.2	Ladder Peak	78.07
8	500	150.00	454.5	Ladder Peak	83.75
9	600	150.00	378.8	Ladder Peak	88.44
10	700	150.00	324.7	Ladder Peak	91.48
11	1,000	150.00	227.3	Ladder Peak	95.54
12	2,000	150.00	113.6	Ladder Peak	101.50
13	3,000	150.00	75.8	Ladder Peak	104.59
14	7,000	150.00	32.5	Ladder Peak	109.58
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 8.2.16.TDM (1:2)

Number of peaks found: 20 Corr. Area 1: 3,171.3
 Noise: 0.3

Peak table for sample 1 : 8.2.16.TDM (1:2)

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	358	497.36	2,102.7		74.76
3	381	179.80	715.8		76.53
4	398	148.91	567.1		77.90
5	458	150.32	496.8		81.39
6	504	95.42	286.7		83.95
7	526	160.83	463.0		84.98
8	561	158.34	427.8		86.60
9	581	92.78	242.1		87.54
10	640	90.70	214.8		89.65
11	674	133.88	301.0		90.68
12	710	97.63	208.2		91.62
13	779	86.44	168.1		92.55
14	863	154.45	271.2		93.68
15	946	532.61	852.7		94.81
16	3,046	6.09	3.0		104.64
17	4,228	12.97	4.6		106.12
18	5,095	11.68	3.5		107.20
19	6,592	9.83	2.3		109.07
20	7,566	12.20	2.4		110.15
21	10,380	75.00	10.9	Upper Marker	113.00
22	12,709	0.00	0.0		115.36

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...

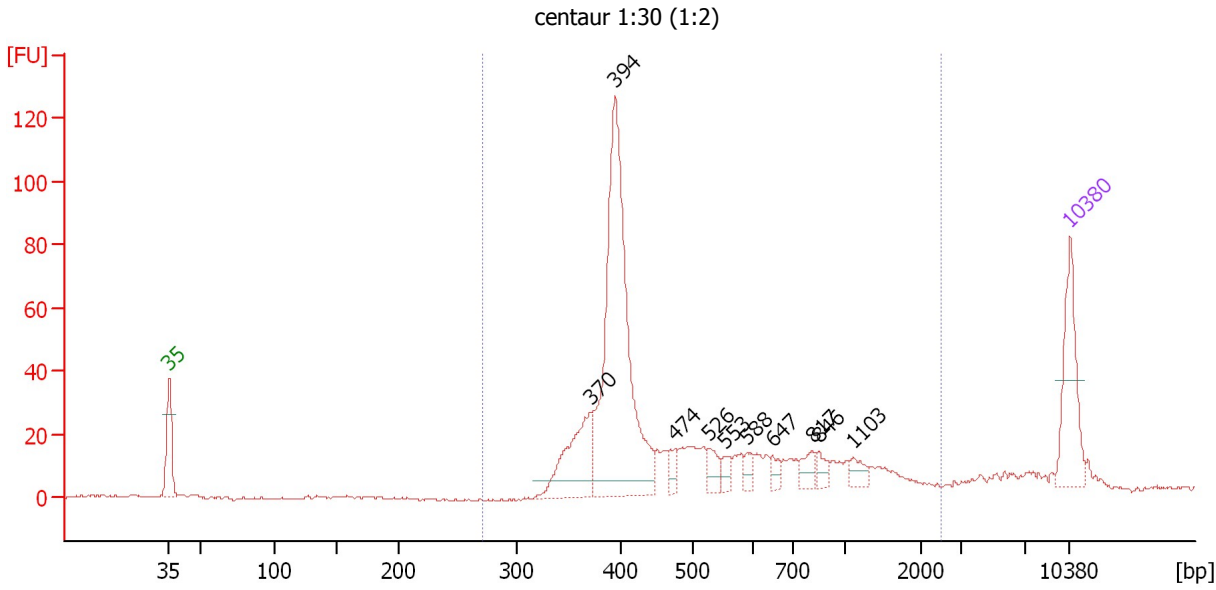
... Region table for sample 1 : **8.2.16.TDM (1:2)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	3,217	697	3,171.3	11,225.5	3,784.17	96	60.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : centaur 1:30 (1:2)

Number of peaks found: 10 Corr. Area 1: 662.3
 Noise: 0.4

Peak table for sample 2 : centaur 1:30 (1:2)

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	370	118.88	486.2		75.72
3	394	446.30	1,714.3		77.63
4	474	16.35	52.2		82.30
5	526	20.86	60.1		84.95
6	553	12.41	34.0		86.23
7	588	13.35	34.4		87.90
8	647	12.56	29.4		89.86
9	817	17.95	33.3		93.06
10	846	12.82	23.0		93.45
11	1,103	13.00	17.9		96.15
12	10,380	75.00	10.9	Upper Marker	113.00

