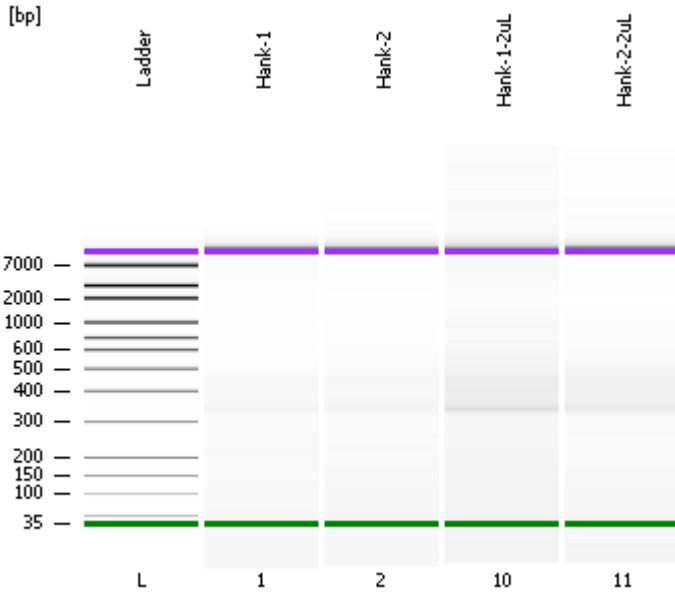


Assay Class: High Sensitivity DNA Assay
Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
Modified: 9/20/2012 3:44:16 PM

Electrophoresis File Run Summary



Instrument Information:

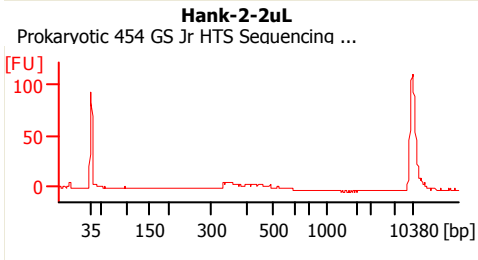
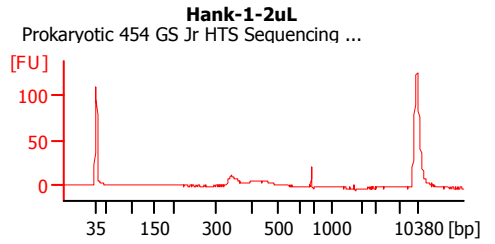
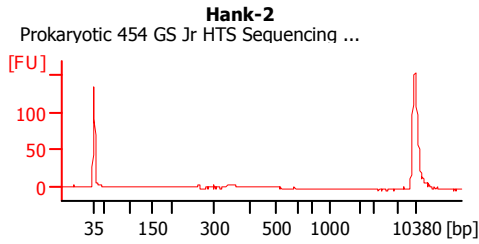
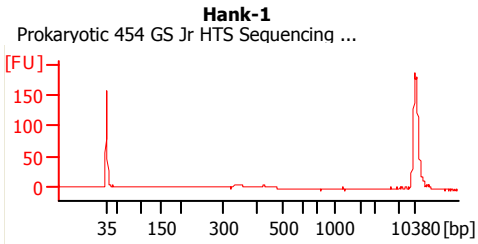
Instrument Name: DE24802282 Firmware: C.01.069
Serial#: DE24802282 Type: G2938B

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 #: QD28BK50
Reagent Kit Lot #: set B
Chip Comments: HS-DNA | High Sensitivity DNA | Hank Raizen | Nik Grundwald | 738-4105 | nraizen@gmail.com | Unknown | Unknown | Submitted by Mark



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\AqilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
 Modified: 9/20/2012 3:44:16 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Hank-1	Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
Hank-2	Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
Hank-1-2uL	Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
Hank-2-2uL	Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2	<input type="checkbox"/>	<input checked="" type="checkbox"/>			
Ladder		<input type="checkbox"/>	<input checked="" type="checkbox"/>			

Chip Lot #

QD28BK50

Reagent Kit Lot #

set B

Chip Comments :

