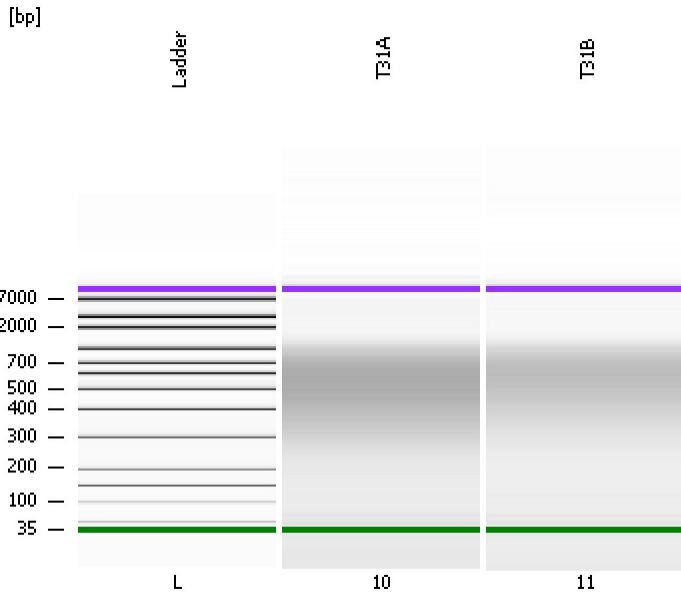


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
Modified: 8/25/2016 4:26:45 PM

Electrophoresis File Run Summary



Instrument Information:

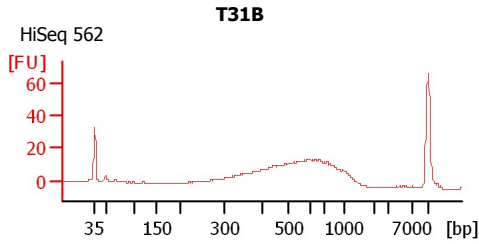
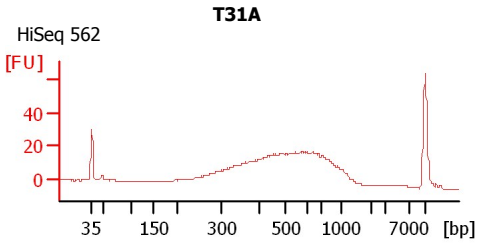
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
 Modified: 8/25/2016 4:26:45 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
T31A	HiSeq 562	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
T31B	HiSeq 562	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
Ladder		<input type="checkbox"/>	<input checked="" type="checkbox"/>			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
Modified: 8/25/2016 4:26:45 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