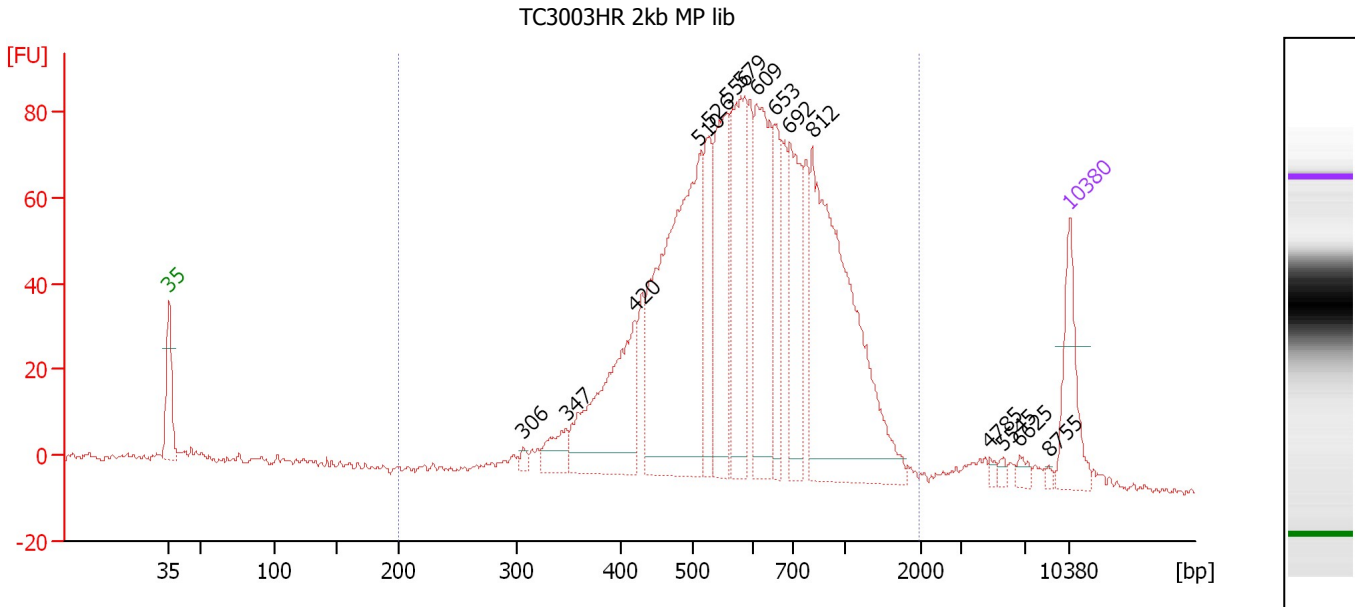
Region table for sample 2 : centaur 1:30 (1:2)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
271	2,466	572	662.3	2,945.6	895.40	90	59.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : TC3003HR 2kb MP lib

Number of peaks found: 15 Corr. Area 1: 1,601.9
 Noise: 0.8

Peak table for sample 3 : TC3003HR 2kb MP lib

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	306	8.95	44.4		70.54
3	347	41.97	183.0		73.88
4	420	244.81	882.6		79.22
5	510	516.64	1,535.2		84.21
6	526	113.18	326.1		84.96
7	556	183.28	499.8		86.36
8	579	211.07	552.4		87.45
9	609	221.04	550.4		88.70
10	653	92.41	214.5		90.05
11	692	152.03	332.7		91.25
12	812	474.52	885.4		92.99
13	4,785	5.40	1.7		106.81
14	5,545	5.39	1.5		107.76
15	6,625	9.08	2.1		109.11
16	8,755	3.64	0.6		111.35
17	10,380	75.00	10.9	Upper Marker	113.00

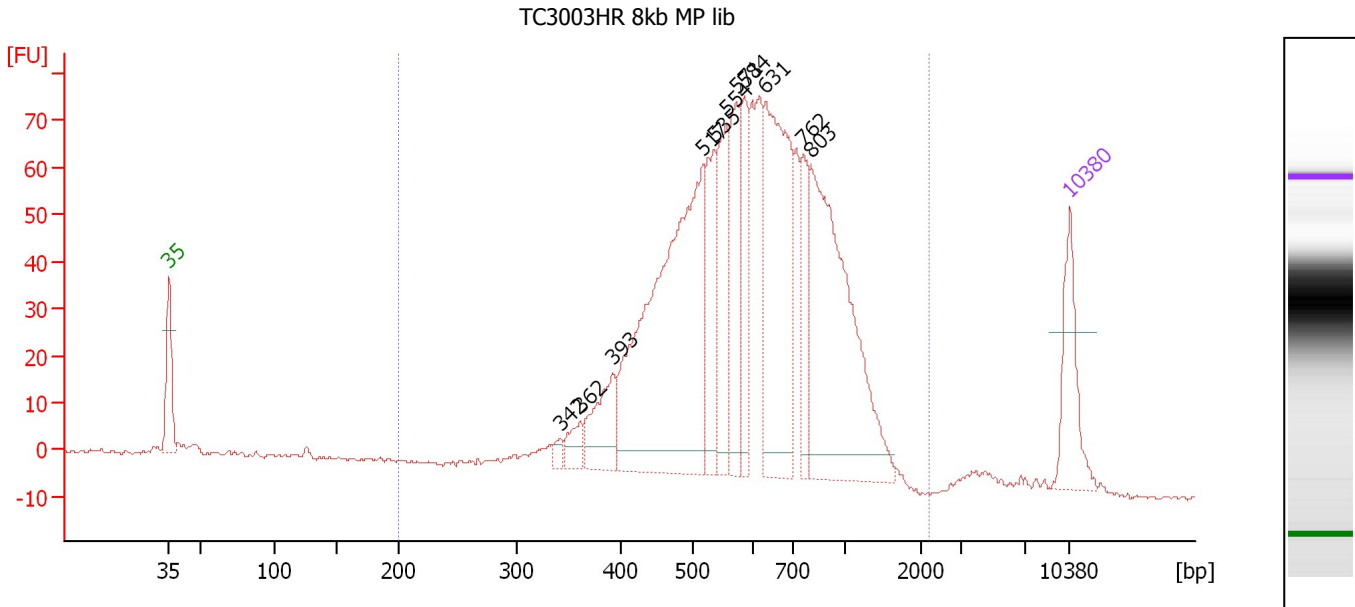
Region table for sample 3 : TC3003HR 2kb MP lib