HS-DNA | High Sensitivity DNA | Hank Raizen | Nik Grundwald | 738-4105 | nraizen@gmail.com | Unknown | Unknown | Submitted by Mark

Assay Class: High Sensitivity DNA Assay
Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
Modified: 9/20/2012 3:44:16 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/ μ l] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/ μ l] : 125
Upper Marker Concentration [pg/ μ l] : 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

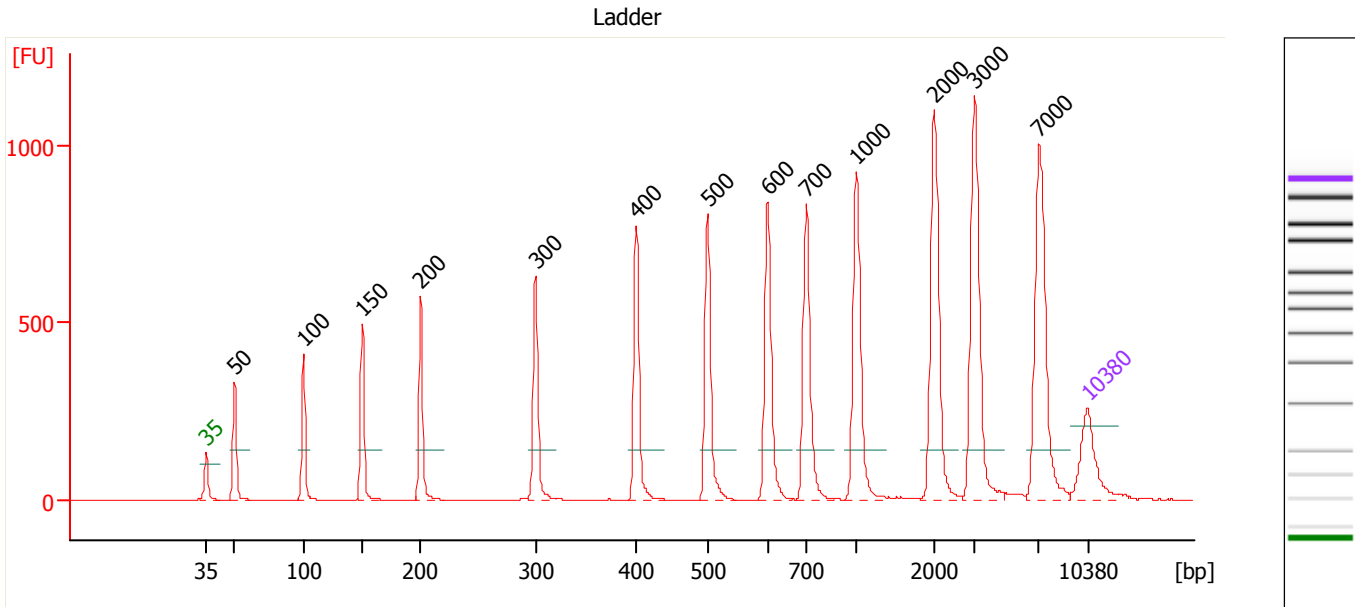
Ladder

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:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