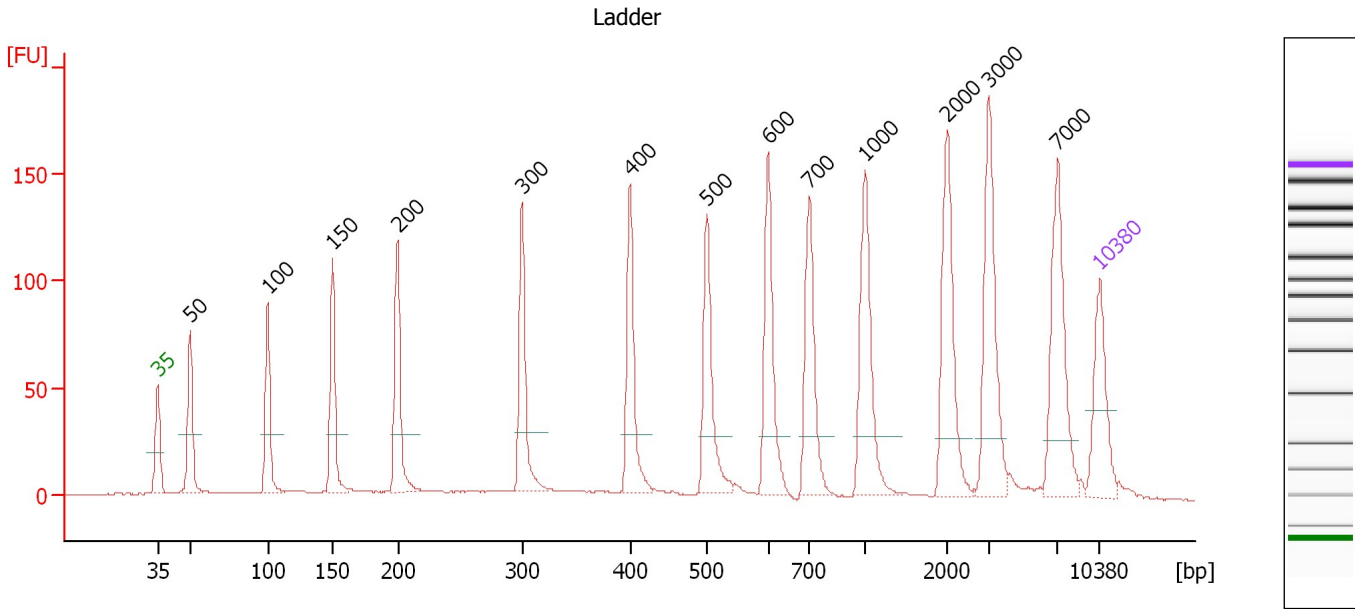
Ladder

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:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
 Modified: 8/25/2016 4:26:45 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

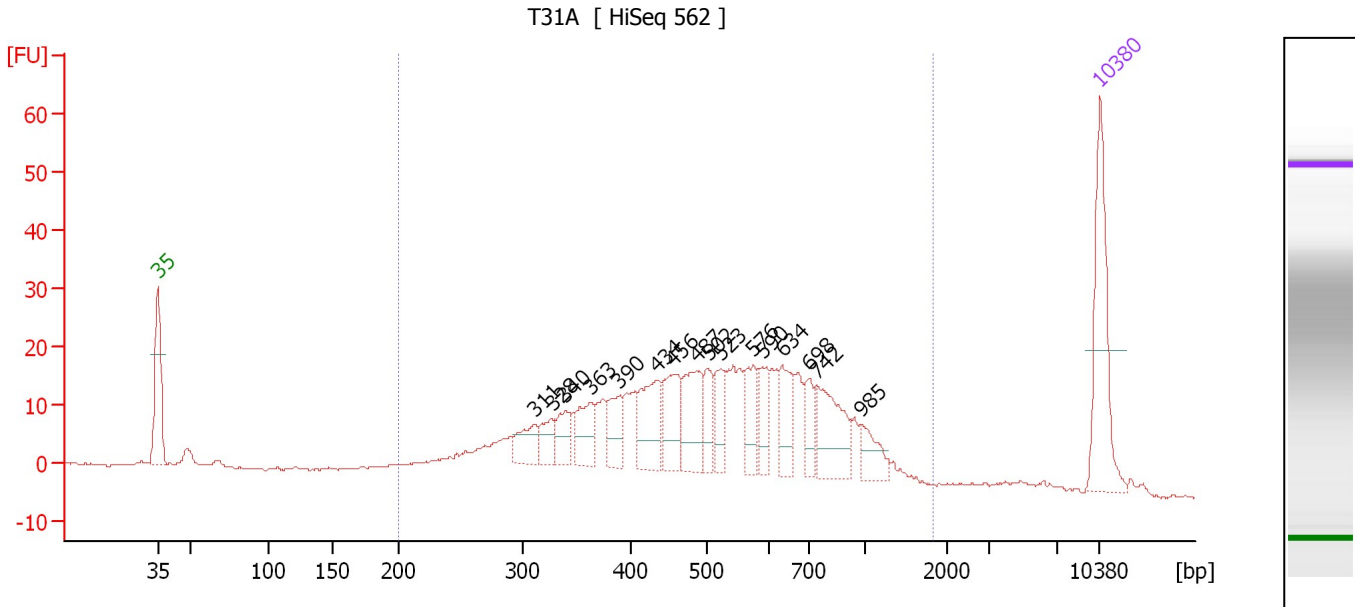
Peak table for Ladder

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	45.44
3	100	150.00	2,272.7	Ladder Peak	51.18
4	150	150.00	1,515.2	Ladder Peak	56.00
5	200	150.00	1,136.4	Ladder Peak	60.83
6	300	150.00	757.6	Ladder Peak	70.08
7	400	150.00	568.2	Ladder Peak	78.10
8	500	150.00	454.5	Ladder Peak	83.79
9	600	150.00	378.8	Ladder Peak	88.36
10	700	150.00	324.7	Ladder Peak	91.41
11	1,000	150.00	227.3	Ladder Peak	95.53
12	2,000	150.00	113.6	Ladder Peak	101.62
13	3,000	150.00	75.8	Ladder Peak	104.77
14	7,000	150.00	32.5	Ladder Peak	109.85
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
 Modified: 8/25/2016 4:26:45 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : T31A