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,965	651	1,601.9	6,860.2	2,554.71	96	37.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : TC3003HR 8kb MP lib

Number of peaks found: 11 Corr. Area 1: 1,380.5
 Noise: 0.4

Peak table for sample 4 : TC3003HR 8kb MP lib

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	12.28	54.4		73.45
3	362	29.27	122.6		75.02
4	393	86.13	332.3		77.49
5	517	580.02	1,701.3		84.52
6	535	111.44	315.7		85.38
7	554	122.30	334.3		86.29
8	571	127.96	339.2		87.10
9	584	93.21	241.7		87.71
10	631	305.10	732.8		89.38
11	762	81.45	162.0		92.31
12	803	396.90	748.9		92.87
13	10,380	75.00	10.9	Upper Marker	113.00

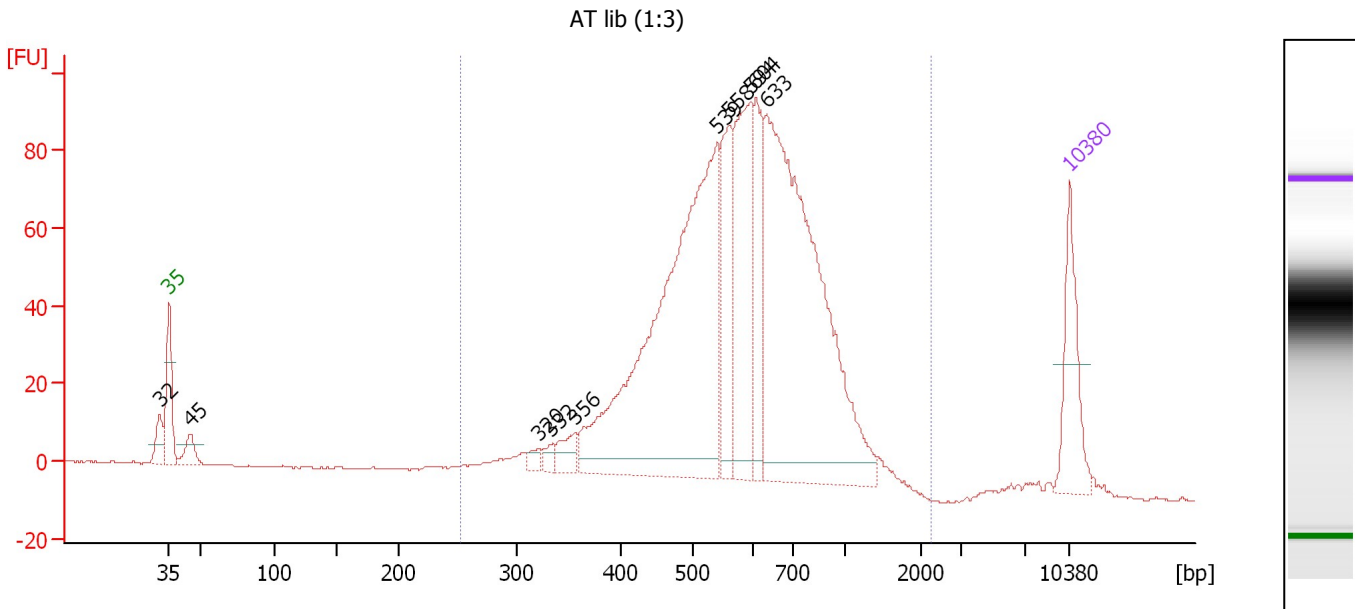
Region table for sample 4 : TC3003HR 8kb MP lib

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	2,183	644	1,380.5	5,907.2	2,209.44	97	34.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : AT lib (1:3)

