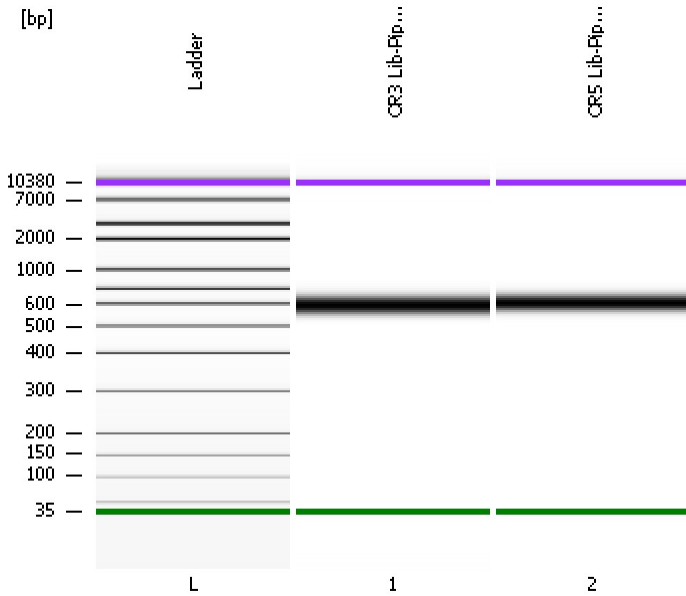


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
Modified: 10/15/2013 5:43:30 PM

Electrophoresis File Run Summary



Instrument Information:

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

Assay Information:

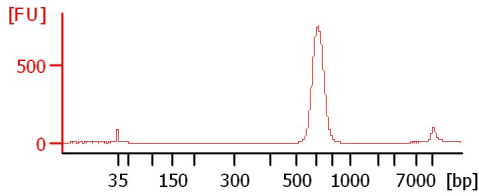
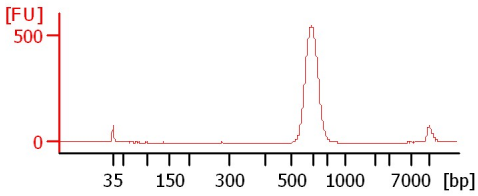
Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy
Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

Chip Information:

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

OR3 Lib-Pipp 1:2

OR5 Lib-Pipp 1:2



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
Modified: 10/15/2013 5:43:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
OR3 Lib-Pipp 1:2		<input type="checkbox"/>	✓			
OR5 Lib-Pipp 1:2		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
 Modified: 10/15/2013 5:43:30 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