Peak table for Ladder

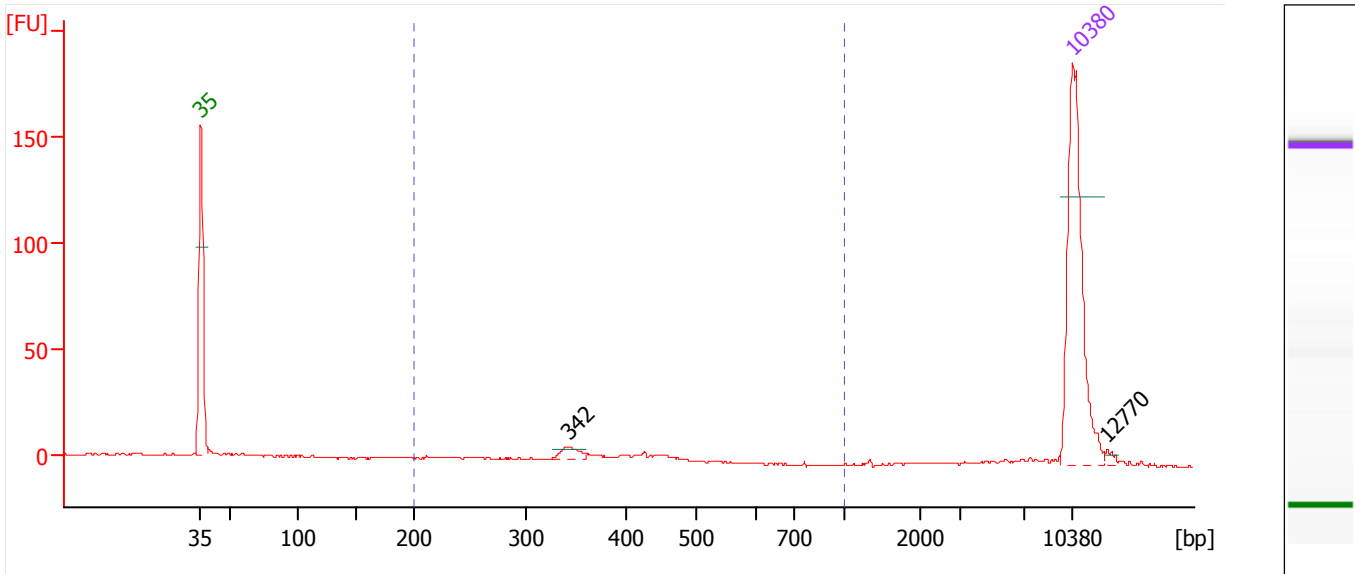
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	Ladder Peak
9	600	150.00	378.8	Ladder Peak
10	700	150.00	324.7	Ladder Peak
11	1,000	150.00	227.3	Ladder Peak
12	2,000	150.00	113.6	Ladder Peak
13	3,000	150.00	75.8	Ladder Peak
14	7,000	150.00	32.5	Ladder Peak
15	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
 Modified: 9/20/2012 3:44:16 PM

Electropherogram Summary Continued ...

Hank-1 [Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2]



Overall Results for sample 1 : Hank-1

Number of peaks found: 2 Corr. Area 1: 57.0
 Noise: 0.2

Peak table for sample 1 : Hank-1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	342	7.66	33.9	
3	10,380	75.00	10.9	Upper Marker
4	12,770	0.00	0.0	

Region table for sample 1 : Hank-1

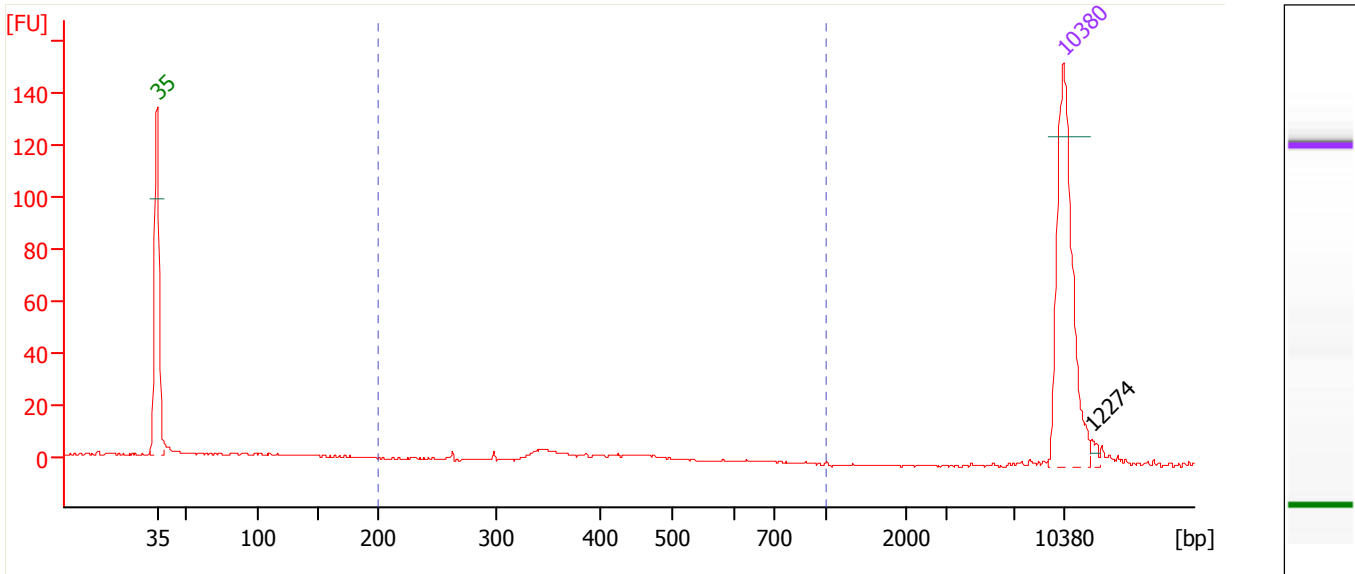
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	57.0	45	366	19.0	27.11	119.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
 Modified: 9/20/2012 3:44:16 PM