Number of peaks found: 10 Corr. Area 1: 1,470.4
 Noise: 0.2

Peak table for sample 5 : AT lib (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.28
2	35	125.00	5,411.3	Lower Marker	43.00
3	45	33.34	1,120.2		44.65
4	320	13.00	61.6		71.70
5	332	12.28	56.1		72.63
6	356	28.40	121.0		74.53
7	539	683.13	1,919.8		85.59
8	558	121.60	330.3		86.46
9	594	196.85	502.2		88.16
10	604	109.05	273.5		88.57
11	633	630.75	1,509.8		89.44
12	10,380	75.00	10.9	Upper Marker	113.00

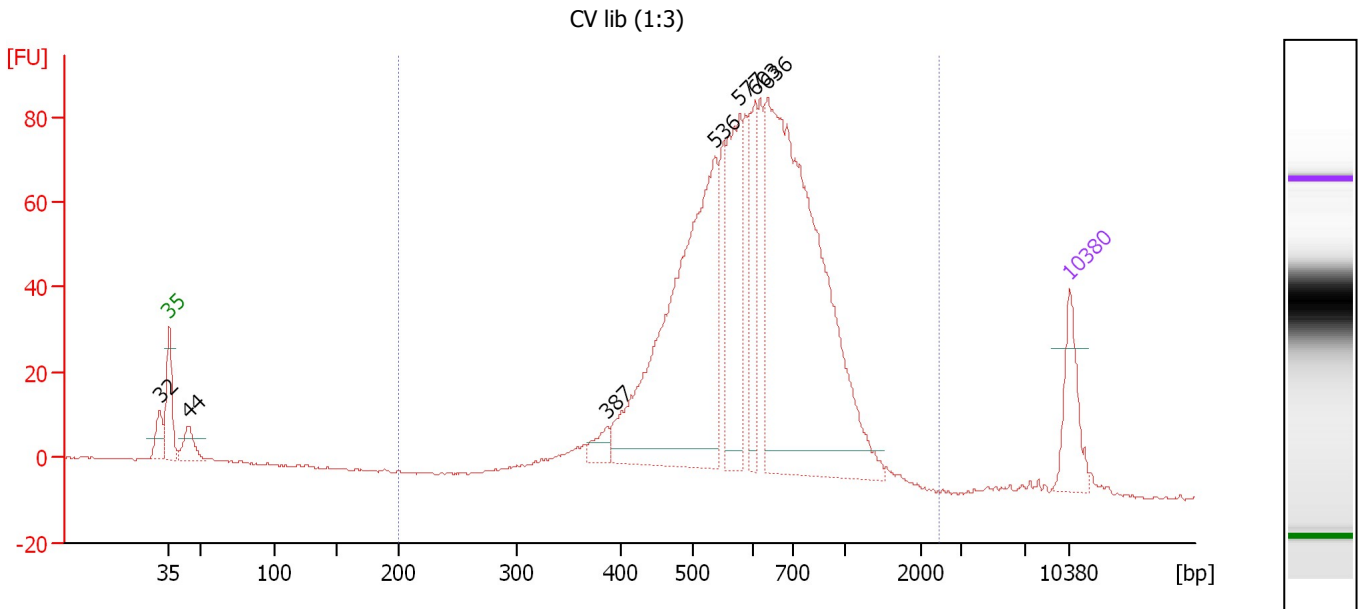
Region table for sample 5 : AT lib (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
253	2,211	607	1,470.4	5,222.3	1,889.12	97	29.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : CV lib (1:3)

Number of peaks found: 7 Corr. Area 1: 1,280.3
 Noise: 0.2

Peak table for sample 6 : CV lib (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.27
2	35	125.00	5,411.3	Lower Marker	43.00
3	44	57.51	1,969.1		44.51
4	387	31.32	122.6		77.04
5	536	734.25	2,077.4		85.42
6	577	241.58	634.9		87.34
7	603	124.16	311.9		88.54
8	636	951.81	2,268.4		89.53
9	10,380	75.00	10.9	Upper Marker	113.00