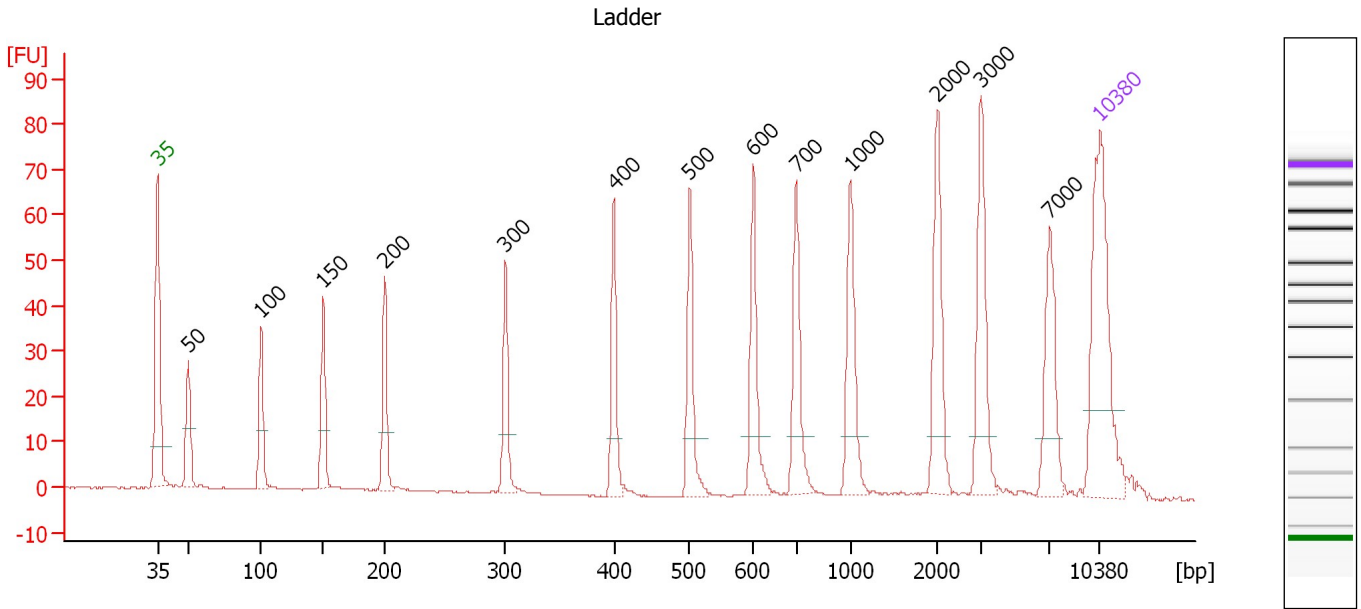
Ladder

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:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

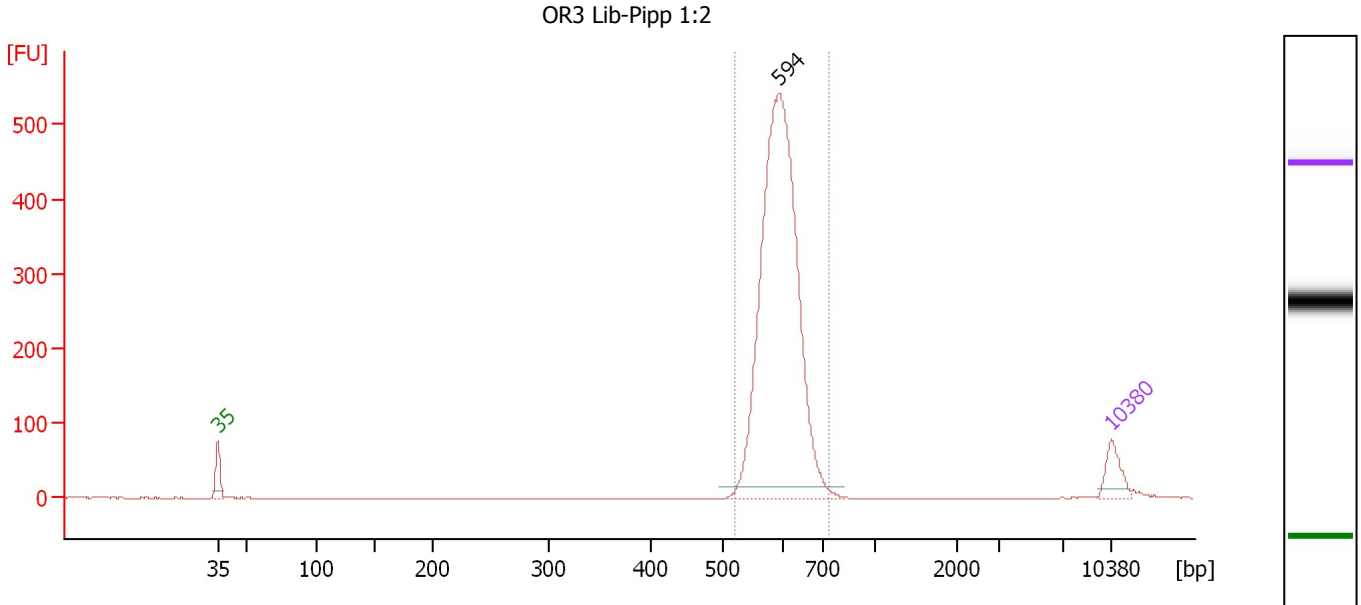
Peak table for Ladder

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:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

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Electropherogram Summary Continued ...