Number of peaks found: 16 Corr. Area 1: 486.0
 Noise: 0.2

Peak table for sample 10 : T31A

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	311	28.51	138.9		70.97
3	328	22.80	105.5		72.29
4	340	24.15	107.7		73.26
5	363	36.45	152.0		75.16
6	390	29.64	115.2		77.28
7	434	53.68	187.4		80.04
8	456	47.50	157.7		81.30
9	487	52.52	163.6		83.02
10	502	26.64	80.4		83.89
11	523	28.26	81.8		84.86
12	576	33.05	86.9		87.27
13	590	26.85	68.9		87.91
14	634	35.83	85.6		89.40
15	698	22.16	48.1		91.35
16	742	59.59	121.7		91.98
17	985	22.63	34.8		95.31
18	10,380	75.00	10.9	Upper Marker	113.00

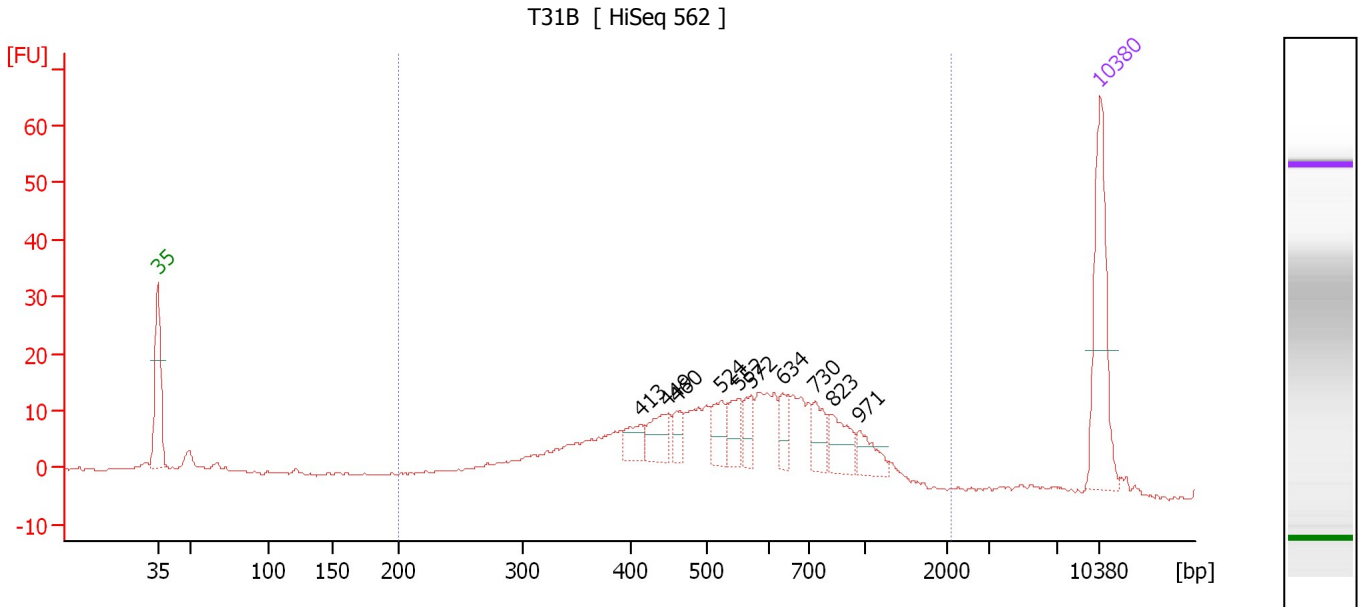
Region table for sample 10 : T31A

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,837	539	486.0	3,002.9	886.18	97	39.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
 Modified: 8/25/2016 4:26:45 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : T31B

Number of peaks found: 10 Corr. Area 1: 328.5
 Noise: 0.2

Peak table for sample 11 : T31B

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	413	19.75	72.4		78.87
3	449	27.84	93.9		80.89
4	460	14.57	48.0		81.53
5	524	22.67	65.6		84.88
6	552	22.03	60.5		86.16
7	572	16.23	43.0		87.08
8	634	16.77	40.1		89.40
9	730	22.00	45.6		91.83
10	823	27.55	50.7		93.10
11	971	18.50	28.9		95.12
12	10,380	75.00	10.9	Upper Marker	113.00