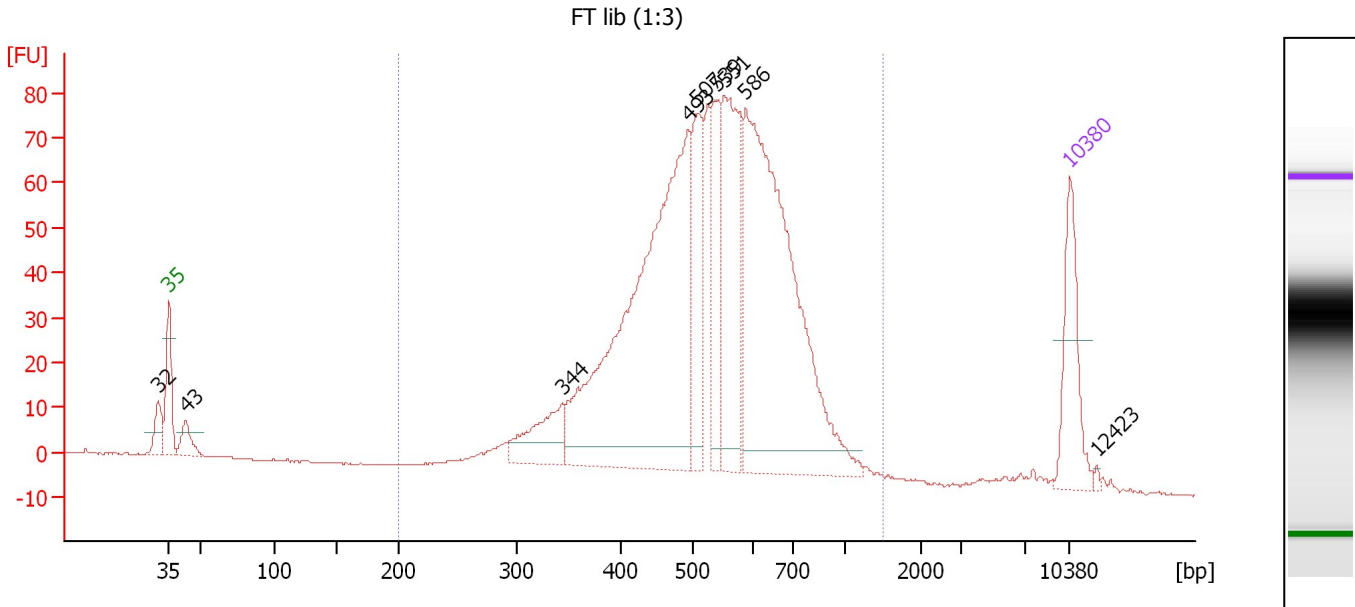
Region table for sample 6 : CV lib (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	2,430	633	1,280.3	6,687.5	2,548.46	97	29.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : FT lib (1:3)

Number of peaks found: 9 Corr. Area 1: 1,345.4
 Noise: 0.2

Peak table for sample 7 : FT lib (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.20
2	35	125.00	5,411.3	Lower Marker	43.00
3	43	32.76	1,150.7		44.33
4	344	75.94	335.0		73.57
5	493	643.74	1,979.6		83.33
6	507	123.20	368.2		84.08
7	539	89.13	250.7		85.56
8	551	207.23	569.8		86.15
9	586	527.70	1,364.1		87.79
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,423	0.00	0.0		115.07

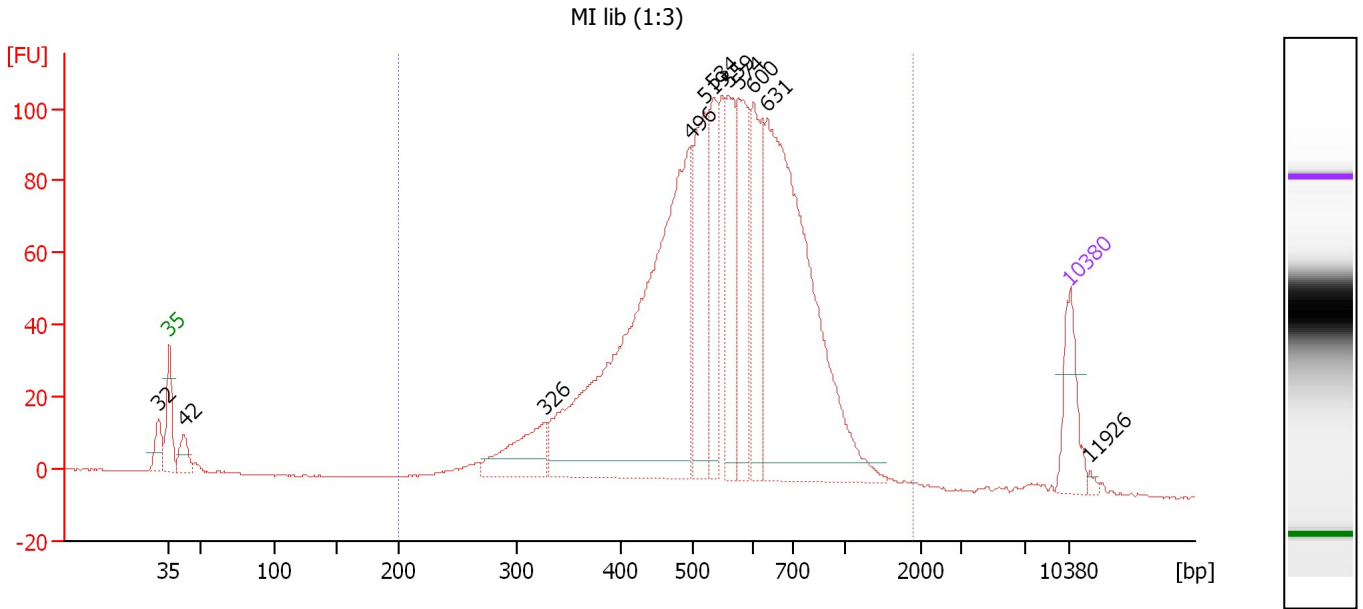
Region table for sample 7 : FT lib (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,488	540	1,345.4	5,692.2	1,863.23	96	25.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : MI lib (1:3)