Electropherogram Summary Continued ...

Hank-2 [Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2]



Overall Results for sample 2 : Hank-2

Number of peaks found: 1 Corr. Area 1: 69.1
 Noise: 0.3

Peak table for sample 2 : Hank-2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	10,380	75.00	10.9	Upper Marker
3	12,274	0.00	0.0	

Region table for sample 2 : Hank-2

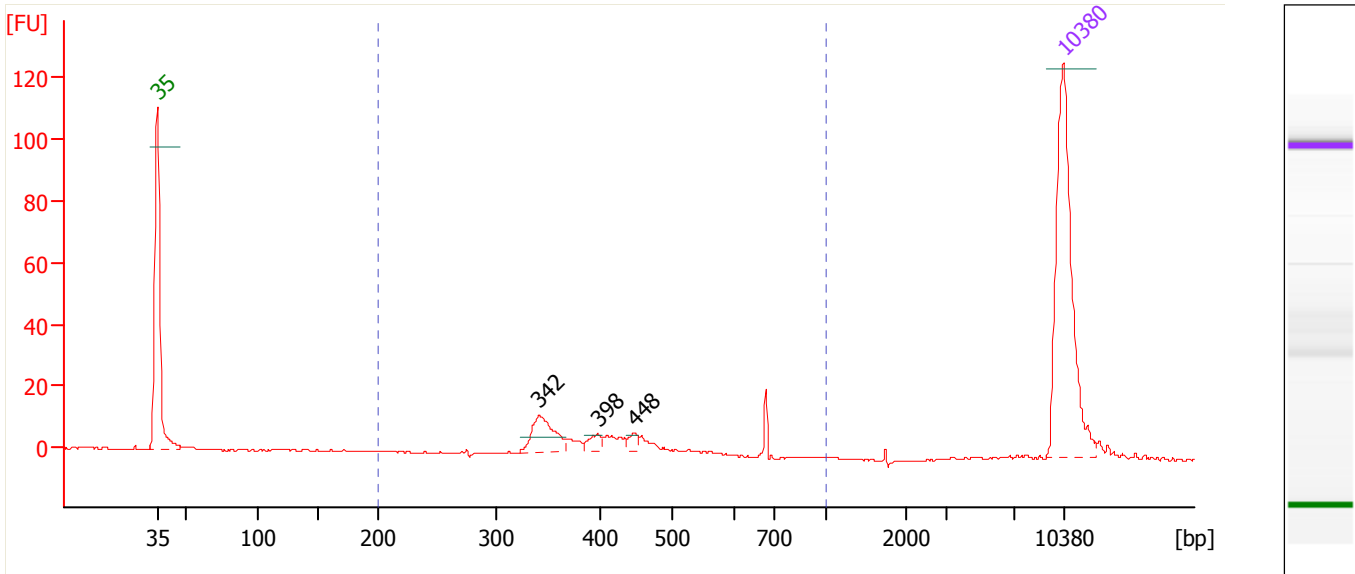
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	69.1	38	408	28.6	38.87	161.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
 Modified: 9/20/2012 3:44:16 PM

Electropherogram Summary Continued ...