Setpoint Deviations for sample 1 : OR3 Lib-Pipp 1:2

Height Threshold [FU] : 15

Overall Results for sample 1 : OR3 Lib-Pipp 1:2

Number of peaks found: 1 Corr. Area 1: 2,180.0
 Noise: 0.2

Peak table for sample 1 : OR3 Lib-Pipp 1:2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	594	2,201.34	5,616.6	
3	10,380	75.00	10.9	Upper Marker

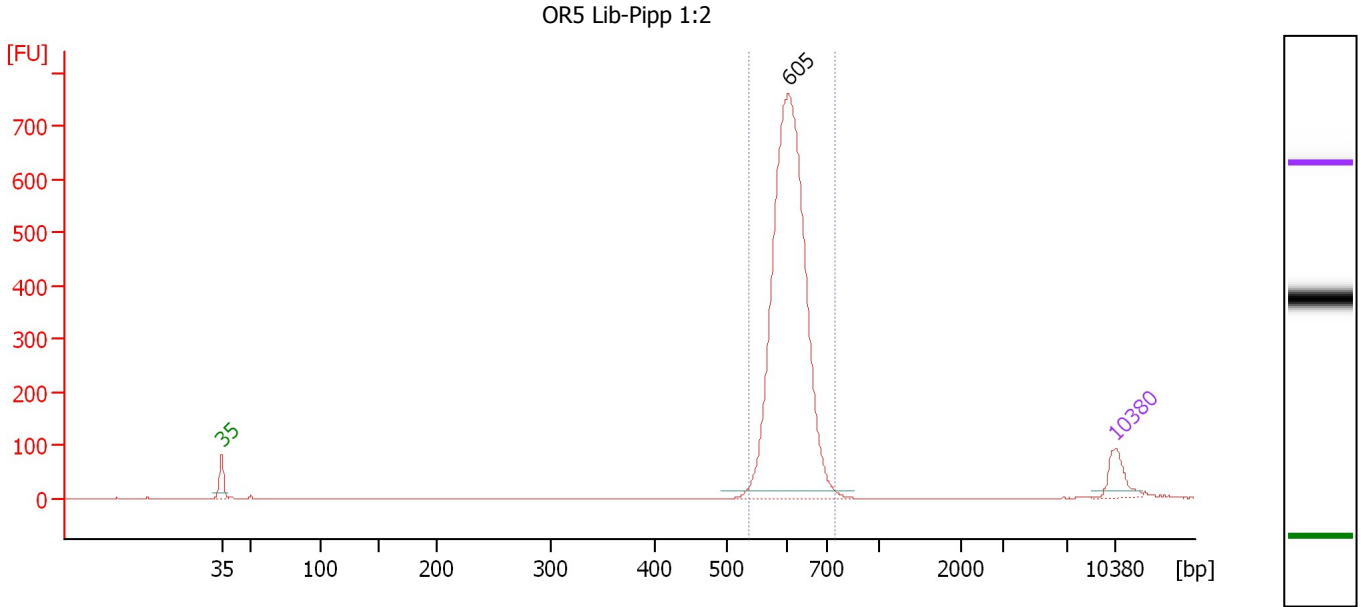
Region table for sample 1 : OR3 Lib-Pipp 1:2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
519	733	599	5,587.9	2,203.47	2,180.0	99	5.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
 Modified: 10/15/2013 5:43:30 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 2 : OR5 Lib-Pipp 1:2

Height Threshold [FU] : 15

Overall Results for sample 2 : OR5 Lib-Pipp 1:2

Number of peaks found: 1 Corr. Area 1: 2,745.7
 Noise: 0.1

Peak table for sample 2 : OR5 Lib-Pipp 1:2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	605	2,140.26	5,361.7	
3	10,380	75.00	10.9	Upper Marker

Region table for sample 2 : OR5 Lib-Pipp 1:2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
538	749	612	5,302.8	2,135.67	2,745.7	99	5.3	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...tings\Bioanalyzer\2013-10-15\2013-10-15_001_OR_Lib_Pipp.xad

Created: 10/15/2013 4:57:58 PM
Modified: 10/15/2013 5:43:30 PM

Gel Image