Number of peaks found: 11 Corr. Area 1: 1,822.3
 Noise: 0.2

Peak table for sample 8 : MI lib (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.19
2	35	125.00	5,411.3	Lower Marker	43.00
3	42	49.02	1,770.2		44.13
4	326	127.85	594.3		72.17
5	496	1,010.12	3,086.9		83.51
6	519	246.35	719.1		84.64
7	534	155.96	442.5		85.35
8	559	162.80	441.0		86.53
9	574	200.17	528.1		87.24
10	600	187.39	473.5		88.42
11	631	764.00	1,833.3		89.40
12	10,380	75.00	10.9	Upper Marker	113.00
13	11,926	0.00	0.0		114.57

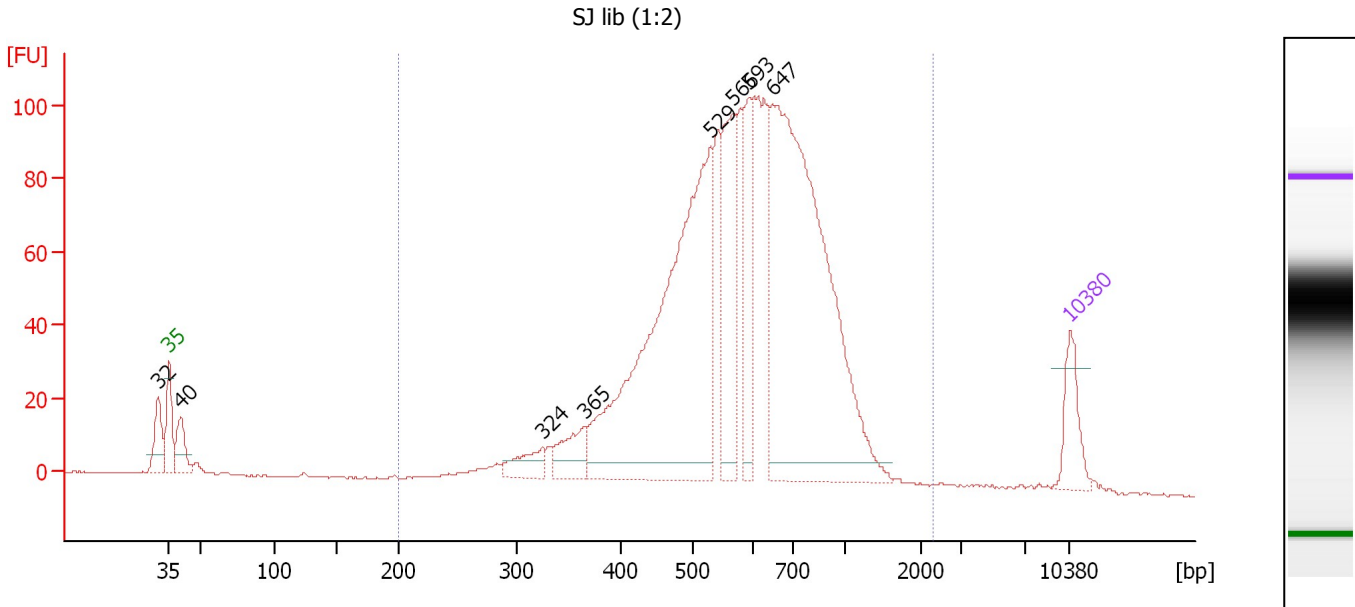
Region table for sample 8 : MI lib (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,874	567	1,822.3	9,381.6	3,146.32	97	30.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : SJ lib (1:2)

Number of peaks found: 8 Corr. Area 1: 1,694.1
 Noise: 0.2

Peak table for sample 9 : SJ lib (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.18
2	35	125.00	5,411.3	Lower Marker	43.00
3	40	85.45	3,207.5		43.88
4	324	67.14	313.7		72.04
5	365	93.05	385.7		75.32
6	529	1,106.19	3,168.3		85.11
7	566	267.94	716.9		86.86
8	593	206.82	528.4		88.12
9	647	1,175.22	2,752.4		89.87
10	10,380	75.00	10.9	Upper Marker	113.00