Hank-1-2uL [Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2]



Overall Results for sample 10 : Hank-1-2uL

Number of peaks found: 3 Corr. Area 1: 84.3
 Noise: 0.3

Peak table for sample 10 : Hank-1-2uL

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	342	23.74	105.3	
3	398	5.59	21.3	
4	448	4.27	14.4	
5	10,380	75.00	10.9	Upper Marker

Region table for sample 10 : Hank-1-2uL

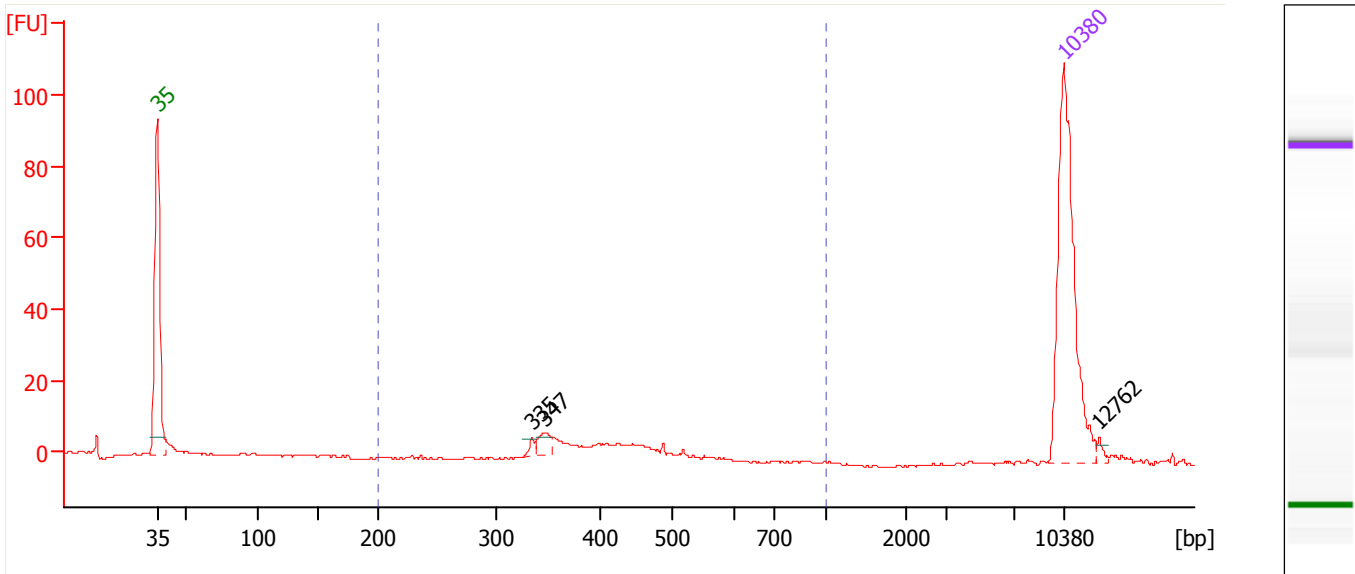
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	84.3	90	418	21.6	58.03	220.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\AgilentData\2012-09-20\HS2005_Raizen.xad

Created: 9/20/2012 2:18:21 PM
 Modified: 9/20/2012 3:44:16 PM

Electropherogram Summary Continued ...

Hank-2-2uL [Prokaryotic 454 GS Jr HTS Sequencing Prep 5 2]



Overall Results for sample 11 : Hank-2-2uL

Number of peaks found: 3 Corr. Area 1: 61.0
 Noise: 0.3

Peak table for sample 11 : Hank-2-2uL

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	335	2.77	12.5	
3	347	7.79	34.0	
4	10,380	75.00	10.9	Upper Marker
5	12,762	0.00	0.0	

Region table for sample 11 : Hank-2-2uL

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	61.0	69	406	13.9	47.18	180.2	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\AqilentData\2012-09-20\HS2005 Raizen.xad

Created: 9/20/2012 2:18:21 PM
Modified: 9/20/2012 3:44:16 PM

Gel Image