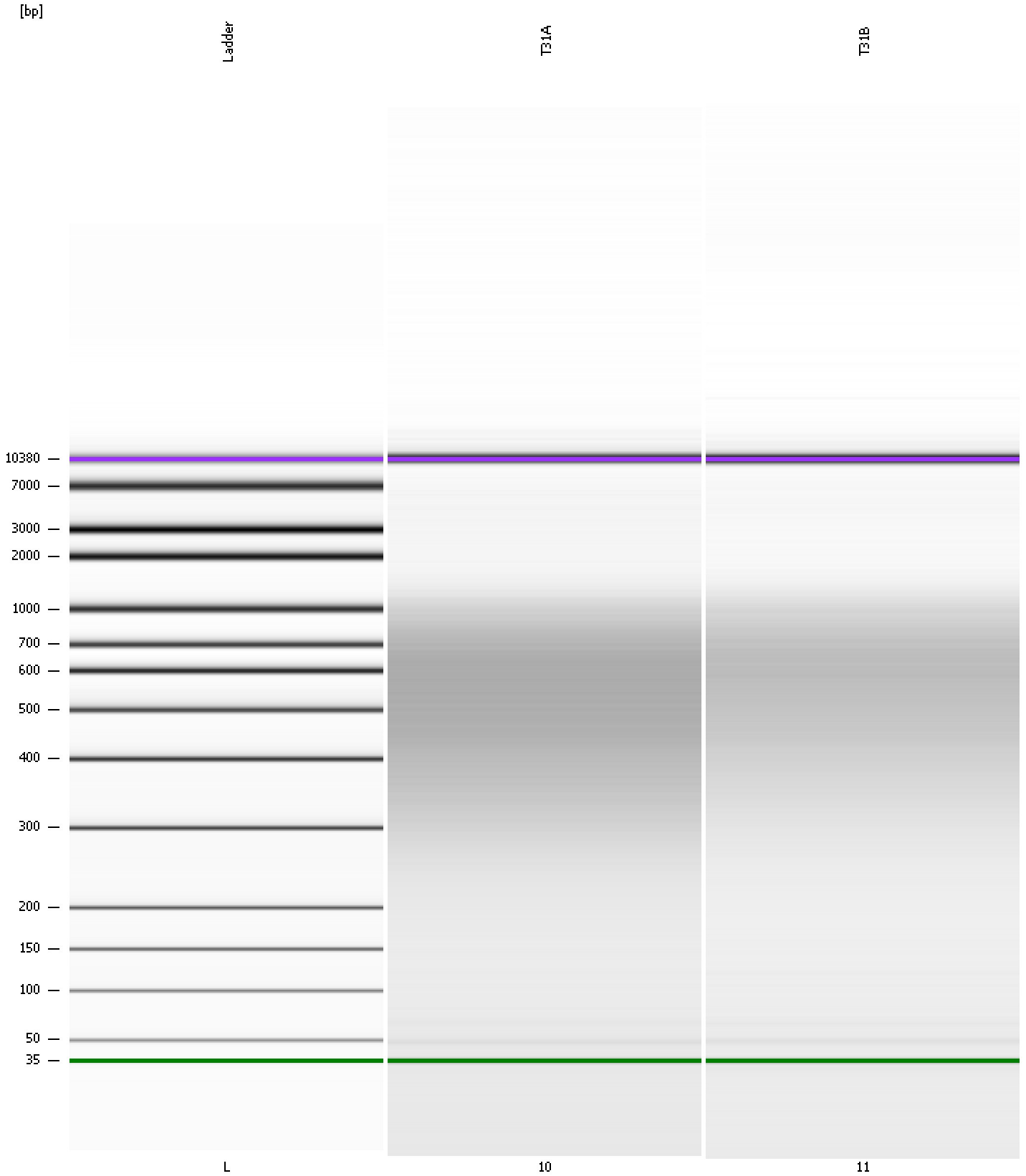
Region table for sample 11 : T31B

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	2,121	580	328.5	1,733.5	557.99	96	38.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad

Created: 8/25/2016 3:37:46 PM
Modified: 8/25/2016 4:26:45 PM

Gel Image



Assay Class: High Sensitivity DNA Assay Created: 8/25/2016 3:37:46 PM
 Data Path: C:\...xpert\data\2016-08-25\2016-08-25_003_HiSeq652_T31A_T31B.xad Modified: 8/25/2016 4:26:45 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		8/25/2016 4:19:04 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\2016-08-25\2016-08-25_003.xad)		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		8/25/2016 3:37:52 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1