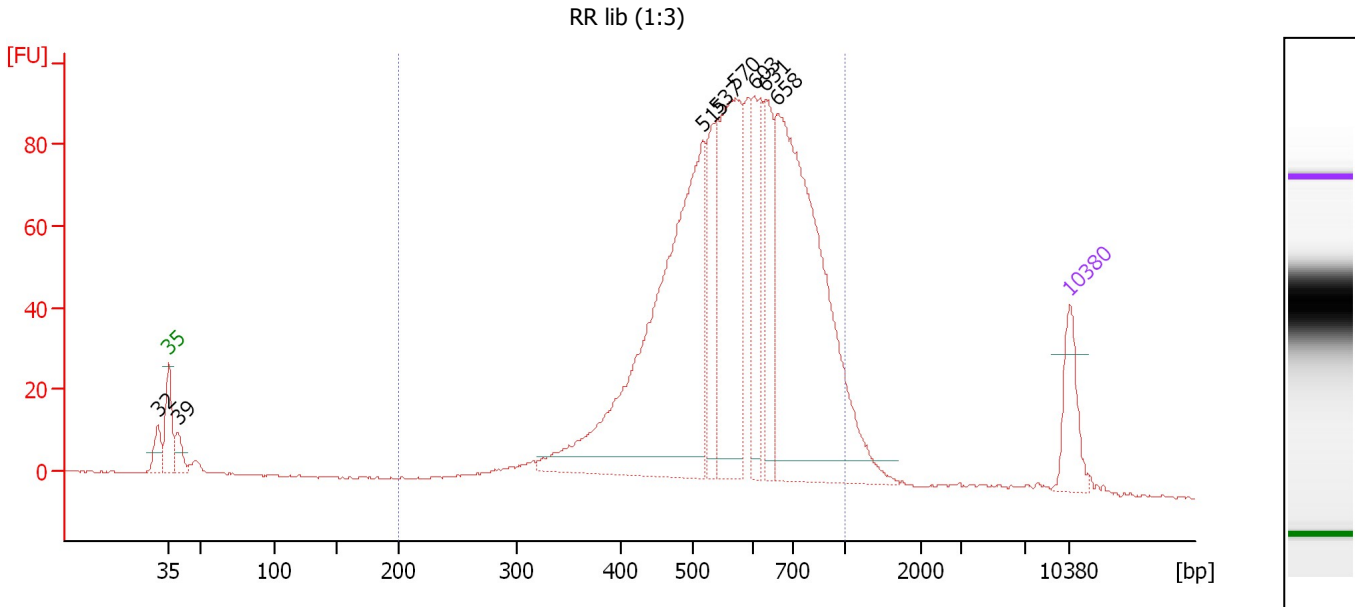
Region table for sample 9 : SJ lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	2,308	619	1,694.1	10,080.5	3,649.42	95	32.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : RR lib (1:3)

Number of peaks found: 8 Corr. Area 1: 1,436.4
 Noise: 0.2

Peak table for sample 10 : RR lib (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.12
2	35	125.00	5,411.3	Lower Marker	43.00
3	39	44.01	1,707.0		43.66
4	515	943.51	2,775.1		84.46
5	537	159.10	448.5		85.51
6	570	415.12	1,102.7		87.05
7	603	158.85	398.9		88.54
8	631	156.36	375.7		89.37
9	658	875.28	2,015.7		90.20
10	10,380	75.00	10.9	Upper Marker	113.00

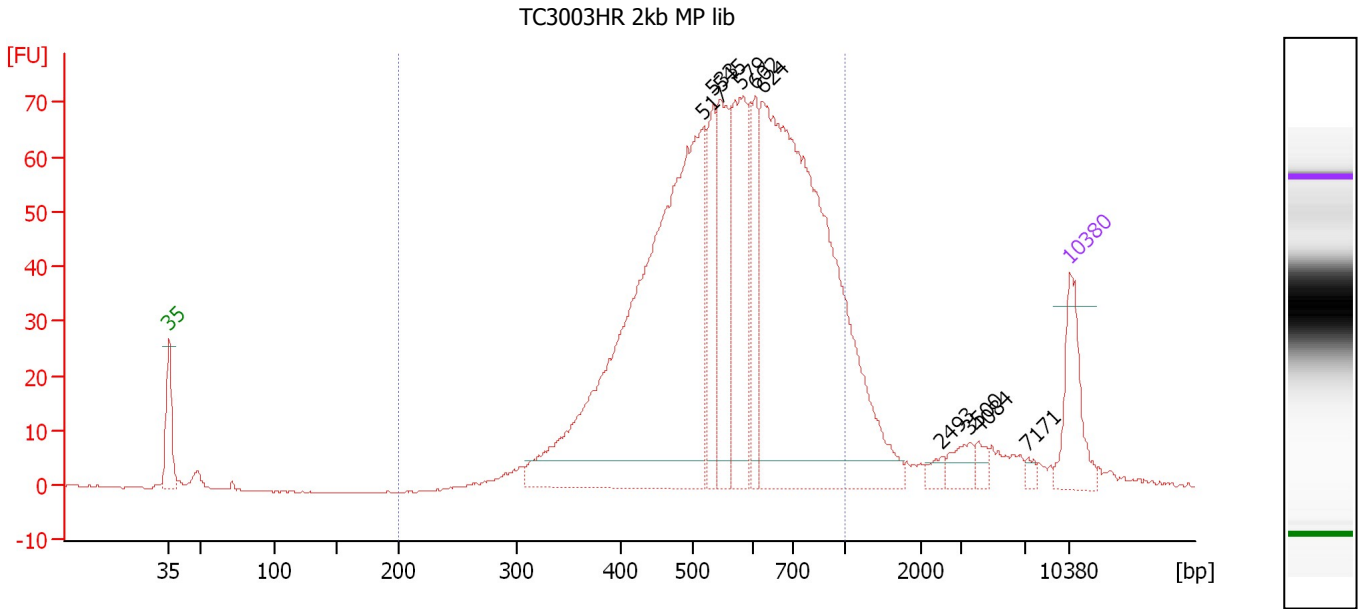
Region table for sample 10 : RR lib (1:3)

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	590	1,436.4	8,481.9	3,046.91	94	24.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
 Modified: 8/4/2016 4:38:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : TC3003HR 2kb MP lib

Number of peaks found: 10 Corr. Area 1: 1,245.4
 Noise: 0.2

Peak table for sample 11 : TC3003HR 2kb MP lib

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	517	1,007.36	2,954.3		84.53
3	533	116.21	330.1		85.31
4	545	171.38	476.2		85.87
5	579	208.63	546.2		87.44
6	602	109.18	274.7		88.51
7	624	968.58	2,350.7		89.18
8	2,493	12.23	7.4		103.02
9	3,500	24.81	10.7		105.21
10	4,084	12.67	4.7		105.94
11	7,171	7.30	1.5		109.75
12	10,380	75.00	10.9	Upper Marker	113.00

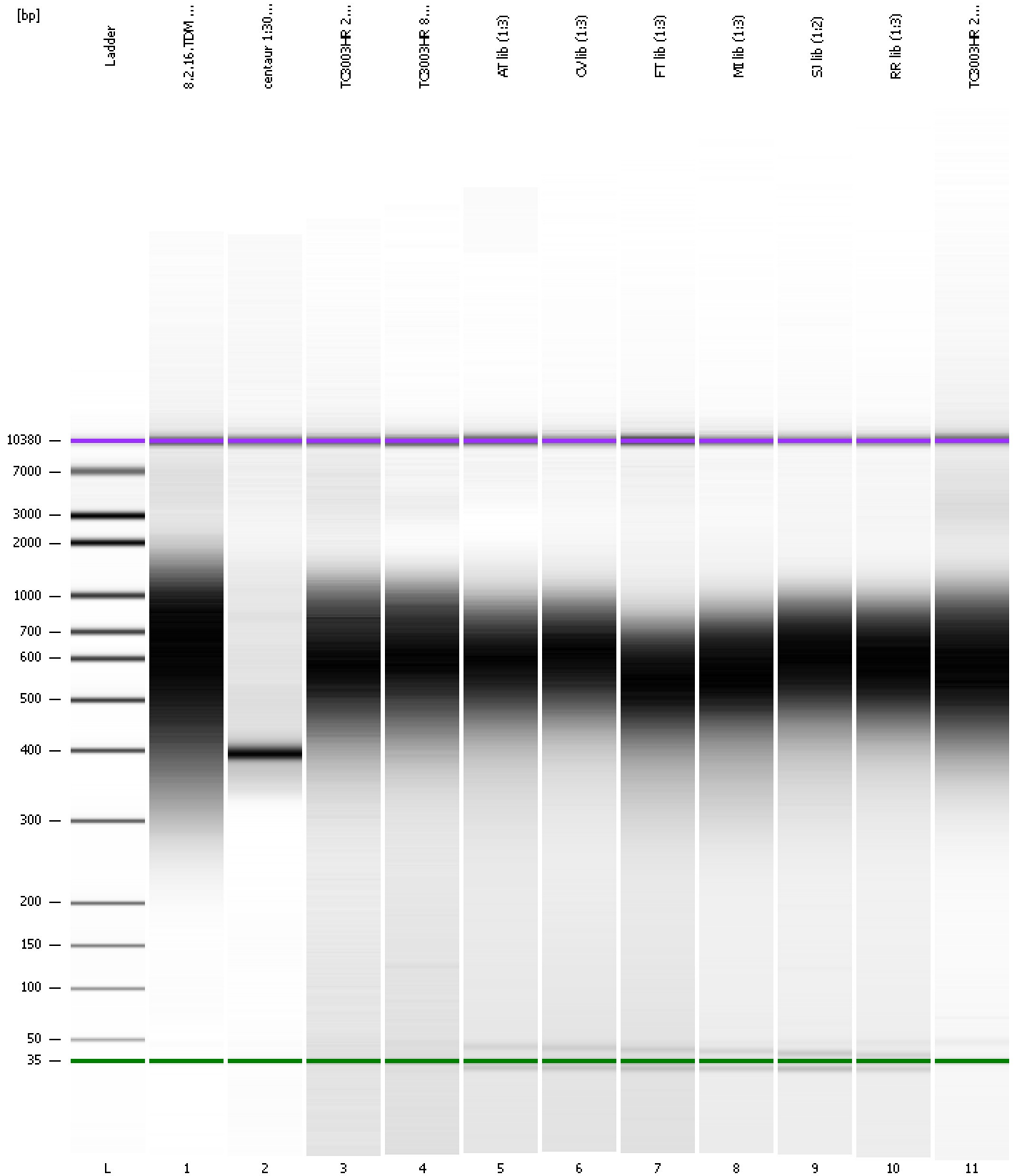
Region table for sample 11 : TC3003HR 2kb MP lib

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	581	1,245.4	7,397.3	2,586.17	87	26.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad

Created: 8/4/2016 3:51:07 PM
Modified: 8/4/2016 4:38:30 PM

Gel Image



Assay Class: High Sensitivity DNA Assay Created: 8/4/2016 3:51:07 PM
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad Modified: 8/4/2016 4:38:30 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		8/4/2016 4:32:27 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-08-04\2016-08-04_004.xad)		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		8/4/2016 3:51:13 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1